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The following documentation is split into four parts. *About Bio-Formats* explains the goal of the software, discusses how it processes metadata, and provides other useful information such as version history and how to report bugs. *User Information* focuses on how to use Bio-Formats as a plugin for ImageJ and Fiji, and also gives details of other software packages which can use Bio-Formats to read and write microscopy formats. *Developer Documentation* covers more in-depth information on using Bio-Formats as a Java library and how to interface from non-Java codes. Finally, *Formats* is a guide to all the file formats currently supported by Bio-Formats.
Part I

About Bio-Formats
Bio-Formats is a standalone Java library for reading and writing life sciences image file formats. It is capable of parsing both pixels and metadata for a large number of formats, as well as writing to several formats.

The primary goal of Bio-Formats is to facilitate the exchange of microscopy data between different software packages and organizations. It achieves this by converting proprietary microscopy data into an open standard called the OME data model\(^1\), particularly into the OME-TIFF\(^2\) file format.

We believe the standardization of microscopy metadata to a common structure is of vital importance to the community. A brief article on the benefits of standardization\(^3\) from thinkstandards.net\(^4\) provides an excellent summary. See also LOCI’s article on open source software in science\(^5\).

\(^1\)http://genomebiology.com/2005/6/5/R47
\(^2\)http://www.openmicroscopy.org/site/support/ome-model/ome-tiff
\(^3\)http://www.thinkstandards.net/benefits.html
\(^4\)http://www.thinkstandards.net/
\(^5\)http://loci.wisc.edu/software/oss
WHY JAVA?

From a practical perspective, Bio-Formats is written in Java because it is cross-platform and widely used, with a vast array of libraries for handling common programming tasks. Java is one of the easiest languages from which to deploy cross-platform software. In contrast to C++, which has a large number of complex platform issues to consider, and Python, which leans heavily on C and C++ for many of its components (e.g., NumPy and SciPy), Java code is compiled one time into platform-independent byte code, which can be deployed as is to all supported platforms. And despite this enormous flexibility, Java manages to provide time performance nearly equal to C++, often better in the case of I/O operations (see further discussion on the comparative speed of Java on the LOCI site¹).

There are also historical reasons associated with the fact that the project grew out of work on the VisAD Java component library². You can read more about the origins of Bio-Formats on the LOCI Bio-Formats homepage³.

¹http://loci.wisc.edu/faq/isnt-java-too-slow
²http://visad.ssec.wisc.edu
³http://loci.wisc.edu/software/bio-formats
BIO-FORMATS METADATA PROCESSING

Pixels in microscopy are almost always very straightforward, stored on evenly spaced rectangular grids. It is the metadata (details about the acquisition, experiment, user, and other information) that can be complex. Using the OME data model enables applications to support a single metadata format, rather than the multitude of proprietary formats available today.

Every file format has a distinct set of metadata, stored differently. Bio-Formats processes and converts each format’s metadata structures into a standard form called the **OME data model**\(^1\), according to the **OME-XML**\(^2\) specification. We have defined an open exchange format called **OME-TIFF**\(^3\) that stores its metadata as OME-XML. Any software package that supports OME-TIFF is also compatible with the dozens of formats listed on the Bio-Formats page, because Bio-Formats can convert your files to OME-TIFF format.

To facilitate support of OME-XML, we have created a library in **Java**\(^4\) for reading and writing OME-XML metadata.

There are three types of metadata in Bio-Formats, which we call core metadata, original metadata, and OME metadata.

1. **Core metadata** only includes things necessary to understand the basic structure of the pixels: image resolution; number of focal planes, time points, channels, and other dimensional axes; byte order; dimension order; color arrangement (RGB, indexed color or separate channels); and thumbnail resolution.

2. **Original metadata** is information specific to a particular file format. These fields are key/value pairs in the original format, with no guarantee of cross-format naming consistency or compatibility. Nomenclature often differs between formats, as each vendor is free to use their own terminology.

3. **OME metadata** is information from #1 and #2 converted by Bio-Formats into the OME data model. **Performing this conversion is the primary purpose of Bio-Formats.** Bio-Formats uses its ability to convert proprietary metadata into OME-XML as part of its integration with the OME and OMERO servers—essentially, they are able to populate their databases in a structured way because Bio-Formats sorts the metadata into the proper places. This conversion is nowhere near complete or bug free, but we are constantly working to improve it. We would greatly appreciate any and all input from users concerning missing or improperly converted metadata fields.

---

2. [http://www.openmicroscopy.org/site/support/ome-model/ome-xml](http://www.openmicroscopy.org/site/support/ome-model/ome-xml)
5. [http://www.openmicroscopy.org/site/support/ome-model/ome-xml](http://www.openmicroscopy.org/site/support/ome-model/ome-xml)
For help, see the Bio-Formats, File Formats and OME-XML and OME-TIFF sections of the OME FAQ for answers to some common questions. Please contact us if you have any questions or problems with Bio-Formats. There is a guide for reporting bugs here.

For advanced users and developers, further information is available on the troubleshooting page.

3.1 Reporting a bug

3.1.1 Before filing a bug report

If you think you have found a bug in Bio-Formats, the first thing to do is update your version of Bio-Formats to the latest version to check if the problem has already been addressed. The Fiji updater will automatically do this for you, while in ImageJ you can select Plugins → Bio-Formats → Update Bio-Formats Plugins.

You can also download the latest version of Bio-Formats. If you are not sure which version you need, select the latest build of the Bio-Formats package bundle from the components table.

3.1.2 Sending a bug report

If you can still reproduce the bug after updating to the latest version of Bio-Formats, please send a bug report to the OME Users mailing list. You can upload files to our QA system or for large files (>2 GB), we can provide you with an FTP server address if you write to the mailing list.

To ensure that any inquiries you make are resolved promptly, please include the following information:

- **Exact error message.** Copy and paste any error messages into the text of your email. Alternatively, attach a screenshot of the relevant windows.

- **Version information.** Indicate which release of Bio-Formats, which operating system, and which version of Java you are using.

- **Non-working data.** If possible, please send a non-working file. This helps us ensure that the problem is fixed for next release and will not reappear in later releases. Note that any data provided is used for internal testing only; we do not make images publicly available unless given explicit permission to do so.

- **Metadata and screenshots.** If possible, include any additional information about your data. We are especially interested in the expected dimensions (width, height, number of channels, Z slices, and timepoints). Screenshots of the image being successfully opened in other software are also useful.

- **Format details.** If you are requesting support for a new format, we ask that you send as much data as you have regarding this format (sample files, specifications, vendor/manufacturer information, etc.). This helps us to better support the format and ensures future versions of the format are also supported.

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1. [http://www.openmicroscopy.org/site/support/faq/bio-formats](http://www.openmicroscopy.org/site/support/faq/bio-formats)
2. [http://www.openmicroscopy.org/site/support/faq/file-formats](http://www.openmicroscopy.org/site/support/faq/file-formats)
4. [http://www.openmicroscopy.org/site/support/faq](http://www.openmicroscopy.org/site/support/faq)
5. [http://www.openmicroscopy.org/site/community/mailing-lists](http://www.openmicroscopy.org/site/community/mailing-lists)
Please be patient - it may be a few days until you receive a response, but we reply to every email inquiry we receive.

3.2 Troubleshooting

This page is aimed at anyone who is responsible for supporting Bio-Formats, but may also be useful for advanced users looking to troubleshoot their own problems. Eventually, it might be best to move some of this to the FAQ or other documentation.

3.2.1 General tips

• Make sure to read the FAQ, particularly the “File Formats”, “Bio-Formats”, and “OME-XML & OME-TIFF” sections.
• If this page doesn’t help, it is worth quickly checking the following places where questions are commonly asked and/or bugs are reported:
  – OME Trac
  – Fiji Bugzilla (for ImageJ/Fiji issues)
  – ome-devel mailing list (searchable using google with ‘site:lists.openmicroscopy.org.uk’)
  – ome-users mailing list (searchable using google with ‘site:lists.openmicroscopy.org.uk’)
  – ImageJ mailing list (for ImageJ/Fiji issues)
• Make sure to ask for a specific error message or description of the unexpected behavior, if one is not provided (“it does not work” is obviously not adequate).
• “My (12, 14, 16)-bit images look all black when I open them” is a common issue. In ImageJ/Fiji, this is almost always fixable by checking the “Autoscale” option; with the command line tools, the “--autoscale -fast” options should work. The problem is typically that the pixel values are very, very small relative to the maximum possible pixel value (4095, 16383, and 65535, respectively), so when displayed the pixels are effectively black.
• If the file is very, very small (4096 bytes) and any exception is generated when reading the file, then make sure it is not a MacOS X resource fork. The ‘file’ command should tell you:

  $ file /path/to/suspicious-file
  suspicious-file: AppleDouble encoded Macintosh file

3.2.2 Tips for ImageJ/Fiji

• The Bio-Formats version being used can be found by selecting “Help > About Plugins > Bio-Formats Plugins”.
• “How do I make the options window go away?” is a common question. There are a few ways to do this:
  – To disable the options window only for files in a specific format, select “Plugins > Bio-Formats > Bio-Formats Plugins Configuration”, then pick the format from the list and make sure the “Windowless” option is checked.
  – To avoid the options window entirely, use the “Plugins > Bio-Formats > Bio-Formats Windowless Importer” menu item to import files.
  – Open files by calling the Bio-Formats importer plugin from a macro.
• A not uncommon cause of problems is that the user has multiple copies of loci_tools.jar in their ImageJ plugins folder, or has a copy of loci_tools.jar and a copy of formats-gpl.jar. It is often difficult to determine for sure that this is the problem - the only error message that pretty much guarantees it is a “NoSuchMethodException”. If the user maintains that they downloaded the latest version and whatever error message/odd behavior they are seeing looks like it was fixed already, then it is worth suggesting that they remove all copies of loci_tools.jar and download a fresh version.

---

9http://www.openmicroscopy.org/site/support/faq
10http://trac.openmicroscopy.org.uk/ome
11http://fiji.sc/cgi-bin/bugzilla/index.cgi
12http://lists.openmicroscopy.org.uk/pipermail/ome-devel
13http://lists.openmicroscopy.org.uk/pipermail/ome-users
14http://imagej.1557.n6.nabble.com/
3.2.3 Tips for command line tools

- When run with no arguments, all of the command line tools will print information on usage.
- When run with the `-version` argument, `showinfo` and `bfconvert` will display the version of Bio-Formats that is being used (version number, build date, and Git commit reference).

3.2.4 Tips by format

3I/Olympus Slidebook (.sld)

- Slidebook support is generally not great, despite a lot of effort. This is the one format for which it is recommended to just export to OME-TIFF from the acquisition software and work with the exported files. Happily, there is free software from 3I which can do the export post-acquisition: https://www.slidebook.com/reader.php

DICOM

- Health care or institutional regulations often prevent users from sending problematic files, so often we have to solve the problem blind. In these cases, it is important to get the exact error message, and inform the user that fixing the problem may be an iterative process (i.e. they might have to try a couple of trunk builds before we can finally fix the problem).

ZVI

- If the ZVI reader plugin is installed in ImageJ/Fiji, then it will be used instead of Bio-Formats to read ZVI files. To check if this is the cause of the problem, make sure that the file opens correctly using “Plugins > Bio-Formats > Bio-Formats Importer”; if that works, then just remove ZVI_Reader.class from the plugins folder.
CHAPTER
FOUR

BIO-FORMATS VERSIONS

Bio-Formats is updated whenever a new version of OMERO1 is released. The version number is three numbers separated by dots; e.g., 4.0.0. See the version history for a list of major changes in each release.

4.1 Version history

4.1.1 5.0.4 (2014 September 3)

• Fix compile and runtime errors under Java 1.8
• Improvements to Nikon .nd2 metadata parsing
• Added support for PicoQuant .bin files (thanks to Ian Munro)

4.1.2 5.0.3 (2014 August 7)

• Many bug fixes for Nikon .nd2 files
• Several other bug fixes, including:
  – LZW image decoding
  – Stage position parsing for Zeiss CZI
  – Exposure time units for ScanR
  – Physical pixel size units for DICOM
  – NDPI and Zeiss LSM files larger than 4GB
  – Z and T dimensions for InCell 6000 plates
  – Export of RGB images in ImageJ
• Improved metadata saving in MATLAB functions

4.1.3 5.0.2 (2014 May 28)

• Many bug fixes for Zeiss .czi files
• Several other bug fixes, including:
  – Gatan .dm3 units and step count parsing
  – Impector .msr 5D image support
  – DICOM reading of nested tags
• Update native-lib-loader version (to 2.0.1)
• Updates and improvements to user documentation

1http://www.openmicroscopy.org/site/support/omero5/
4.1.4 5.0.1 (2014 Apr 7)

- Added image pyramid support for CellSens .vsi data
- Several bug fixes, including:
  - Woolz import into OMERO
  - Cellomics file name parsing (thanks to Lee Kamentsky)
  - Olympus FV1000 timestamp support (thanks to Lewis Kraft and Patrick Riley)
  - (A)PNG large image support
  - Zeiss .czi dimension detection for SPIM datasets
- Performance improvements for Becker & Hickl .sdt file reading (thanks to Ian Munro)
- Performance improvements to directory listing over NFS
- Update slf4j and logback versions (to 1.7.6 and 1.1.1 respectively)
- Update goodies-forms version (to 1.7.2)

4.1.5 5.0.0 (2014 Feb 25)

- New bundled ‘bioformats_package.jar’ for ImageJ
- Now uses logback as the slf4j binding by default
- Updated component names, .jar file names, and Maven artifact names
- Fixed support for Becker & Hickl .sdt files with multiple blocks
- Fixed tiling support for TIFF, Hamamatsu .ndpi, JPEG, and Zeiss .czi files
- Improved continuous integration testing
- Updated command line documentation

4.1.6 5.0.0-RC1 (2013 Dec 19)

- Updated Maven build system and launched new Artifactory repository (http://artifacts.openmicroscopy.org)
- Added support for:
  - Bio-Rad SCN
  - Yokogawa CellVoyager (thanks to Jean-Yves Tinevez)
  - LaVision Inspector
  - PCORAW
  - Woolz (thanks to Bill Hill)
- Added support for populating and parsing ModuloAlong{Z, C, T} annotations for FLIM/SPIM data
- Updated netCDF and slf4j version requirements - netCDF 4.3.19 and slf4j 1.7.2 are now required
- Updated and improved MATLAB users and developers documentation
- Many bug fixes including for Nikon ND2, Zeiss CZI, and CellWorX formats

4.1.7 5.0.0-beta1 (2013 June 20)

- Updated to 2013-06 OME-XML schema
- Improved the performance in tiled formats

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2http://www.openmicroscopy.org/site/support/ome-model/
• Added caching of Reader metadata using http://code.google.com/p/kryo/
• Added support for:
  – Aperio AFI
  – Inveon
  – MPI-BPC Inspector
• Many bug fixes, including:
  – Add ZEN 2012/Lightsheet support to Zeiss CZI
  – Improved testing of autogenerated code
  – Moved OME-XML specification into Bio-Formats repository

4.1.8 4.4.10 (2014 Jan 15)
• Bug fixes including CellWorx, Metamorph and Zeiss CZI
• Updates to MATLAB documentation

4.1.9 4.4.9 (2013 Oct 16)
• Many bug fixes including improvements to support for ND2 format
• Java 1.6 is now the minimum supported version; Java 1.5 is no longer supported

4.1.10 4.4.8 (2013 May 2)
• No changes - release to keep version numbers in sync with OMERO

4.1.11 4.4.7 (2013 April 25)
• Many bug fixes to improve support for more than 20 formats
• Improved export to multi-file datasets
• Now uses slf4j for logging rather than using log4j directly, enabling other logging implementations to be used, for example when Bio-Formats is used as a component in other software using a different logging system.

4.1.12 4.4.6 (2013 February 11)
• Many bug fixes
• Further documentation improvements

4.1.13 4.4.5 (2012 November 13)
• Restructured and improved documentation
• Many bug fixes, including:
  – File grouping in many multi-file formats
  – Maven build fixes
  – ITK plugin fixes
4.1.14 4.4.4 (2012 September 24)

• Many bug fixes

4.1.15 4.4.2 (2012 August 22)

• Security fix for OMERO plugins for ImageJ

4.1.16 4.4.1 (2012 July 20)

• Fix a bug that prevented BigTIFF files from being read
• Fix a bug that prevented PerkinElmer .flex files from importing into OMERO

4.1.17 4.4.0 (2012 July 13)

• Many, many bug fixes
• Added support for:
  – .nd2 files from Nikon Elements version 4
  – PerkinElmer Operetta data
  – MJPEG-compressed AVIs
  – MicroManager datasets with multiple positions
  – Zeiss CZI data
  – IMOD data

4.1.18 4.3.3 (2011 October 18)

• Many bug fixes, including:
  – Speed improvements to HCImage/SimplePCI and Zeiss ZVI files
  – Reduce memory required by Leica LIF reader
  – More accurately populate metadata for Prairie TIFF datasets
  – Various fixes to improve the security of the OMERO plugin for ImageJ
  – Better dimension detection for Bruker MRI datasets
  – Better thumbnail generation for histology (SVS, NDPI) datasets
  – Fix stage position parsing for Metamorph TIFF datasets
  – Correctly populate the channel name for PerkinElmer Flex files

4.1.19 4.3.2 (2011 September 15)

• Many bug fixes, including:
  – Better support for Volocity datasets that contain compressed data
  – More accurate parsing of ICS metadata
  – More accurate parsing of cellSens .vsi files
• Added support for a few new formats
  – .inr
  – Canon DNG
• Updated Zeiss LSM reader to parse application tags
• Various performance improvements, particularly for reading/writing TIFFs
• Updated OMERO ImageJ plugin to work with OMERO 4.3.x

4.1.20 4.3.1 (2011 July 8)

• Several bug fixes, including:
  – Fixes for multi-position Deltavision files
  – Fixes for MicroManager 1.4 data
  – Fixes for 12 and 14-bit JPEG-2000 data
  – Various fixes for reading Volocity .mvd2 datasets
• Added various options to the ‘showinf’ and ‘bfconvert’ command line tools
• Added better tests for OME-XML backwards compatibility
• Added the ability to roughly stitch tiles in a multi-position dataset

4.1.21 4.3.0 (2011 June 14)

• Many bug fixes, including:
  – Many fixes for reading and writing sub-images
  – Fixes for stage position parsing in the Zeiss formats
  – File type detection fixes
• Updated JPEG-2000 reading and writing support to be more flexible
• Added support for 9 new formats:
  – InCell 3000
  – Trestle
  – Hamamatsu .ndpi
  – Hamamatsu VMS
  – SPIDER
  – Volocity .mvd2
  – Olympus SIS TIFF
  – IMAGIC
  – cellSens VSI
• Updated to 2011-06 OME-XML schema
• Minor speed improvements in many formats
• Switched version control system from SVN to Git
• Moved all Trac tickets into the OME Trac: http://trac.openmicroscopy.org.uk
• Improvements to testing frameworks
• Added Maven build system as an alternative to the existing Ant build system
• Added pre-compiled C++ bindings to the download page

4.1.22 4.2.2 (2010 December 6)

• Several bug fixes, notably:
  – Metadata parsing fixes for Zeiss LSM, Metamorph STK, and FV1000
  – Prevented leaked file handles when exporting to TIFF/OME-TIFF
  – Fixed how BufferedImage are converted to byte arrays
• Proper support for OME-XML XML annotations
• Added support for SCANDO Medical .aim files
• Minor improvements to ImageJ plugins
• Added support for reading JPEG-compressed AVI files

4.1.23 4.2.1 (2010 November 12)

• Many, many bug fixes
• Added support for 7 new formats:
  – CellWorX .pnl
  – ECAT7
  – Varian FDF
  – Perkin Elmer Densitometer
  – FEI TIFF
  – Compix/SimplePCI TIFF
  – Nikon Elements TIFF
• Updated Zeiss LSM metadata parsing, with generous assistance from Zeiss, FMI, and MPI-CBG
• Lots of work to ensure that converted OME-XML validates
• Improved file stitching functionality; non-numerical file patterns and limited regular expression-style patterns are now supported

4.1.24 4.2.0 (2010 July 9)

• Fixed many, many bugs in all aspects of Bio-Formats
• Reworked ImageJ plugins to be more user- and developer-friendly
• Added many new unit tests
• Added support for approximately 25 new file formats, primarily in the SPM domain
• Rewrote underlying I/O infrastructure to be thread-safe and based on Java NIO
• Rewrote OME-XML parsing/generation layer; OME-XML 2010-06 is now supported
• Improved support for exporting large images
• Improved support for exporting to multiple files
• Updated logging infrastructure to use slf4j and log4j
4.1.25 4.1.1 (2009 December 3)

- Fixed many bugs in popular file format readers

4.1 (2009 October 21):

- Fixed many bugs in most file format readers
- Significantly improved confocal and HCS metadata parsing
- Improved C++ bindings
- Eliminated references to Java AWT classes in core Bio-Formats packages
- Added support for reading Flex datasets from multiple servers
- Improved OME-XML generation; generated OME-XML is now valid
- Added support for Olympus ScanR data
- Added OSGi information to JARs
- Added support for Amira Mesh files
- Added support for LI-FLIM files
- Added more informative exceptions
- Added support for various types of ICS lifetime data
- Added support for Nikon EZ-C1 TIFFs
- Added support for Maia Scientific MIAS data

4.1.26 4.0.1 (2009 June 1)

- Lots of bug fixes in most format readers and writers
- Added support for Analyze 7.1 files
- Added support for Nifti files
- Added support for Cellomics .c01 files
- Refactored ImageJ plugins
- Bio-Formats, the common package, and the ImageJ plugins now require Java 1.5
- Changed license from GPLv3 or later to GPLv2 or later
- Updated Olympus FV1000, Zeiss LSM, Zeiss ZVI and Nikon ND2 readers to parse ROI data
- Added option to ImageJ plugin for displaying ROIs parsed from the chosen dataset
- Fixed BufferedImage construction for signed data and unsigned int data

4.1.27 4.0.0 (2009 March 3)

- Improved OME data model population for Olympus FV1000, Nikon ND2, Metamorph STK, Leica LEI, Leica LIF, InCell 1000 and MicroManager
- Added TestNG tests for format writers
- Added option to ImageJ plugin to specify custom colors when customizing channels
- Added ability to upgrade the ImageJ plugin from within ImageJ
- Fixed bugs in Nikon ND2, Leica LIF, BioRad PIC, TIFF, PSD, and OME-TIFF
- Fixed bugs in Data Browser and Exporter plugins
- Added support for Axon Raw Format (ARF), courtesy of Johannes Schindelin
• Added preliminary support for IPLab-Mac file format

4.1.28 2008 December 29

• Improved metadata support for Deltavision, Zeiss LSM, MicroManager, and Leica LEI
• Restructured code base/build system to be component-driven
• Added support for JPEG and JPEG-2000 codecs within TIFF, OME-TIFF and OME-XML
• Added support for 16-bit compressed Flex files
• Added support for writing JPEG-2000 files
• Added support for Minolta MRW format
• Added support for the 2008-09 release of OME-XML
• Removed dependency on JMagick
• Re-added caching support to data browser plugin
• Updated loci.formats.Codec API to be more user-friendly
• Expanded loci.formats.MetadataStore API to better represent the OME-XML model
• Improved support for Nikon NEF
• Improved support for TillVision files
• Improved ImageJ import options dialog
• Fixed bugs with Zeiss LSM files larger than 4 GB
• Fixed minor bugs in most readers
• Fixed bugs with exporting from an Image5D window
• Fixed several problems with virtual stacks in ImageJ

4.1.29 2008 August 30

• Fixed bugs in many file format readers
• Fixed several bugs with swapping dimensions
• Added support for Olympus CellIR/APL files
• Added support for MINC MRI files
• Added support for Aperio SVS files compressed with JPEG 2000
• Added support for writing OME-XML files
• Added support for writing APNG files
• Added faster LZW codec
• Added drag and drop support to ImageJ shortcut window
• Re-integrated caching into the data browser plugin

4.1.30 2008 July 1

• Fixed bugs in most file format readers
• Fixed bugs in OME and OMERO download functionality
• Fixed bugs in OME server-side import
• Improved metadata storage/retrieval when uploading to and downloading from the OME Perl server
• Improved Bio-Formats ImageJ macro extensions
• Major updates to MetadataStore API
• Updated OME-XML generation to use 2008-02 schema by default
• Addressed time and memory performance issues in many readers
• Changed license from LGPL to GPL
• Added support for the FEI file format
• Added support for uncompressed Hamamatsu Aquacosmos NAF files
• Added support for Animated PNG files
• Added several new options to Bio-Formats ImageJ plugin
• Added support for writing ICS files

4.1.31 2008 April 17

• Fixed bugs in Slidebook, ND2, FV1000 OIB/OIF, Perkin Elmer, TIFF, Prairie, Openlab, Zeiss LSM, MNG, Molecular Dynamics GEL, and OME-TIFF
• Fixed bugs in OME and OMERO download functionality
• Fixed bugs in OME server-side import
• Fixed bugs in Data Browser
• Added support for downloading from OMERO 2.3 servers
• Added configuration plugin
• Updates to MetadataStore API
• Updates to OME-XML generation - 2007-06 schema used by default
• Added support for Li-Cor L2D format
• Major updates to TestNG testing framework
• Added support for writing multi-series OME-TIFF files
• Added support for writing BigTIFF files

4.1.32 2008 Feb 12

• Fixed bugs in QuickTime, SimplePCI and DICOM
• Fixed a bug in channel splitting logic

4.1.33 2008 Feb 8

• Many critical bugfixes in format readers and ImageJ plugins
• Newly reborn Data Browser for 5D image visualization
  – some combinations of import options do not work yet

4.1.34 2008 Feb 1

• Fixed bugs in Zeiss LSM, Metamorph STK, FV1000 OIB/OIF, Leica LEI, TIFF, Zeiss ZVI, ICS, Prairie, Openlab LIFF, Gatan, DICOM, QuickTime
• Fixed bug in OME-TIFF writer
• Major changes to MetadataStore API
• Added support for JPEG-compressed TIFF files
• Added basic support for Aperio SVS files
  – JPEG2000 compression is still not supported
• Improved “crop on import” functionality
• Improvements to bfconvert and bfview
• Improved OME-XML population for several formats
• Added support for JPEG2000-compressed DICOM files
• EXIF data is now parsed from TIFF files

4.1.35 2007 Dec 28

• Fixed bugs in Leica LEI, Leica TCS, SDT, Leica LIF, Visitech, DICOM, Imaris 5.5 (HDF), and Slidebook readers
• Better parsing of comments in TIFF files exported from ImageJ
• Fixed problem with exporting 48-bit RGB data
• Added logic to read multi-series datasets spread across multiple files
• Improved channel merging in ImageJ - requires ImageJ 1.39l
• Support for hyperstacks and virtual stacks in ImageJ - requires ImageJ 1.39l
• Added API for reading directly from a byte array or InputStream
• Metadata key/value pairs are now stored in ImageJ’s “Info” property
• Improved OMERO download plugin - it is now much faster
• Added “open all series” option to ImageJ importer
• ND2 reader based on Nikon’s SDK now uses our own native bindings
• Fixed metadata saving bug in ImageJ
• Added sub-channel labels to ImageJ windows
• Major updates to 4D Data Browser
• Minor updates to automated testing suite

4.1.36 2007 Dec 1

• Updated OME plugin for ImageJ to support downloading from OMERO
• Fixed bug with floating point TIFFs
• Fixed bugs in Visitech, Zeiss LSM, Imaris 5.5 (HDF)
• Added alternate ND2 reader that uses Nikon’s native libraries
• Fixed calibration and series name settings in importer
• Added basic support for InCell 1000 datasets

4.1.37 2007 Nov 21

• Fixed bugs in ND2, Leica LIF, DICOM, Zeiss ZVI, Zeiss LSM, FV1000 OIB, FV1000 OIF, BMP, Evotec Flex, BioRad PIC, Slidebook, TIFF
• Added new ImageJ plugins to slice stacks and do “smart” RGB merging
• Added “windowless” importer plugin
  – uses import parameters from IJ_Prefs.txt, without prompting the user
• Improved stack slicing and colorizing logic in importer plugin
• Added support for DICOM files compressed with lossless JPEG
  – requires native libraries
• Fixed bugs with signed pixel data
• Added support for Imaris 5.5 (HDF) files
• Added 4 channel merging to importer plugin
• Added API methods for reading subimages
• Major updates to the 4D Data Browser

4.1.38 2007 Oct 17

• Critical OME-TIFF bugfixes
• Fixed bugs in Leica LIF, Zeiss ZVI, TIFF, DICOM, and AVI readers
• Added support for JPEG-compressed ZVI images
• Added support for BigTIFF
• Added importer plugin option to open each plane in a new window
• Added MS Video 1 codec for AVI

4.1.39 2007 Oct 1

• Added support for compressed DICOM images
• Added support for uncompressed LIM files
• Added support for Adobe Photoshop PSD files
• Fixed bugs in DICOM, OME-TIFF, Leica LIF, Zeiss ZVI, Visitech, PerkinElmer and Metamorph
• Improved indexed color support
• Addressed several efficiency issues
• Fixed how multiple series are handled in 4D data browser
• Added option to reorder stacks in importer plugin
• Added option to turn off autoscaling in importer plugin
• Additional metadata convenience methods

4.1.40 2007 Sept 11

• Major improvements to ND2 support; lossless compression now supported
• Support for indexed color images
• Added support for Simple-PCI .cxd files
• Command-line OME-XML validation
• Bugfixes in most readers, especially Zeiss ZVI, Metamorph, PerkinElmer and Leica LEI
• Initial version of Bio-Formats macro extensions for ImageJ
4.1.41 2007 Aug 1

- Added support for latest version of Leica LIF
- Fixed several issues with Leica LIF, Zeiss ZVI
- Better metadata mapping for Zeiss ZVI
- Added OME-TIFF writer
- Added MetadataRetrieve API for retrieving data from a MetadataStore
- Miscellaneous bugfixes

4.1.42 2007 July 16

- Fixed several issues with ImageJ plugins
- Better support for Improvision and Leica TCS TIFF files
- Minor improvements to Leica LIF, ICS, QuickTime and Zeiss ZVI readers
- Added searchable metadata window to ImageJ importer

4.1.43 2007 July 2

- Fixed issues with ND2, Openlab LIFF and Slidebook
- Added support for Visitech XYS
- Added composite stack support to ImageJ importer

4.1.44 2007 June 18

- Fixed issues with ICS, ND2, MicroManager, Leica LEI, and FV1000 OIF
- Added support for large (> 2 GB) ND2 files
- Added support for new version of ND2
- Minor enhancements to ImageJ importer
- Implemented more flexible logging
- Updated automated testing framework to use TestNG
- Added package for caching images produced by Bio-Formats

4.1.45 2007 June 6

- Fixed OME upload/download bugs
- Fixed issues with ND2, EPS, Leica LIF, and OIF
- Added support for Khoros XV
- Minor improvements to the importer

4.1.46 2007 May 24

- Better Slidebook support
- Added support for Quicktime RPZA
- Better Leica LIF metadata parsing
- Added support for BioRad PIC companion files
• Added support for bzip2-compressed files
• Improved ImageJ plugins
• Native support for FITS and PGM

4.1.47 2007 May 2

• Added support for NRRD
• Added support for Evotec Flex (requires LuraWave Java SDK with license code)
• Added support for gzip-compressed files
• Added support for compressed QuickTime headers
• Fixed QuickTime Motion JPEG-B support
• Fixed some memory issues (repeated small array allocations)
• Fixed issues reading large (> 2 GB) files
• Removed “ignore color table” logic, and replaced with Leica-specific solution
• Added status event reporting to readers
• Added API to toggle metadata collection
• Support for multiple dimensions rasterized into channels
• Deprecated reader and writer methods that accept the ‘id’ parameter
• Deprecated IFormatWriter.save in favor of saveImage and saveBytes
• Moved dimension swapping and min/max calculation logic to delegates
• Separate GUI logic into isolated loci.formats.gui package
• Miscellaneous bugfixes and tweaks in most readers and writers
• Many other bugfixes and improvements

4.1.48 2007 Mar 16

• Fixed calibration bugs in importer plugin
• Enhanced metadata support for additional formats
• Fixed LSM bug

4.1.49 2007 Mar 7

• Added support for Micro-Manager file format
• Fixed several bugs – Leica LIF, Leica LEI, ICS, ND2, and others
• Enhanced metadata support for several formats
• Load series preview thumbnails in the background
• Better implementation of openBytes(String, int, byte[]) for most readers
• Expanded unit testing framework

4.1.50 2007 Feb 28

• Better series preview thumbnails
• Fixed bugs with multi-channel Leica LEI
• Fixed bugs with “ignore color tables” option in ImageJ plugin
4.1.51 2007 Feb 26

- Many bugfixes: Leica LEI, ICS, FV1000 OIB, OME-XML and others
- Better metadata parsing for BioRad PIC files
- Enhanced API for calculating channel minimum and maximum values
- Expanded MetadataStore API to include more semantic types
- Added thumbnails to series chooser in ImageJ plugin
- Fixed plugins that upload and download from an OME server

4.1.52 2007 Feb 7

- Added plugin for downloading images from OME server
- Improved HTTP import functionality
- Added metadata filtering – unreadable metadata is no longer shown
- Better metadata table for multi-series datasets
- Added support for calibration information in Gatan DM3
- Eliminated need to install JAI Image I/O Tools to read ND2 files
- Fixed ZVI bugs: metadata truncation, and other problems
- Fixed bugs in Leica LIF: incorrect calibration, first series labeling
- Fixed memory bug in Zeiss LSM
- Many bugfixes: PerkinElmer, Deltavision, Leica LEI, LSM, ND2, and others
- IFormatReader.close(boolean) method to close files temporarily
- Replaced Compression utility class with extensible Compressor interface
- Improved testing framework to use .bioformats configuration files

4.1.53 2007 Jan 5

- Added support for Prairie TIFF
- Fixed bugs in Zeiss LSM, OIB, OIF, and ND2
- Improved API for writing files
- Added feature to read files over HTTP
- Fixed bugs in automated testing framework
- Miscellaneous bugfixes

4.1.54 2006 Dec 22

- Expanded ImageJ plugin to optionally use Image5D or View5D
- Improved support for ND2 and JPEG-2000 files
- Added automated testing framework
- Fixed bugs in Zeiss ZVI reader
- Miscellaneous bugfixes
4.1.55 2006 Nov 30

- Added support for ND2/JPEG-2000
- Added support for MRC
- Added support for MNG
- Improved support for floating-point images
- Fixed problem with 2-channel Leica LIF data
- Minor tweaks and bugfixes in many readers
- Improved file stitching logic
- Allow ImageJ plugin to be called from a macro

4.1.56 2006 Nov 2

- Bugfixes and improvements for Leica LIF, Zeiss LSM, OIF and OIB
- Colorize channels when they are split into separate windows
- Fixed a bug with 4-channel datasets

4.1.57 2006 Oct 31

- Added support for Imaris 5 files
- Added support for RGB ICS images

4.1.58 2006 Oct 30

- Added support for tiled TIFFs
- Fixed bugs in ICS reader
- Fixed importer plugin deadlock on some systems

4.1.59 2006 Oct 27

- Multi-series support for Slidebook
- Added support for Alicona AL3D
- Fixed plane ordering issue with FV1000 OIB
- Enhanced dimension detection in FV1000 OIF
- Added preliminary support for reading NEF images
- Added option to ignore color tables
- Fixed ImageJ GUI problems
- Fixed spatial calibration problem in ImageJ
- Fixed some lingering bugs in Zeiss ZVI support
- Fixed bugs in OME-XML reader
- Tweaked ICS floating-point logic
- Fixed memory leaks in all readers
- Better file stitching logic
### 4.1.60 2006 Oct 6

- Support for 3i SlideBook format (single series only for now)
- Support for 16-bit RGB palette TIFF
- Fixed bug preventing import of certain Metamorph STK files
- Fixed some bugs in PerkinElmer UltraView support
- Fixed some bugs in Leica LEI support
- Fixed a bug in Zeiss ZVI support
- Fixed bugs in Zeiss LSM support
- Fixed a bug causing slow identification of Leica datasets
- Fixed bugs in the channel merging logic
- Fixed memory leak for OIB format
- Better scaling of 48-bit RGB data to 24-bit RGB
- Fixed duplicate channels bug in “open each channel in a separate window”
- Fixed a bug preventing PICT import into ImageJ
- Better integration with HandleExtraFileTypes
- Better virtual stack support in Data Browser plugin
- Fixed bug in native QuickTime random access
- Keep aspect ratio for computed thumbnails
- Much faster file stitching logic

### 4.1.61 2006 Sep 27

- PerkinElmer: support for PE UltraView
- Openlab LIFF: support for Openlab v5
- Leica LEI: bug fixes, and support for multiple series
- ZVI, OIB, IPW: more robust handling of these formats (eliminated custom OLE parsing logic in favor of Apache POI)
- OIB: better metadata parsing (but maybe still not perfect?)
- LSM: fixed a bug preventing import of certain LSMs
- Metamorph STK: fixed a bug resulting in duplicate image planes
- User interface: use of system look & feel for file chooser dialog when available
- Better notification when JAR libraries are missing

### 4.1.62 2006 Sep 6

- Leica LIF: multiple distinct image series within a single file
- Zeiss ZVI: fixes and improvements contributed by Michel Boudinot
- Zeiss LSM: fixed bugs preventing the import of certain LSM files
- TIFF: fixed a bug preventing import of TIFFs created with Bio-Rad software

### 4.1.63 2006 Mar 31

- First release
Part II

User Information
The following sections explain the features of Bio-Formats and how to use it within ImageJ and Fiji:

# 5.1 ImageJ overview

**ImageJ** is an image processing and analysis application written in Java, widely used in the life sciences fields, with an extensible plugin infrastructure. You can use Bio-Formats as a plugin for ImageJ to read and write images in the formats it supports.

## 5.1.1 Installation

Download [bioformats_package.jar](http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/bioformats_package.jar) and drop it into your `ImageJ/plugins` folder. Next time you run ImageJ, a new Bio-Formats submenu with several plugins will appear in the Plugins menu, including the Bio-Formats Importer and Bio-Formats Exporter.

## 5.1.2 Usage

The Bio-Formats Importer plugin can display image stacks in several ways:

- In a standard ImageJ window (including as a hyperstack)
- Using the LOCI Data Browser plugin (included)
- With Joachim Walter’s Image5D plugin (if installed)
- With Rainer Heintzmann’s View5D plugin (if installed)

ImageJ v1.37 and later automatically (via `HandleExtraFileTypes`) calls the Bio-Formats logic, if installed, as needed when a file is opened within ImageJ, i.e. when using File → Open instead of explicitly choosing Plugins → Bio-Formats → Bio-Formats Importer from the menu.

For a more detailed description of each plugin, see the Bio-Formats page of the Fiji wiki.

## 5.1.3 Upgrading

To upgrade, just overwrite the old `bioformats_package.jar` with the latest one.

You may want to download the latest version of ImageJ first, to take advantage of new features and bug-fixes.

As of the 4.0.0 release, you can also upgrade the Bio-Formats plugin directly from ImageJ. Select Plugins → Bio-Formats → Update Bio-Formats Plugins from the ImageJ menu, then select which release you would like to use. You will then need to restart ImageJ to complete the upgrade process.

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2. [http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/bioformats_package.jar](http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/bioformats_package.jar)
3. [http://loci.wisc.edu/software/data-browser](http://loci.wisc.edu/software/data-browser)
4. [http://developer.imagej.net/plugins/image5d](http://developer.imagej.net/plugins/image5d)
5. [http://www.nanoimaging.de/View5D](http://www.nanoimaging.de/View5D)
5.1.4 Macros and plugins

Bio-Formats is fully scriptable in a macro, and callable from a plugin. To use in a macro, use the Macro Recorder to record a call to the Bio-Formats Importer with the desired options. You can also perform more targeted metadata queries using the Bio-Formats macro extensions.

Here are some example ImageJ macros and plugins that use Bio-Formats to get you started:

- **basicMetadata.txt** - A macro that uses the Bio-Formats macro extensions to print the chosen file’s basic dimensional parameters to the Log.
- **planeTimings.txt** - A macro that uses the Bio-Formats macro extensions to print the chosen file’s plane timings to the Log.
- **recursiveTiffConvert.txt** - A macro for recursively converting files to TIFF using Bio-Formats.
- **bfOpenAsHyperstack.txt** - This macro from Wayne Rasband opens a file as a hyperstack using only the Bio-Formats macro extensions (without calling the Bio-Formats Importer plugin).
- **zvi2HyperStack.txt** - This macro from Sebastien Huart reads in a ZVI file using Bio-Formats, synthesizes the LUT using emission wavelength metadata, and displays the result as a hyperstack.
- **dvSplitTimePoints.txt** - This macro from Sebastien Huart splits timepoints/channels on all DV files in a folder.
- **batchTiffConvert.txt** - This macro converts all files in a directory to TIFF using the Bio-Formats macro extensions.
- **Read_Image** - A simple plugin that demonstrates how to use Bio-Formats to read files into ImageJ.
- **Mass_Importer** - A simple plugin that demonstrates how to open all image files in a directory using Bio-Formats, grouping files with similar names to avoid opening the same dataset more than once.

5.2 Fiji overview

Fiji is an image processing package. It can be described as a distribution of ImageJ together with Java, Java 3D and a lot of plugins organized into a coherent menu structure. Fiji compares to ImageJ as Ubuntu compares to Linux.

Fiji works with Bio-Formats out of the box, because it comes bundled with the Bio-Formats ImageJ plugins.

For further details on Bio-Formats in Fiji, see the Bio-Formats Fiji wiki page.

5.2.1 Upgrading

Upgrading Bio-Formats within Fiji is as simple as invoking the “Update Fiji” command from the Help menu. By default, Fiji even automatically checks for updates every time it is launched, so you will always be notified when new versions of Bio-Formats (or any other bundled plugin) are available.

5.3 Bio-Formats features in ImageJ and Fiji

When you select Bio-Formats under the Plugin menu, you will see the following features:

- The **Bio-Formats Importer** is a plugin for loading images into ImageJ or Fiji. It can read over 100 proprietary life sciences formats and standardizes their acquisition metadata into the common OME data model. It will also extract and set basic metadata values such as spatial calibration if they are available in the file.

[8]https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/bio-formats-plugins/utils/macros/basicMetadata.txt


[18]http://fiji.sc/Plugins_Menu


5.2. Fiji overview
• The **Bio-Formats Exporter** is a plugin for exporting data to disk. It can save to the open OME-TIFF\(^{21}\) file format, as well as several movie formats (e.g. QuickTime, AVI) and graphics formats (e.g. PNG, JPEG).

• The **Bio-Formats Remote Importer** is a plugin for importing data from a remote URL. It is likely to be less robust than working with files on disk, so we recommend downloading your data to disk and using the regular Bio-Formats Importer whenever possible.

• The **Bio-Formats Windowless Importer** is a version of the Bio-Formats Importer plugin that runs with the last used settings to avoid any additional dialogs beyond the file chooser. If you always use the same import settings, you may wish to use the windowless importer to save time (Learn more [here](#)).

• The **Bio-Formats Macro Extensions** plugin prints out the set of commands that can be used to create macro extensions. The commands and the instructions for using them are printed to the ImageJ log window.

• The **Stack Slicer** plugin is a helper plugin used by the Bio-Formats Importer. It can also be used to split a stack across channels, focal planes or time points.

• The **Bio-Formats Plugins Configuration** dialog is a useful way to configure the behavior of each file format. The Formats tab lists supported file formats and toggles each format on or off, which is useful if your file is detected as the wrong format. It also toggles whether each format bypasses the importer options dialog through the “Windowless” checkbox. You can also configure any specific option for each format. The Libraries tab provides a list of available helper libraries used by Bio-Formats.

• The **Bio-Formats Plugins Shortcut Window** opens a small window with a quick-launch button for each plugin. Dragging and dropping files onto the shortcut window opens them quickly using the **Bio-Formats Importer** plugin.

• The **Update Bio-Formats Plugins** command will check for Bio-Formats Plugins updates. We recommend you update to the latest build as soon as you think you may have discovered a bug.

### 5.4 Installing Bio-Formats in ImageJ

**Note:** Since FIJI is essentially ImageJ with plugins like Bio-Formats already built in, people who install Fiji can skip this section. If you are also using the OMERO plugin for ImageJ, you may find the set-up guide on the new [user help site](http://help.openmicroscopy.org/imagej.html) useful for getting you started with both plugins at the same time.

Once you download\(^{23}\) and install ImageJ, you can install the Bio-Formats plugin by going to the Bio-Formats [download page](http://downloads.openmicroscopy.org/latest/bio-formats5.0/). For most end-users, we recommend downloading the **bioformats_package.jar** complete bundle.

However, you must decide which version of it you want to install. There are three primary versions of Bio-Formats: the latest builds, the daily builds, and the release versions. Which version you should install depends on your needs:

• The **latest build** is automatically updated every time any change is made to the source code on the main “dev_5.0” branch in Git, Bio-Formats’ software version control system. This build has the latest bug fixes, but it is not well tested and may have also introduced new bugs.

• The **daily build** is a compilation of that day’s changes that occurs daily around midnight. It is not any better tested than the latest build; but if you download it multiple times in a day, you can be sure you will get the same version each time.

• The **release** is thoroughly tested and has documentation to match. The list of supported formats on the Bio-Formats site corresponds to the most recent release. We do not add new formats to the list until a release containing support for that format has been completed. The release is less likely to contain bugs.

The release version is also more useful to programmers because they can link their software to a known, fixed version of Bio-Formats. Bio-Formats’ behavior will not be changing “out from under them” as they continue developing their own programs.

**Note:** There are currently two release version of Bio-Formats as we are maintaining support for the 4.4.x series while only actively developing the new 5.x series. Unless you are using Bio-Formats with the OMERO ImageJ plugin and an OMERO 4.4.x server, we recommend you use Bio-Formats 5. A new 4.4.x version will only be released if a major bug fix is required.

\(^{21}\)http://www.openmicroscopy.org/site/support/ome-model/ome-tiff

\(^{22}\)http://help.openmicroscopy.org/imagej.html

\(^{23}\)http://rsbweb.nih.gov/ij/download.html

\(^{24}\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/
We often **recommend that most people simply use the latest build** for two reasons. First, it may contain bug-fixes or new features you want anyway; secondly, you will have to reproduce any bug you encounter in Bio-Formats against the latest build before submitting a bug report. Rather than using the release until you find a bug that requires you to upgrade and reproduce it, why not just use the latest build to begin with?

Once you decide which version you need, go to the Bio-Formats download page\(^25\) and save the appropriate `bioformats_package.jar` to the Plugins directory within ImageJ.

![Figure 5.1: Plugin Directory for ImageJ: Where in ImageJ’s file structure you should place the file once you downloaded it.](image)

You may have to quit and restart ImageJ. Once you restart it, you will find Bio-Formats in the Bio-Formats option under the Plugins menu:

\(^25\) [http://downloads.openmicroscopy.org/latest/bio-formats5.0/](http://downloads.openmicroscopy.org/latest/bio-formats5.0/)
You are now ready to start using Bio-Formats.

5.5 Using Bio-Formats to load images into ImageJ

This section will explain how to use Bio-Formats to import files into ImageJ and how to use the settings on the Bio-Formats Import Options screen.

5.5.1 Opening files

There are three ways you can open a file using Bio-Formats:

1. Select the Bio-Formats Importer under the Bio-Formats plugins menu.
2. Drag and drop it onto the Bio-Formats Plugins Shortcut window.
3. Use the Open command in the File menu.

Unless you used the Bio-Formats Plugins Configuration dialog to open the file type windowlessly, you know you used Bio-Formats to open a file when you see a screen like this:

![Bio-Formats Import Options Screen](image)

If you used the File > Open command and did not see the Bio-Formats Import Options screen, ImageJ/Fiji probably used another plugin instead of Bio-Formats to open the file. If this happens and you want to open a file using Bio-Formats, use one of the other two methods instead.

5.5.2 Opening files windowlessly

When you open a file with Bio-Formats, the Import Options Screen automatically recalls the settings you last used to open a file with that specific format (e.g. JPG, TIF, LSM, etc.). If you always choose the same options whenever you open files in a specific file format, you can save yourself time by bypassing the Bio-Formats Import Options screen. You can accomplish this two ways:

1. You can select the **Bio-Formats Windowless Importer**, located in the Bio-Formats menu under ImageJ’s Plugin menu. When you select this option, Bio-Formats will import the file using the same settings you used the last time you imported a file with the same format.

2. If you invariably use the same settings when you open files in a specific format, you can always bypass the Import Options Screen by changing the settings in the **Bio-Formats Plugins Configuration** option, which is also located in the Bio-Formats menu under ImageJ’s Plugin menu.
Once you select this option, select the file format you are interested in from the list on the left side of the screen. Check both the Enabled and Windowless boxes. Once you do this, whenever you open a file using the Bio-Formats Windowless Importer, the Bio-Formats Importer, or the drag-and-drop method described in the previous section, the file will always open the same way using the last setting used.

Please note that if you want to change any of the import settings once you enable this windowless option, you will have to go back to the Bio-Formats Plugins Configuration screen, unselect the windowless option, open a file using the regular Bio-Formats Importer, select your settings, and re-select the windowless option.

5.5.3 Group files with similar names

One of the most important features of Bio-Formats is to combine multiple files from a dataset into one coherent, multi-dimensional image.

To demonstrate how to use the Group files with similar names feature, you can use the dub data set available under LOCI’s Sample Data page. You will notice that it is a large dataset: each of the 85 files shows the specimen at 33 optical sections along the z-plane at a specific time.

If you open just one file in ImageJ/Fiji using the Bio-Formats Importer, you will get an image incorporating three dimensions (x, y, z). However, if you select Group files with similar names from the Bio-Formats Import Options screen, you will be able to create a 4-D image (x, y, z, and t) incorporating the 85 files.

After clicking OK, you will see a screen like this:

![Screen showing list of files to be grouped]

The list of files to be grouped can be specified in one of three ways:

- **Axis 1 number of images**: 85
- **Axis 1 axis first image**: 1
- **Axis 1 axis increment**: 1

This screen allows you to select which files within the 85-file cluster to use to create that 4-D image. Some information will be pre-populated in the fields. Unless you want to change the settings in that field, there is no need to change or delete it. If you click OK at this point, you will load all 85 files.

However, you can specify which files you want to open by adjusting the “axis information”, the file “name contains”, or the “pattern” sections. Even though there are three options, you only need to need to make changes to one of them. Since Bio-Format’s precedence for processing data is from top to bottom, only the uppermost section that you made changes to will be used. If you change multiple boxes, any information you enter into lower boxes will be ignored.

To return to the example involving the dub data set, suppose you want to open the first image and only every fifth image afterwards (i.e. dub01, dub06, dub11 . . . dub81). This would give you 17 images. There are different ways to accomplish this:

You can use the Axis Settings only when your files are numbered in sequential order and you want to open only a subset of the files that have similar names. Since the dub data set is numbered sequentially, you can use this feature.

**Axis 1 number of images** refers to the total number of images you want to open. Since you want to view 17 images, enter 17. **Axis 1 axis first image** specifies which image in the set you want to be the first. Since you want to start with dub01, enter 1 in that box. You also want to view only every fifth image, so enter 5 in the **Axis 1 axis increment** box.

---

26http://loci.wisc.edu/sample-data/dub
27http://loci.wisc.edu/software/sample-data
The **File name contains** box should be used if all of the files that you want to open have common text. This is especially useful when the files are not numbered. For example, if you have “Image_Red.tif”, “Image_Green.tif”, and “Image_Blue.tif” you could enter “Image_” in the box to group them all.

To continue the example involving the dub dataset, you cannot use the **file name contains** box to open every fifth image. However, if you only wanted to open dub10 thorough dub19, you could enter “dub1” in the **file name contains** box.

The **pattern** box can be used to do either of the options listed above or much more. This box can accept a single file name like “dub01.pic”. It can also contain a pattern that use “<” and “>” to specify what numbers or text the file names contain.

There are three basic forms to the “< >” blocks:

- **Text enumeration** - “Image_<Red,Green,Blue>.tif” is the pattern for Image_Red.tif, Image_Green.tif, Image_Blue.tif. (Note that the order you in which you enter the file names is the order in which they will be loaded.)
- **Number range** - “dub<1-85>.pic” is the pattern for “dub1.pic”, “dub2.pic”, “dub3.pic” . . . “dub85.pic”.
- **Number range with step** - “dub<1-85:5>.pic” is the pattern for “dub1.pic”, “dub6.pic”, “dub11.pic”, “dub16.pic” . . . “dub85.pic”.

It can also accept a [Java regular expression](http://download.oracle.com/javase/1.5.0/docs/api/java/util/regex/Pattern.html).

### 5.5.4 Autoscale

**Autoscale** helps increase the brightness and contrast of an image by adjusting the range of light intensity within an image to match the range of possible display values. Note that Autoscale does not change your data. It just changes how it is displayed.

Each pixel in an image has a numerical value ascribed to it to describe its intensity. The bit depth—the number of possible values—depends on the number of bits used in the image. Eight bits, for example, gives 256 values to express intensity where 0 is completely black, 255 is completely white, and 1 through 254 display increasingly lighter shades of grey.

ImageJ can collect the intensity information about each pixel from an image or stack and create a histogram (you can see it by selecting Histogram under the Analyze menu). Here is the histogram of a one particular image:

![Histogram of 15test.ome](image)

Notice that the histogram heavily skews right. Even though there are 256 possible values, only 0 thorough 125 are being used.
Autoscale adjusts the image so the smallest and largest number in that image or stack’s histogram become the darkest and brightest settings. For this image, pixels with the intensity of 125 will be displayed in pure white. The other values will be adjusted too to help show contrast between values that were too insignificant to see before.

Here is one image Bio-Formats imported with and without using Autoscale:

![Image 1](image1.png)

![Image 2](image2.png)

Autoscale readjusts the image based on the highest value in the entire data set. This means if the highest value in your dataset is close to maximum display value, Autoscale’s adjusting may be undetectable to the eye.

ImageJ/Fiji also has its own tools for adjusting the image, which are available by selecting Brightness/Contrast, which is under the Adjust option in the Image menu.

### 5.6 Managing memory in ImageJ/Fiji using Bio-Formats

When dealing with a large stack of images, you may receive a warning like this:
This means the allotted memory is less than what Bio-Formats needs to load all the images. If you have a very large data set, you may have to:

- View your stack with Data Browser
- Crop the view area
- Open only a subset of images
- Use Virtual Stack
- Increase ImageJ/Fiji’s memory.

If your files contain JPEG or JPEG-2000 images, you may see this memory warning even if your file size is smaller than the amount of allocated memory. This is because compressed images like JPEG need to be decompressed into memory before being displayed and require more memory than their file size suggests. If you are having this issue, try utilizing one of the memory management tools below.

### 5.6.1 View your stack with Data Browser

Data Browser is another part of Bio-Formats that enables users to view large 3, 4, or 5-D datasets by caching a subset of all the images available. This enables users to view a stack that is bigger than the computer's memory.

You can select Data Browser as an option for View stack with, the leftmost, uppermost option in the Bio-Formats Import Options screen.
Note that when you use Data Browser, other features like cropping and specifying range are not available. You can, however, adjust the size of the image cache in the Data Browser after you open the files. You can read more about it on LOCI’s Data Browser page.29

5.6.2 Cropping the view area

*Crop on Import* is useful if your images are very large and you are only interested in one specific section of the stack you are importing. If you select this feature, you will see a screen where you can enter the height and width (in pixels) of the part of image you want to see. Note that these measurements are from the top left corner of the image.

5.6.3 Opening only a subset of images

The *Specify Range for Each Series* option is useful for viewing a portion of a data set where all the plane images are encapsulated into one file (e.g. the Zeiss LSM format). If your file has a large quantity of images, you can specify which channels, Z-planes, and times you want to load.

5.6.4 Use Virtual Stack

*Virtual Stack* conserves memory by not loading specific images until necessary. Note that unlike Data Browser, Virtual Stack does not contain a buffer and may produce choppy animations.

5.6.5 Increasing ImageJ/Fiji’s memory

Finally, you can also increase the amount of the computer memory devoted to ImageJ/Fiji by selecting *Memory & Threads* under the *Edit* menu.

29http://loci.wisc.edu/software/data-browser
Generally, allocating more than 75% of the computer’s total memory will cause ImageJ/Fiji to become slow and unstable.

**Please note** that unlike the other three features, ImageJ/Fiji itself provides this feature and not Bio-Formats. You can find out more about this feature by looking at ImageJ’s documentation[^30].

[^30]: http://rsbweb.nih.gov/ij/docs/menus/edit.html#options
The Bio-Formats Command line tools (bftools.zip) provide a complete package for carrying out a variety of tasks:

## 6.1 Command line tools introduction

There are several scripts for using Bio-Formats on the command line.

### 6.1.1 Installation

Download `bftools.zip`, unzip it into a new folder.

**Note:** As of Bio-Formats 5.0.0, this zip now contains the bundled jar and you no longer need to download `loci_tools.jar` or the new `bioformats_package.jar` separately.

The zip file contains both Unix scripts and Windows batch files.

### 6.1.2 Tools available

Currently available tools include:

- **showinf**: Prints information about a given image file to the console, and displays the image itself in the Bio-Formats image viewer (see Displaying images and metadata for more information).
- **ijview**: Displays the given image file in ImageJ using the Bio-Formats Importer plugin.
- **bfconvert**: Converts an image file from one format to another. Bio-Formats must support writing to the output file (see Converting a file to different format for more information).
- **formatlist**: Displays a list of supported file formats in HTML, plaintext or XML.
- **xmlindent**: A simple XML prettifier similar to `xmllint --format` but more robust in that it attempts to produce output regardless of syntax errors in the XML.
- **xmlvalid**: A command-line XML validation tool, useful for checking an OME-XML document for compliance with the OME-XML schema.
- **tiffcomment**: Dumps the comment from the given TIFF file’s first IFD entry; useful for examining the OME-XML block in an OME-TIFF file (also see Editing XML in an OME-TIFF).

Some of these tools also work in combination, for example Validating XML in an OME-TIFF uses both `tiffcomment` and `xmlvalid`.

Running any of these commands without any arguments will print usage information to help you.

---

1. [http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/bftools.zip](http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/bftools.zip)
6.1.3 Using the tools directly from source

Firstly, obtain a copy of the sources and build them (see *Obtaining and building Bio-Formats*). You can configure the scripts to use your source tree instead of `bioformats_package.jar` in the same directory by following these steps:

1. Point your CLASSPATH to the checked-out directory and the JAR files in the `jar` folder.
   - E.g. on Windows with Java 1.6 or later, if you have checked out the source at C:\code\bio-formats, set your CLASSPATH environment variable to the value `C:\code\bio-formats\jar\*;C:\code\bio-formats`.
   - You can access the environment variable configuration area by right-clicking on My Computer, choosing Properties, Advanced tab, Environment Variables button.

2. Compile the source with `ant compile`.

3. Set the `BF_DEVEL` environment variable to any value (the variable just needs to be defined).

6.1.4 Version checker

If you run bftools outside of the OMERO environment, you may encounter an issue with the automatic version checker causing a tool to crash when trying to connect to upgrade.openmicroscopy.org.uk. The error message will look something like this:

```
Failed to compare version numbers
java.io.IOException: Server returned HTTP response code: 400 for URL: http://upgrade.openmicroscopy.org.uk?version=4.4.8;os.name=Linux;os.version=2.6.32-358.6.2.el6.x86_64;os.arch=amd64;java.runtime.version=1.6.0_24-b24;java.vm.vendor=Sun+Microsystems+Inc.;bioformats.caller= Bio-Formats+utilities
```

To avoid this issue, call the tool with the `-no-upgrade` parameter.

6.2 Displaying images and metadata

The `showinf` command line tool can be used to show the images and metadata contained in a file.

If no options are specified, `showinf` displays a summary of available options.

To simply display images:

```
showinf /path/to/file
```

All of the images in the first ‘series’ (or 5 dimensional stack) will be opened and displayed in a simple image viewer. The number of series, image dimensions, and other basic metadata will be printed to the console.

To display a different series, for example the second one:

```
showinf -series 1 /path/to/file
```

Note that series numbers begin with 0.

To display the OME-XML metadata for a file on the console:

```
showinf -omexml /path/to/file
```

Image reading can be suppressed if only the metadata is needed:

```
showinf -nopix /path/to/file
```
A subset of images can also be opened instead of the entire stack, by specifying the start and end plane indices (inclusive):

```
showinf -range 0 0 /path/to/file
```

That opens only the first image in first series in the file.

For very large images, it may also be useful to open a small tile from the image instead of reading everything into memory. To open the upper-left-most 512x512 tile from the images:

```
showinf -crop 0,0,512,512 /path/to/file
```

The parameter to `--crop` is of the format `x, y, width, height`. The `(x, y)` coordinate `(0, 0)` is the upper-left corner of the image; `x + width` must be less than or equal to the image width and `y + height` must be less than or equal to the image height.

By default, `showinf` will check for a new version of Bio-Formats. This can take several seconds (especially on a slow internet connection); to save time, the update check can be disabled:

```
showinf --no-update /path/to/file
```

Similarly, if OME-XML is displayed then it will automatically be validated. On slow or missing internet connections, this can take some time, and so can be disabled:

```
showinf --novalid /path/to/file
```

### 6.3 Converting a file to different format

The `bfconvert` command line tool can be used to convert files between supported formats.

`bfconvert` with no options displays a summary of available options.

To convert a file to single output file (e.g. TIFF):

```
bfconvert /path/to/input output.tiff
```

The output file format is determined by the extension of the output file, e.g. `.tiff` for TIFF files, `.ome.tiff` for OME-TIFF, `.png` for PNG.

All images in the input file are converted by default. To convert only one series:

```
bfconvert -series 0 /path/to/input output-first-series.tiff
```

To convert only one timepoint:

```
bfconvert -timepoint 0 /path/to/input output-first-timepoint.tiff
```

To convert only one channel:

```
bfconvert -channel 0 /path/to/input output-first-channel.tiff
```

To convert only one Z section:
bfconvert -z 0 /path/to/input output-first-z.tif

To convert images between certain indices (inclusive):

bfconvert -range 0 2 /path/to/input output-first-3-images.tif

Images can also be written to multiple files by specifying a pattern string in the output file. For example, to write one series, timepoint, channel, and Z section per file:

bfconvert /path/to/input output_series_%s_Z%z_C%c_T%t.tif

%s is the series index, %z is the Z section index, %c is the channel index, and %t is the timepoint index (all indices begin at 0).

By default, all images will be written uncompressed. Supported compression modes vary based upon the output format, but when multiple modes are available the compression can be changed using the -compression option. For example, to use LZW compression in a TIFF file:

bfconvert -compression LZW /path/to/input output-lzw.tif

### 6.4 Validating XML in an OME-TIFF

The XML stored in an OME-TIFF file can be validated using the command line tools. Both the tiffcomment and xmlvalid commands are used; tiffcomment extracts the XML from the file and xmlvalid validates the XML and prints any errors to the console.

For example:

```
tiffcomment /path/to/file.ome.tif | xmlvalid -
```

will perform the extraction and validation all at once.

Typical successful output is:

```
[~/Work/bftools]$ ./xmlvalid sample.ome
Parsing schema path
http://www.openmicroscopy.org/Schemas/OME/2010-06/ome.xsd
Validating sample.ome
No validation errors found.
[~/Work/bftools]$
```

If any errors are found they are reported. When correcting errors it is usually best to work from the top of the file as errors higher up can cause extra errors further down. In this example the output shows 3 errors but there are only 2 mistakes in the file:

```
[~/Work/bftools]$ ./xmlvalid broken.ome
Parsing schema path
http://www.openmicroscopy.org/Schemas/OME/2010-06/ome.xsd
Validating broken.ome
  cvc-complex-type.4: Attribute ‘SizeY’ must appear on element ‘Pixels’.
  cvc-enumeration-valid: Value ‘Non Zero’ is not facet-valid with respect to enumeration ‘[EvenOdd, NonZero]’. It must be a value from the enumeration.
  cvc-attribute.3: The value ‘Non Zero’ of attribute ‘FillRule’ on element ‘ROI:Shape’ is not valid with respect to its type, ‘null’.
```

### 6.4. Validating XML in an OME-TIFF
Error validating document: 3 errors found
[~/Work/bftools]$

If the XML is found to have validation errors, the `tiffcomment` command can be used to overwrite the XML in the OME-TIFF file with corrected XML. The XML can be displayed in an editor window:

```
tiffcomment -edit /path/to/file.ome.tif
```

or the new XML can be read from a file:

```
tiffcomment -set new-comment.xml /path/to/file.ome.tif
```

### 6.5 Editing XML in an OME-TIFF

To edit the XML in an OME-TIFF file you can use `tiffcomment`, one of the Bio-Formats tools.

To use the built in editor run:

```
tiffcomment -edit sample.ome.tif
```

To extract or view the XML run:

```
tiffcomment sample.ome.tif
```

To inject replacement XML into a file run:

```
tiffcomment -set 'newmetadata.xml' sample.ome.tif
```
OMERO 5 uses Bio-Formats to read original files from over 130 file formats. Please refer to the OMERO documentation\(^1\) for further information.

\(^1\)http://www.openmicroscopy.org/site/support/omero5/
8.1 BISQUE

The BISQUE\(^1\) (Bio-Image Semantic Query User Environment) Database, developed at the Center for Bio-Image Informatics at UCSB, was developed for the exchange and exploration of biological images. The Bisque system supports several areas useful for imaging researchers from image capture to image analysis and querying. The bisque system is centered around a database of images and metadata. Search and comparison of datasets by image data and content is supported. Novel semantic analyses are integrated into the system allowing high level semantic queries and comparison of image content.

Bisque integrates with Bio-Formats by calling the \textit{showinf} command line tool.

8.2 OME Server

OME\(^2\) is a set of software that interacts with a database to manage images, image metadata, image analysis and analysis results. The OME system is capable of leveraging Bio-Formats to import files.

Please note - the OME server is no longer maintained and has now been superseded by the OMERO server\(^3\). Support for the OME server has been entirely removed in the 5.0.0 version of Bio-Formats; the following instructions can still be used with the 4.4.x versions.

8.2.1 Installation

For OME Perl v2.6.1\(^4\) and later, the command line installer automatically downloads the latest \texttt{loci_tools.jar} and places it in the proper location. This location is configurable, but is \texttt{/OME/java/loci_tools.jar} by default.

For a list of what was recognized for a particular import into the OME server, go to the Image details page in the web interface, and click the “Image import” link in the upper right hand box.

Bio-Formats is capable of parsing original metadata for supported formats, and standardizes what it can into the OME data model. For the rest, it expresses the metadata in OME terms as key/value pairs using an OriginalMetadata custom semantic type. However, this latter method of metadata representation is of limited utility, as it is not a full conversion into the OME data model.

Bio-Formats is enabled in OME v2.6.1 for all formats except:

- OME-TIFF
- Metamorph HTD
- Deltavision DV
- Metamorph STK
- Bio-Rad PIC
- Zeiss LSM
- TIFF

\(^1\)http://www.bioimage.ucsb.edu/bisque
\(^2\)http://openmicroscopy.org/site/support/legacy/ome-server
\(^3\)http://www.openmicroscopy.org/site/support/omero5/
\(^4\)http://downloads.openmicroscopy.org/ome/2.6.1/
The above formats have their own Perl importers that override Bio-Formats, meaning that Bio-Formats is not used to process them by default. However, you can override this behavior (except for Metamorph HTD, which Bio-Formats does not support) by editing an OME database configuration value:

```sql
% psql ome
```

To see the current file format reader list:

```sql
ome=# select value from configuration where name='import_formats';
```

```sql
value
------------------------------------------------------------------------------
["OME::ImportEngine::OMETIFFreader","OME::ImportEngine::MetamorphHTDFormat",
 "OME::ImportEngine::DVreader","OME::ImportEngine::STKreader",
 "OME::ImportEngine::BioradReader","OME::ImportEngine::LSMreader",
 "OME::ImportEngine::TIFFreader","OME::ImportEngine::BMPreader",
 "OME::ImportEngine::DICOMreader","OME::ImportEngine::XMLreader",
 "OME::ImportEngine::BioFormats"]
```

To remove extraneous readers from the list:

```sql
ome=# update configuration set value='["OME::ImportEngine::MetamorphHTDFormat",
 ","OME::ImportEngine::XMLreader","OME::ImportEngine::BioFormats"]' where
 name='import_formats';
```

```sql
UPDATE 1
```

```sql
ome=# select value from configuration where name='import_formats';
```

```sql
value
------------------------------------------------------------------------------
["OME::ImportEngine::MetamorphHTDFormat","OME::ImportEngine::XMLreader",
 "OME::ImportEngine::BioFormats"]
```

To reset things back to how they were:

```sql
ome=# update configuration set value='["OME::ImportEngine::OMETIFFreader",
 ","OME::ImportEngine::MetamorphHTDFormat",
 ","OME::ImportEngine::DVreader",
 ","OME::ImportEngine::STKreader",
 ","OME::ImportEngine::BioradReader",
 ","OME::ImportEngine::LSMreader",
 ","OME::ImportEngine::TIFFreader",
 ","OME::ImportEngine::BMPreader",
 ","OME::ImportEngine::DICOMreader",
 ","OME::ImportEngine::XMLreader",
 "OME::ImportEngine::BioFormats"]' where
 name='import_formats';
```

Lastly, please note that Li-Cor L2D files cannot be imported into an OME server (see this [Trac ticket](http://dev.loci.wisc.edu/trac/software/ticket/266) for details). Since the OME perl server has been discontinued, we have no plans to fix this limitation.

### 8.2.2 Upgrading

You can upgrade your OME server installation to take advantage of a new Bio-Formats release by overwriting the old `loci_tools.jar` with the new one.

5http://dev.loci.wisc.edu/trac/software/ticket/266

6http://downloads.openmicroscopy.org/latest/bio-formats5.0/
8.2.3 Source Code

The source code for the Bio-Formats integration with OME server spans three languages, using piped system calls in both directions to communicate, with imported pixels written to OMEIS pixels files. The relevant source files are:

- OmeisImporter.java\textsuperscript{7} – omebf Java command line tool
- BioFormats.pm\textsuperscript{8} – Perl module for OME Bio-Formats importer
- omeis.c\textsuperscript{9} – OMEIS C functions for Bio-Formats (search for “bioformats” case insensitively to find relevant sections)

\textsuperscript{7}http://github.com/openmicroscopy/bioformats/tree/v4.4.10/components/scifio/src/loci/formats/ome/OmeisImporter.java
\textsuperscript{8}http://downloads.openmicroscopy.org/ome/code/BioFormats.pm
\textsuperscript{9}http://downloads.openmicroscopy.org/ome/code/omeis.c
9.1 FARSIGHT

FARSIGHT\(^1\) is a collection of modules for image analysis created by LOCI’s collaborators at the University of Houston\(^2\). These open source modules are built on the ITK library and thus can take advantage of ITK’s support for Bio-Formats to process otherwise unsupported image formats.

The principal FARSIGHT module that benefits from Bio-Formats is the Nucleus Editor\(^3\), though in principle any FARSIGHT-based code that reads image formats via the standard ITK mechanism will be able to leverage Bio-Formats.

See also:
FARSIGHT Downloads page\(^4\)
FARSIGHT HowToBuild tutorial\(^5\)

9.2 i3dcore

i3dcore\(^6\), also known as the CBIA 3D image representation library, is a 3D image processing library developed at the Centre for Biomedical Image Analysis\(^7\). Together with i3dalgo\(^8\) and i4dcore\(^9\), i3dcore forms a continuously developed templated cross-platform C++ suite of libraries for multidimensional image processing and analysis.

i3dcore is capable of reading images with Bio-Formats using Java for C++\(^10\) (java4cpp).

See also:
Download i3dcore\(^11\)
CBIA Software Development\(^12\)

9.3 ImgLib

ImgLib\(^13\) is a multidimensional image processing library. It provides a general mechanism for writing image analysis algorithms, without writing case logic for bit depth\(^14\), or worrying about the source of the pixel data (arrays in memory, files on disk, etc.).

---
\(^1\)http://www.farsight-toolkit.org/
\(^2\)http://www.uh.edu/
\(^3\)http://www.farsight-toolkit.org/wiki/NucleusEditor
\(^4\)http://www.farsight-toolkit.org/wiki/Special:FarsightDownloads
\(^5\)http://www.farsight-toolkit.org/wiki/FARSIGHT_HowToBuild
\(^6\)http://cbia.fi.muni.cz/user_dirs/i3dlib_doc/i3dcore/index.html
\(^7\)http://cbia.fi.muni.cz/software-development.html
\(^8\)http://cbia.fi.muni.cz/user_dirs/i3dlib_doc/i3dalgo/index.html
\(^9\)http://cbia.fi.muni.cz/user_dirs/of_doc/libi4d.html
\(^10\)http://java4cpp.kapott.org/
\(^11\)http://cbia.fi.muni.cz/user_dirs/i3dlib_doc/i3dcore/index.html#download
\(^12\)http://cbia.fi.muni.cz/software-development.html
\(^13\)http://imglib2.net/
\(^14\)http://en.wikipedia.org/wiki/Color_depth
The SCIFIO\(^{15}\) project provides an ImgOpener\(^{16}\) utility class for reading data into ImgLib2 data structures using Bio-Formats.

### 9.4 ITK

The Insight Toolkit\(^{17}\) (ITK) is an open-source, cross-platform system that provides developers with an extensive suite of software tools for image analysis. Developed through extreme programming methodologies, ITK employs leading-edge algorithms for registering and segmenting multidimensional data.

ITK provides an ImageIO plug-in structure that works via discovery through a dependency injection scheme. This allows a program built on ITK to load plug-ins for reading and writing different image types without actually linking to the ImageIO libraries required for those types. Such encapsulation automatically grants two major boons: firstly, programs can be easily extended just by virtue of using ITK (developers do not have to specifically accommodate or anticipate what plug-ins may be used). Secondly, the architecture provides a distribution method for open source software, like Bio-Formats, which have licenses that might otherwise exclude them from being used with other software suites.

The SCIFIO ImageIO\(^{18}\) plugin provides an ITK imageIO base that uses Bio-Formats to read and write supported life sciences file formats. This plugin allows any program built on ITK to read any of the image types supported by Bio-Formats.

### 9.5 Qu for MATLAB

Qu for MATLAB\(^{19}\) is a MATLAB toolbox for the visualization and analysis of N-dimensional datasets targeted to the field of biomedical imaging, developed by Aaron Ponti.

- Uses Bio-Formats to read files
- Open source software available under the Mozilla Public License

See also:

Qu for MATLAB download page\(^{20}\)

### 9.6 Subimager

Subimager\(^{21}\), the SUBprocess IMAGE servER, is an HTTP server that uses Bio-Formats as a back-end to serve .TIF images. Subimager is designed to be run as a subprocess of CellProfiler to provide CellProfiler with the capability to read and write a variety of image formats. It can be used as a stand-alone image server. It was developed by the Broad Institute\(^{22}\) to facilitate integration with their CellProfiler\(^{23}\) image analysis application.

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\(^{15}\)http://scif.io/

\(^{16}\)https://github.com/scifio/scifio/blob/master/src/main/java/io/scif/img/ImgOpener.java

\(^{17}\)http://itk.org/

\(^{18}\)https://github.com/scifio/scifio-imageio

\(^{19}\)http://www.scs2.net/home/index.php?option=com_content&view=article&id=46%3Aqu-for-matlab&catid=34%3Aqu&Itemid=55

\(^{20}\)http://www.scs2.net/home/index.php?option=com_content&view=article&id=46%3Aqu-for-matlab&catid=34%3Aqu&Itemid=55&limitstart=3

\(^{21}\)https://github.com/CellProfiler/subimager

\(^{22}\)http://www.broadinstitute.org/

\(^{23}\)http://www.cellprofiler.org/
CHAPTER TEN

NUMERICAL DATA PROCESSING APPLICATIONS

10.1 IDL

IDL\(^1\) (Interactive Data Language) is a popular data visualization and analysis platform used for interactive processing of large amounts of data including images.

IDL possesses the ability to interact with Java applications via its IDL-Java bridge. Karsten Rodenacker has written a script that uses Bio-Formats to read in image files to IDL.

10.1.1 Installation

Download the \texttt{ij\_read\_bio\_formats.pro}\(^2\) script from Karsten Rodenacker’s IDL goodies (\textit{?})\(^3\) web site. See the comments at the top of the script for installation instructions and caveats.

10.1.2 Upgrading

To use a newer version of Bio-Formats, overwrite the requisite JAR files with the newer version\(^4\) and restart IDL.

10.2 KNIME

KNIME\(^5\) (Konstanz Information Miner) is a user-friendly and comprehensive open-source data integration, processing, analysis, and exploration platform. KNIME supports image import using Bio-Formats using the KNIME Image Processing\(^6\) (a.k.a. KNIP) plugin.

10.3 MATLAB

MATLAB\(^7\) is a high-level language and interactive environment that facilitates rapid development of algorithms for performing computationally intensive tasks.

Calling Bio-Formats from MATLAB is fairly straightforward, since MATLAB has built-in interoperability with Java. We have created a set of scripts\(^8\) for reading image files. Note the minimum supported MATLAB version is R2007b (7.5).

\(^1\)http://www.exelisvis.com/ProductsServices/IDL.aspx
\(^2\)http://kar03.bplaced.net/karo/IDL/_pro/ij\_read\_bio\_formats.pro
\(^3\)http://kar03.bplaced.net/karo/ro\_embed.php?file=IDL/index.html
\(^4\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/
\(^5\)http://www.knime.org/
\(^6\)http://tech.knime.org/community/image-processing
\(^7\)http://www.mathworks.com/products/matlab/
\(^8\)https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/formats-gpl/matlab
10.3.1 Installation

Download the MATLAB toolbox from the Bio-Formats downloads page. Unzip bfmatlab.zip and add the unzipped bf-matlab folder to your MATLAB path.

**Note:** As of Bio-Formats 5.0.0, this zip now contains the bundled jar and you no longer need to download loci_tools.jar or the new bioformats_package.jar separately.

10.3.2 Usage

Please see Using Bio-Formats in MATLAB for usage instructions. If you intend to extend the existing .m files, please also see the developer page for more information on how to use Bio-Formats in general.

10.3.3 Performance

In our tests (MATLAB R14 vs. java 1.6.0_20), the script executes at approximately half the speed of our showinf command line tool, due to overhead from copying arrays.

10.3.4 Upgrading

To use a newer version of Bio-Formats, overwrite the content of the bfmatlab folder with the newer version of the toolbox and restart MATLAB.

10.3.5 Alternative scripts

Several other groups have developed their own MATLAB scripts that use Bio-Formats, including the following:

- https://github.com/prakatmac/bf-tools/
- imread for multiple life science image file formats

10.4 VisAD

The VisAD visualization toolkit is a Java component library for interactive and collaborative visualization and analysis of numerical data. VisAD uses Bio-Formats to read many image formats, notably TIFF.

10.4.1 Installation

The visad.jar file has Bio-Formats bundled inside, so no further installation is necessary.

10.4.2 Upgrading

It should be possible to use a newer version of Bio-Formats by putting the latest bioformats_package.jar or formats-gpl.jar before visad.jar in the class path. Alternately, you can create a “VisAD Lite” using the make lite command from VisAD source, and use the resultant visad-lite.jar, which is a stripped down version of VisAD without sample applications or Bio-Formats bundled in.

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12. [http://www.ssec.wisc.edu/~billh/visad.html](http://www.ssec.wisc.edu/~billh/visad.html)
11.1 Bitplane Imaris

Imaris\(^1\) is Bitplane’s core scientific software module that delivers all the necessary functionality for data visualization, analysis, segmentation and interpretation of 3D and 4D microscopy datasets. Combining speed, precision and ease-of-use, Imaris provides a complete set of features for working with three- and four-dimensional multi-channel images of any size, from a few megabytes to multiple gigabytes in size.

As of version 7.2\(^2\), Imaris integrates with Fiji overview, which includes Bio-Formats. See this page\(^3\) for a detailed list of Imaris’ features.

11.2 CellProfiler

CellProfiler\(^4\)—developed by the Broad Institute\(^5\)’s Imaging Platform\(^6\)—is free open-source software designed to enable biologists without training in computer vision or programming to quantitatively measure phenotypes from thousands of images automatically. CellProfiler uses Bio-Formats to read images from disk, as well as write movies.

11.2.1 Installation

The CellProfiler distribution comes with Bio-Formats included, so no further installation is necessary.

11.2.2 Upgrading

It should be possible to use a newer version of Bio-Formats by replacing the bundled loci_tools.jar with a newer version.

- For example, on Mac OS X, Ctrl+click the CellProfiler icon, choose Show Package Contents, and replace the following files:
  - Contents/Resources/bioformats/loci_tools.jar
  - Contents/Resources/lib/python2.5/bioformats/loci_tools.jar

See also:

CellProfiler web site\(^7\)

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\(^1\)http://www.bitplane.com/
\(^2\)http://www.bitplane.com/releasenotes.aspx?product=Imaris&version=7.2&patch=0
\(^3\)http://www.bitplane.com/imaris/imaris
\(^4\)http://www.cellprofiler.org/
\(^5\)http://www.broadinstitute.org/
\(^6\)http://www.broadinstitute.org/science/platforms/imaging/imaging-platform
\(^7\)http://www.cellprofiler.org/
11.3 Comstat2

Comstat2 is a Java-based computer program for the analysis and treatment of biofilm images in 3D. It is the Master’s project of Martin Vorregaard. Comstat2 uses the Bio-Formats Importer plugin for ImageJ to read files in TIFF and Leica LIF formats.

11.4 Endrov

Endrov (or http://www.endrov.net) (EV) is a multi-purpose image analysis program developed by the Thomas Burglin group at Karolinska Institute, Department of Biosciences and Nutrition.

11.4.1 Installation

The EV distribution comes bundled with the core Bio-Formats library (bio-formats.jar), so no further installation is necessary.

11.4.2 Upgrading

It should be possible to use a newer version of Bio-Formats by downloading the latest formats-gpl.jar and putting it into the libs folder of the EV distribution, overwriting the old file.

You could also include some optional libraries, to add support for additional formats, if desired.

11.5 FocalPoint

FocalPoint is an image browser, similar to Windows Explorer or other file manager application, specifically designed to work with more complex image types. FocalPoint uses Bio-Formats to generate thumbnails for some formats.

11.5.1 Installation

FocalPoint is bundled with Bio-Formats, so no further installation is necessary.

11.5.2 Upgrading

It should be possible to use a newer version of Bio-Formats by overwriting the old loci_tools.jar within the FocalPoint distribution. For Mac OS X, you will have to control click the FocalPoint program icon, choose “Show Package Contents” and navigate into Contents/Resources/Java to find the loci_tools.jar file.

11.6 Graphic Converter

Graphic Converter is a Mac OS application for opening, editing, and organizing photos. Versions 6.4.1 and later use Bio-Formats to open all file formats supported by Bio-Formats.

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8http://www.comstat.dk/  
9https://github.com/mahogny/Endrov  
10http://www.biosci.ki.se/groups/tbu  
11http://www.ki.se/  
12http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/formats-gpl.jar  
13http://www.bioinformatics.bbsrc.ac.uk/projects/focalpoint/  
15http://en.wikipedia.org/wiki/File_manager  
16http://downloads.openmicroscopy.org/latest/bio-formats5.0/  
17http://www.lemkesoft.com
11.7 Icy

Icy\textsuperscript{18} is an open-source image analysis and visualization software package that combines a user-friendly graphical interface with the ability to write scripts and plugins that can be uploaded to a centralized website. It uses Bio-Formats internally to read images and acquisition metadata, so no further installation is necessary.

11.8 imago

Mayachitra imago\textsuperscript{19} is an advanced desktop image management package that enables scientists to easily store, manage, search, and analyze 5D biological images and their analysis results. imago integrates flexible annotation and metadata management with advanced image analysis tools.

imago uses Bio-Formats to read files in some formats, including Bio-Rad PIC, Image-Pro Workspace, Metamorph TIFF, Leica LCS LEI, Olympus FluoView FV1000, Nikon NIS-Elements ND2, and Zeiss LSM.

A free 30-day trial version of imago is available here\textsuperscript{20}.

11.9 Iqm

Iqm\textsuperscript{21} is an image processing application written in Java. It is mainly constructed around the Java JAI library and furthermore it incorporates the functionality of the popular ImageJ image processing software.

Because iqm integrates with ImageJ, it can take advantage of the \textit{Bio-Formats ImageJ plugin} to read image data.

11.10 Macnification

Macnification\textsuperscript{22} is a Mac OS X application for organizing, editing, analyzing and annotating microscopic images, designed for ease of use. It is being developed by Orbicule\textsuperscript{23}.

Macnification uses Bio-Formats to read files in some formats, including Gatan DM3, ICS, ImagePro SEQ, ImagePro IPW, Metamorph STK, OME-TIFF and Zeiss LSM.

See also:

Free trial download\textsuperscript{24}

11.11 MIPAV

The MIPAV\textsuperscript{25} (Medical Image Processing, Analysis, and Visualization) application—developed at the Center for Information Technology\textsuperscript{26} at the National Institutes of Health\textsuperscript{27}—enables quantitative analysis and visualization of medical images of numerous modalities such as PET, MRI, CT, or microscopy. You can use Bio-Formats as a plugin for MIPAV to read images in the formats it supports.

\textsuperscript{18}http://icy.bioimageanalysis.org/
\textsuperscript{19}http://mayachitra.com/imago/index.html
\textsuperscript{20}http://mayachitra.com/imago/download-trial.php
\textsuperscript{21}http://code.google.com/p/iqm/
\textsuperscript{22}http://www.orbicule.com/macnification/
\textsuperscript{23}http://www.orbicule.com
\textsuperscript{24}http://www.orbicule.com/macnification/download
\textsuperscript{25}http://mipav.cit.nih.gov/
\textsuperscript{26}http://cit.nih.gov/
\textsuperscript{27}http://nih.gov/
11.11.1 Installation

Follow these steps to install the Bio-Formats plugin for MIPAV:

1. Download `bioformats_package.jar` and drop it into your MIPAV folder.
2. Download the plugin source code into your `user/mipav/plugins` folder.
3. From the command line, compile the plugin with:

   ```
   cd mipav/plugins
   javac -cp $MIPAV:$MIPAV/bioformats\_package.jar \ PlugInBioFormatsImporter.java
   ```

4. where $MIPAV is the location of your MIPAV installation.
5. Add `bioformats_package.jar` to MIPAV’s class path:
   - How to do so depends on your platform.
   - E.g., in Mac OS X, edit the `mipav.app/Contents/Info.plist` file.

See the readme file for more information.

To upgrade, just overwrite the old `bioformats_package.jar` with the latest one. You may want to download the latest version of MIPAV first, to take advantage of new features and bug-fixes.

11.12 Vaa3D

Vaa3D, developed by the Peng Lab at the HHMI Janelia Farm Research Campus, is a handy, fast, and versatile 3D/4D/5D Image Visualization & Analysis System for Bioimages & Surface Objects.

Vaa3D can use Bio-Formats via the Bio-Formats C++ bindings to read images.

11.13 VisBio

VisBio is a biological visualization tool designed for easy visualization and analysis of multidimensional image data. VisBio uses Bio-Formats to import files as the Bio-Formats library originally grew out of our efforts to continually expand the file format support within VisBio.

11.13.1 Installation

VisBio is bundled with Bio-Formats, so no further installation is necessary.

11.13.2 Upgrading

It should be possible to use a newer version of Bio-Formats by overwriting the old `bio-formats.jar` and optional libraries within the VisBio distribution. For Mac OS X, you’ll have to control click the VisBio program icon, choose “Show Package Contents” and navigate into Contents/Resources/Java to find the JAR files.

---

28 http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/bioformats_package.jar
29 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/utils/mipav/PlugInBioFormatsImporter.java
30 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/utils/mipav/readme.txt
31 http://downloads.openmicroscopy.org/latest/bio-formats5.0/
32 http://vaa3d.org
33 http://penglab.janelia.org/
34 http://www.hhmi.org/janelia/
36 http://loci.wisc.edu/software/visbio
37 http://downloads.openmicroscopy.org/latest/bio-formats5.0/
11.14 XuvTools

XuvTools\(^{38}\) is automated 3D stitching software for biomedical image data. As of release 1.8.0, XuvTools uses Bio-Formats to read image data.

\(^{38}\)http://www.xuvtools.org
Part III

Developer Documentation
12.1 An in-depth guide to using Bio-Formats

12.1.1 Overview

This document describes various things that are useful to know when working with Bio-Formats. It is recommended that you obtain the Bio-Formats source by following the directions in the Source code section. Having a copy of the Javadocs near by is recommended—the notes that follow will make more sense when you see the API.

For a complete list of supported formats, see the Bio-Formats supported formats table.

For a few working examples of how to use Bio-Formats, see these Github pages.

12.1.2 Using Gradle, Maven or Ivy

All released .jar artifacts may be obtained through the OME Artifactory server. The “Client Settings” section of the Artifactory main page provides example code snippets for inclusion into your Gradle, Maven or Ivy project, which will enable the use of this repository.

Example snippets for using the Bio-Formats 5.0.2 release formats-gpl artifact are available for Gradle and for Maven. These may be copied into your project to enable the use of the Bio-Formats library components, and may be adjusted to use different components or different release or development versions of Bio-Formats.

12.1.3 Basic file reading

Bio-Formats provides several methods for retrieving data from files in an arbitrary (supported) format. These methods fall into three categories: raw pixels, core metadata, and format-specific metadata. All methods described here are present and documented in loci.formats.IFormatReader-it is advised that you take a look at the source and/or the Javadocs. In general, it is recommended that you read files using an instance of ImageReader. While it is possible to work with readers for a specific format, ImageReader contains additional logic to automatically detect the format of a file and delegate subsequent calls to the appropriate reader.

Prior to retrieving pixels or metadata, it is necessary to call setId(String) on the reader instance, passing in the name of the file to read. Some formats allow multiple series (5D image stacks) per file; in this case you may wish to call setSeries(int) to change which series is being read.

Raw pixels are always retrieved one plane at a time. Planes are returned as raw byte arrays, using one of the openBytes methods.

Core metadata is the general term for anything that might be needed to work with the planes in a file. A list of core metadata fields is given below, with the appropriate accessor method in parentheses:

- image width (getSizeX())

1http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/  
2https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/formats-gpl/utils  
3http://artifacts.openmicroscopy.org/artifactory  
4https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/IFormatReader.java  
5https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/ImageReader.java  
6http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatHandler.html#setId(java.lang.String)  
7http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#setSeries(int)  
8http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getSizeX()
• image height ($\text{getSizeY()}^9$)
• number of series per file ($\text{getSeriesCount()}^{10}$)
• total number of images per series ($\text{getImageCount()}^{11}$)
• number of slices in the current series ($\text{getSizeZ()}^{12}$)
• number of timepoints in the current series ($\text{getSizeT()}^{13}$)
• number of actual channels in the current series ($\text{getSizeC()}^{14}$)
• number of channels per image ($\text{getRGBChannelCount()}^{15}$)
• the ordering of the images within the current series ($\text{getDimensionOrder()}^{16}$)
• whether each image is RGB ($\text{isRGB()}^{17}$)
• whether the pixel bytes are in little-endian order ($\text{isLittleEndian()}^{18}$)
• whether the channels in an image are interleaved ($\text{isInterleaved()}^{19}$)
• the type of pixel data in this file ($\text{getPixelType()}^{20}$)

All file formats are guaranteed to accurately report core metadata.

Format-specific metadata refers to any other data specified in the file - this includes acquisition and hardware parameters, among other things. This data is stored internally in a java.util.Hashtable, and can be accessed in one of two ways: individual values can be retrieved by calling $\text{getMetadataValue(String)}^{21}$, which gets the value of the specified key. Note that the keys in this Hashtable are different for each format, hence the name “format-specific metadata”.

See Bio-Formats metadata processing for more information on the metadata capabilities that Bio-Formats provides.

12.1.4 File reading extras

The previous section described how to read pixels as they are stored in the file. However, the native format is not necessarily convenient, so Bio-Formats provides a few extras to make file reading more flexible.

• There are a few “wrapper” readers (that implement IFormatReader) that take a reader in the constructor, and manipulate the results somehow, for convenience. Using them is similar to the java.io InputStream/OutputStream model: just layer whichever functionality you need by nesting the wrappers.
  – BufferedImageReader$^{22}$ extends IFormatReader, and allows pixel data to be returned as BufferedImages instead of raw byte arrays.
  – FileStitcher$^{23}$ extends IFormatReader, and uses advanced pattern matching heuristics to group files that belong to the same dataset.
  – ChannelSeparator$^{24}$ extends IFormatReader, and makes sure that all planes are grayscale - RGB images are split into 3 separate grayscale images.
  – ChannelMerger$^{25}$ extends IFormatReader, and merges grayscale images to RGB if the number of channels is greater than 1.
  – ChannelFiller$^{26}$ extends IFormatReader, and converts indexed color images to RGB images.

---

9 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getSizeY()
10 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getSeriesCount()
11 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getImageCount()
12 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getSizeZ()
13 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getSizeT()
14 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getSizeC()
15 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getType()
16 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getDimensionOrder()
17 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#isRGB()
18 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#isLittleEndian()
19 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#isInterleaved()
20 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getType()
21 http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#getValue(java.lang.String)
22 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/gui/BufferedImageReader.java
23 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/FileStitcher.java
24 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/ChannelSeparator.java
25 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/ChannelMerger.java
26 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/ChannelFiller.java
- MinMaxCalculator<sup>27</sup> extends IFormatReader, and provides an API for retrieving the minimum and maximum pixel values for each channel.
- DimensionSwapper<sup>28</sup> extends IFormatReader, and provides an API for changing the dimension order of a file.

- ImageTools<sup>29</sup> and loci.formats.gui.AWTImageTools<sup>30</sup> provide a number of methods for manipulating BufferedImages and primitive type arrays. In particular, there are methods to split and merge channels in a BufferedImage/array, as well as converting to a specific data type (e.g. convert short data to byte data).

### 12.1.5 Writing files

The following file formats can be written using Bio-Formats:

- TIFF (uncompressed, LZW, JPEG, or JPEG-2000)
- OME-TIFF (uncompressed, LZW, JPEG, or JPEG-2000)
- JPEG
- PNG
- AVI (uncompressed)
- QuickTime (uncompressed is supported natively; additional codecs use QTJava)
- Encapsulated PostScript (EPS)
- OME-XML (not recommended)

The writer API (see loci.formats.IFormatWriter<sup>31</sup>) is very similar to the reader API, in that files are written one plane at time (rather than all at once).

All writers allow the output file to be changed before the last plane has been written. This allows you to write to any number of output files using the same writer and output settings (compression, frames per second, etc.), and is especially useful for formats that do not support multiple images per file.

Please see loci.formats.tools.ImageConverter<sup>32</sup> and this guide to exporting to OME-TIFF files for examples of how to write files.

### 12.1.6 Arcane notes and implementation details

Known oddities:

- Importing multi-file formats (Leica LEI, PerkinElmer, FV1000 OIF, ICS, and Prairie TIFF, to name a few) can fail if any of the files are renamed. There are “best guess” heuristics in these readers, but they are not guaranteed to work in general. Please do not rename files in these formats.
- If you are working on a Macintosh, make sure that the data and resource forks of your image files are stored together. Bio-Formats does not handle separated forks (the native QuickTime reader tries, but usually fails).

### 12.2 Obtaining and building Bio-Formats

#### 12.2.1 Source code

The source code for this Bio-Formats release is available from the download page<sup>33</sup>. This release and the latest Bio-Formats source code are also available from the Git repository. This may be accessed using the repository path:

```plaintext
27 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/MinMaxCalculator.java
28 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/DimensionSwapper.java
31 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/IFormatWriter.java
33 http://downloads.openmicroscopy.org/latest/bio-formats5.0/
```
More information about Git and client downloads are available from the Git project website\(^{34}\). You can also browse the Bio-Formats source on GitHub\(^{35}\).

**Note:** Windows users must set git to use `core.autocrlf=input` to ensure that Bio-Formats uses LF rather than CRLF line endings, otherwise the build will fail (Genshi can’t process code templates with CRLF line endings, leading to broken sources being generated). This can be set globally in the registry when installing `msysgit` or by editing `etc/gitconfig` in the git installation directory. Annoyingly, these settings appear to override per-user and per-repository configuration values, requiring these to be set globally.

Lastly, you can browse the Bio-Formats Javadocs online\(^{36}\), or generate them yourself using the “docs” Ant target.

### 12.2.2 Source code structure

The Bio-Formats code is divided into several projects. Core components are located in subfolders of the `components`\(^{37}\) folder, with some components further classified into `components/forks`\(^{38}\) or `components/stubs`\(^{39}\), depending on the nature of the project.

Each project has a corresponding Maven POM file, which can be used to work with the project in your favorite IDE, or from the command line, once you have cloned the source.

### 12.2.3 Building from source

Instructions for several popular options follow. In all cases, make sure that the prerequisites are installed before you begin.

If you are interested in working on the Bio-Formats source code itself, you can load it into your favorite IDE, or develop with your favorite text editor.

**Prerequisites**

In addition to the Bio-Formats source code, the following programs and packages are also required:

- Python\(^2\), version 2.6 or later (note: not version 3)
- Genshi\(^{41}\) 0.5 or later (0.7 recommended)

**Note:** Genshi may be installed (in order of decreasing preference) with some Linux distributions’ package managers, `pip` (pip install genshi), by downloading a compatible .egg for your system from the Genshi download page\(^{42}\), or from source. If using a .egg, make sure it is added to your PYTHONPATH environment variable.

**NetBeans**

NetBeans comes with Maven support built in. To import the Bio-Formats source, perform the following steps:

1. Choose **File → Open Project** from the menu
2. Select the top-level folder of your Bio-Formats working copy
3. Expand the Modules folder and double-click desired project(s) to work with them

---

\(^{34}\) [http://git-scm.com/](http://git-scm.com/)
\(^{35}\) [https://github.com/openmicroscopy/bioformats](https://github.com/openmicroscopy/bioformats)
\(^{36}\) [http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/](http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/)
\(^{37}\) [https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/](https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/)
\(^{38}\) [https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/forks/](https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/forks/)
\(^{39}\) [https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/stubs/](https://github.com/openmicroscopy/bioformats/tree/v5.0.4/components/stubs/)
\(^{40}\) [http://python.org](http://python.org)
\(^{41}\) [http://genshi.edgewall.org](http://genshi.edgewall.org)
\(^{42}\) [http://genshi.edgewall.org/wiki/Download](http://genshi.edgewall.org/wiki/Download)
Alternately, you can clone the source directly from NetBeans into a project by selecting Team → Git → Clone Other... from the menu.

**Eclipse**

Eclipse uses the “Maven Integration for Eclipse” (m2e) plugin to work with Maven projects. It is more flexible than Eclipse’s built-in project management because m2e transparently converts between project dependencies and JAR dependencies (stored in the Maven repository in ~/.m2/repository) on the build path, depending on which projects are currently open.

We recommend using Eclipse 4.3 (Kepler), specifically - “Eclipse IDE for Java developers”. It comes with m2e installed (http://eclipse.org/downloads/compare.php?release=kepler).

You can then import the Bio-Formats source by choosing File → Import → Existing Maven Projects from the menu and browsing to the top-level folder of your Bio-Formats working copy.

**Command line**

If you prefer developing code with a text editor such as vim or emacs, you can use the Ant or Maven command line tools to compile Bio-Formats. The Bio-Formats source tree provides parallel build systems for both Ant and Maven, so you can use either one to build the code.

For a list of Ant targets, run:

```
ant -p
```

In general, `ant jars` or `ant tools` is the correct command.

When using Maven, Bio-Formats is configured to run the “install” target by default, so all JARs will be copied into your local Maven repository in ~/.m2/repository. Simply run:

```
mvn
```

With either Ant or Maven, you can use similar commands in any subproject folder to build just that component.

### 12.3 Generating test images

Sometimes it is nice to have a file of a specific size or pixel type for testing. To generate a file (that contains gradient images):

```
touch "my-special-test-file\pixelType=uint8\sizeX=8192\sizeY=8192.fake"
```

Whatever is before the & is the image name; remaining key value pairs should be pretty self-explanatory. Just replace the values with whatever you need for testing.

There are a few other keys that can be added as well:

<table>
<thead>
<tr>
<th>Key</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td><code>sizeZ</code></td>
<td>number of Z sections</td>
</tr>
<tr>
<td><code>sizeC</code></td>
<td>number of channels</td>
</tr>
<tr>
<td><code>sizeT</code></td>
<td>number of timepoints</td>
</tr>
<tr>
<td><code>bitsPerPixel</code></td>
<td>number of valid bits (&lt;= number of bits implied by pixel type)</td>
</tr>
<tr>
<td><code>rgb</code></td>
<td>number of channels that are merged together</td>
</tr>
<tr>
<td><code>dimOrder</code></td>
<td>dimension order (e.g. XYZCT)</td>
</tr>
<tr>
<td><code>little</code></td>
<td>whether or not the pixel data should be little-endian</td>
</tr>
<tr>
<td><code>interleaved</code></td>
<td>whether or not merged channels are interleaved</td>
</tr>
<tr>
<td><code>indexed</code></td>
<td>whether or not a color lookup table is present</td>
</tr>
<tr>
<td><code>falseColor</code></td>
<td>whether or not the color lookup table is just for making the image look pretty</td>
</tr>
<tr>
<td><code>series</code></td>
<td>number of series (Images)</td>
</tr>
<tr>
<td><code>lutLength</code></td>
<td>number of entries in the color lookup table</td>
</tr>
</tbody>
</table>

12.3. Generating test images
You can often work with the .fake file directly, but in some cases support for those files is disabled and so you will need to convert the file to something else. Make sure that you have Bio-Formats built and the JARs in your CLASSPATH (individual JARs or just bioformats_package.jar):

```
bfconvert test&pixelType=uint8&sizeX=8192&sizeY=8192.fake test.tiff
```

If you do not have the command line tools installed, substitute `lociformats.tools.ImageConverter` for `bfconvert`.

---

13.1 API documentation

13.1.1 Using Bio-Formats as a Java library

If you wish to make use of Bio-Formats within your own software, you can download formats-gpl.jar\(^1\) to use it as a library. Just add formats-gpl.jar to your CLASSPATH or build path. You will also need common.jar for common I/O functions, ome-xml.jar for metadata standardization, and SLF4J\(^2\) for logging.

There are also certain packages that if present will be utilized to provide additional functionality. To include one, just place it in the same folder.

<table>
<thead>
<tr>
<th>Package</th>
<th>Filename</th>
<th>License</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>Apache Jakarta POI(^3) library, OME fork</td>
<td>ome-poi.jar(^4)</td>
<td>Apache</td>
<td>For OLE-based formats (zvi, oib, ipw, cxd)</td>
</tr>
<tr>
<td>MDB Tools project(^5) Java port, OME fork</td>
<td>mdbtools-java.jar(^6)</td>
<td>LGPL</td>
<td>For Olympus CellR and Zeiss LSM metadata (mdb)</td>
</tr>
<tr>
<td>JAI Image I/O Tools(^7) pure Java implementation, OME fork</td>
<td>jai_imageio.jar(^8)</td>
<td>BSD</td>
<td>For JPEG2000-based formats (nd2, jp2)</td>
</tr>
<tr>
<td>NetCDF Java library(^9)</td>
<td>netcdf-4.3.19.jar(^10)</td>
<td>LGPL</td>
<td>For HDF5-based formats (Imaris 5.5, MINC MRI)</td>
</tr>
<tr>
<td>QuickTime for Java(^11)</td>
<td>QTJava.zip</td>
<td>Commercial</td>
<td>For additional QuickTime codecs</td>
</tr>
</tbody>
</table>

See the list in the Bio-Formats toplevel build file\(^21\) for a complete and up-to-date list of all optional libraries, which can all be found in our Git repository\(^22\).

Examples of usage

**ImageConverter**\(^23\) - A simple command line tool for converting between formats.

\(^1\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/formats-gpl.jar  
\(^2\)http://slf4j.org/  
\(^3\)http://jakarta.apache.org/poi/  
\(^4\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/ome-poi.jar  
\(^5\)http://sourceforge.net/projects/mdbtools  
\(^6\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/mdbtools-java.jar  
\(^7\)http://java.net/projects/jai-imageio  
\(^8\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/jai_imageio.jar  
\(^9\)http://www.unidata.ucar.edu/software/netcdf-java/  
\(^10\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/netcdf-4.3.19.jar  
\(^12\)http://jakarta.apache.org/poi/  
\(^13\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/ome-poi.jar  
\(^14\)http://sourceforge.net/projects/mdbtools  
\(^15\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/mdbtools-java.jar  
\(^16\)http://java.net/projects/jai-imageio  
\(^17\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/jai_imageio.jar  
\(^18\)http://www.unidata.ucar.edu/software/netcdf-java/  
\(^19\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/artifacts/netcdf-4.3.19.jar  
\(^20\)http://www.apple.com/quicktime/download/standalone.html  
\(^21\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/build.xml  
\(^22\)https://github.com/openmicroscopy/bioformats/tree/v5.0.4/jar  
\(^23\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/bio-formats-tools/src/loci/formats/tools/ImageConverter.java
ImageInfo\textsuperscript{24} - A more involved command line utility for thoroughly reading an input file, printing some information about it, and displaying the pixels on screen using the Bio-Formats viewer.

MinimumWriter\textsuperscript{25} - A command line utility demonstrating the minimum amount of metadata needed to write a file.

PrintTimestamps\textsuperscript{26} - A command line example demonstrating how to extract timestamps from a file.

Simple_Read\textsuperscript{27} - A simple ImageJ plugin demonstrating how to use Bio-Formats to read files into ImageJ (see ImageJ overview).

Read_Image\textsuperscript{28} - An ImageJ plugin that uses Bio-Formats to build up an image stack, reading image planes one by one (see ImageJ overview).

Mass_Importer\textsuperscript{29} - A simple plugin for ImageJ that demonstrates how to open all image files in a directory using Bio-Formats, grouping files with similar names to avoid opening the same dataset more than once (see ImageJ overview).

### A Note on Java Web Start (bioformats_package.jar vs. formats-gpl.jar)

To use Bio-Formats with your Java Web Start application, we recommend using \texttt{formats-gpl.jar} rather than \texttt{bioformats_package.jar}—the latter is merely a bundle of \texttt{formats-gpl.jar} plus all its optional dependencies.

The \texttt{bioformats_package.jar} bundle is intended as a convenience (e.g. to simplify installation as an ImageJ plugin), but is by no means the only solution for developers. We recommend using \texttt{formats-gpl.jar} as a separate entity depending on your needs as a developer.

The bundle is quite large because we have added support for several formats that need large helper libraries (e.g. Imaris’ HDF-based format). However, these additional libraries are optional; Bio-Formats has been coded using reflection so that it can both compile and run without them.

When deploying a JNLP-based application, using \texttt{bioformats_package.jar} directly is not the best approach, since every time Bio-Formats is updated, the server would need to feed another 15+ MB JAR file to the client. Rather, Web Start is a case where you should keep the JARs separate, since JNLP was designed to make management of JAR dependencies trivial for the end user. By keeping \texttt{formats-gpl.jar} and the optional dependencies separate, only a <1 MB JAR needs to be updated when \texttt{formats-gpl.jar} changes.

As a developer, you have the option of packaging \texttt{formats-gpl.jar} with as many or as few optional libraries as you wish, to cut down on file size as needed. You are free to make whatever kind of “stripped down” version you require. You could even build a custom \texttt{formats-gpl.jar} that excludes certain classes, if you like.

For an explicit enumeration of all the optional libraries included in \texttt{bioformats_package.jar}, see the package.libraries variable of the \texttt{ant/toplevel.properties}\textsuperscript{30} file of the distribution. You can also read our notes about each in the source distribution’s Ant build.xml\textsuperscript{31} script.

Also see Bio-Formats Javadocs\textsuperscript{32}

### 13.2 Examples

#### 13.2.1 Exporting files using Bio-Formats

This guide pertains to version 4.2 and later.

**Basic conversion**

The first thing we need to do is set up a reader:
// create a reader that will automatically handle any supported format
IFormatReader reader = new ImageReader();
// tell the reader where to store the metadata from the dataset
MetadataStore metadata;

try {
    ServiceFactory factory = new ServiceFactory();
    OMEXMLService service = factory.getInstance(OMEXMLService.class);
    metadata = service.createOMEXMLMetadata();
} catch (DependencyException exc) {
    throw new FormatException("Could not create OME-XML store.", exc);
} catch (ServiceException exc) {
    throw new FormatException("Could not create OME-XML store.", exc);
}

reader.setMetadataStore(metadata);
// initialize the dataset
reader.setId("/path/to/file");

Now, we set up our writer:

// create a writer that will automatically handle any supported output format
IFormatWriter writer = new ImageWriter();
// give the writer a MetadataRetrieve object, which encapsulates all of the
// dimension information for the dataset (among many other things)
writer.setMetadataRetrieve(MetadataTools.asRetrieve(reader.getMetadataStore()));
// initialize the writer
writer.setId("/path/to/output/file");

Note that the extension of the file name passed to ‘writer.setId(…)’ determines the file format of the exported file.

Now that everything is set up, we can start writing planes:

for (int series=0; series<reader.getSeriesCount(); series++) {
    reader.setSeries(series);
    writer.setSeries(series);

    for (int image=0; image<reader.getImageCount(); image++) {
        writer.saveBytes(image, reader.openBytes(image));
    }
}

Finally, make sure to close both the reader and the writer. Failure to do so can cause:

- file handle leaks
- memory leaks
- truncated output files

Fortunately, closing the files is very easy:

reader.close();
writer.close();

### Converting large images

The flaw in the previous example is that it requires an image plane to be fully read into memory before it can be saved. In many cases this is fine, but if you are working with very large images (especially > 4 GB) this is problematic. The solution is to break
each image plane into a set of reasonably-sized tiles and save each tile separately - thus substantially reducing the amount of memory required for conversion.

For now, we’ll assume that your tile size is 1024 x 1024, though in practice you will likely want to adjust this. Assuming you have an IFomatReader and IFomatWriter set up as in the previous example, let’s start writing planes:

```java
int tileWidth = 1024;
int tileHeight = 1024;

for (int series=0; series<reader.getSeriesCount(); series++) {
    reader.setSeries(series);
    writer.setSeries(series);

    // determine how many tiles are in each image plane
    // for simplicity, we’ll assume that the image width and height are
    // multiples of 1024

    int tileRows = reader.getSizeY() / tileHeight;
    int tileColumns = reader.getSizeX() / tileWidth;

    for (int image=0; image<reader.getImageCount(); image++) {
        for (int row=0; row<tileRows; row++) {
            for (int col=0; col<tileColumns; col++) {
                // open a tile - in addition to the image index, we need to specify
                // the (x, y) coordinate of the upper left corner of the tile,
                // along with the width and height of the tile

                int xCoordinate = col * tileWidth;
                int yCoordinate = row * tileHeight;
                byte[] tile =
                    reader.openBytes(image, xCoordinate, yCoordinate, tileWidth, tileHeight);
                writer.saveBytes(
                    image, tile, xCoordinate, yCoordinate, tileWidth, tileHeight);
            }
        }
    }
}
```

As noted, the example assumes that the width and height of the image are multiples of the tile dimensions. Be careful, as this is not always the case; the last column and/or row may be smaller than preceding columns/rows. An exception will be thrown if you attempt to read or write a tile that is not completely contained by the original image plane. Most writers perform best if the tile width is equal to the image width, although specifying any valid width should work.

As before, you need to close the reader and writer.

### Converting to multiple files

The recommended method of converting to multiple files is to use a single IFomatWriter, like so:

```java
// you should have set up a reader as in the first example
ImageWriter writer = new ImageWriter();
writer.setMetadataRetrieve(MetadataTools.asRetrieve(reader.getMetadataStore()));
// replace this with your own filename definitions
// in this example, we’re going to write half of the planes to one file
// and half of the planes to another file
String[] outputFiles =
    new String[] {"/path/to/file/1.tiff", "/path/to/file/2.tiff"};
writer.setId(outputFiles[0]);

int planesPerFile = reader.getImageCount() / outputFiles.length;
for (int file=0; file<outputFiles.length; file++) {
    writer.changeOutputFile(outputFiles[file]);
}
```
The advantage here is that the relationship between the files is preserved when converting to formats that support multi-file datasets internally (namely OME-TIFF). If you are only converting to graphics formats (e.g. JPEG, AVI, MOV), then you could also use a separate IFormatWriter for each file, like this:

```java
// again, you should have set up a reader already
String[] outputFiles = new String[] {"/path/to/file/1.avi", "/path/to/file/2.avi"};
int planesPerFile = reader.getImageCount() / outputFiles.length;
for (int file=0; file<outputFiles.length; file++) {
    ImageWriter writer = new ImageWriter();
    writer.setMetadataRetrieve(MetadataTools.asRetrieve(reader.getMetadataStore()));
    writer.setId(outputFiles[file]);
    for (int image=0; image<planesPerFile; image++) {
        int index = file * planesPerFile + image;
        writer.saveBytes(image, reader.openBytes(index));
    }
    writer.close();
}
```

### Known issues

List of Trac tickets

#### 13.2.2 Further details on exporting raw pixel data to OME-TIFF files

This document explains how to export pixel data to OME-TIFF using Bio-Formats version 4.2 and later.

The first thing that must happen is we must create the object that stores OME-XML metadata. This is done as follows:

```java
ServiceFactory factory = new ServiceFactory();
OMEXMLService service = factory.getInstance(OMEXMLService.class);
IMetadata omexml = service.createOMEXMLMetadata();
```

The `omexml` object can now be used in our code to store OME-XML metadata, and by the file format writer to retrieve OME-XML metadata.

Now that we have somewhere to put metadata, we need to populate as much metadata as we can. The minimum amount of metadata required is:

- endianness of the pixel data
- the order in which dimensions are stored
- the bit depth of the pixel data
- the number of channels
- the number of timepoints
- the number of Z sections

• the width (in pixels) of an image
• the height (in pixels) of an image
• the number of samples per channel (3 for RGB images, 1 otherwise)

We populate that metadata as follows:

```java
omexml.setImageID("Image:0", 0);
omexml.setPixelsID("Pixels:0", 0);

// specify that the pixel data is stored in big-endian order
// replace 'TRUE' with 'FALSE' to specify little-endian order
omexml.setPixelsBinDataBigEndian(Boolean.TRUE, 0, 0);

omexml.setPixelsDimensionOrder(DimensionOrder.XYCZT, 0);
omexml.setPixelsType(PixelType.UINT16, 0);
omexml.setPixelsSizeX(new PositiveInteger(width), 0);
omexml.setPixelsSizeY(new PositiveInteger(height), 0);
omexml.setPixelsSizeZ(new PositiveInteger(zSectionCount), 0);
omexml.setPixelsSizeC(new PositiveInteger(channelCount * samplesPerChannel), 0);
omexml.setPixelsSizeT(new PositiveInteger(timepointCount), 0);
```

for (int channel=0; channel<channelCount; channel++) {
    omexml.setChannelID("Channel:0:" + channel, 0, channel);
    omexml.setChannelSamplesPerPixel(new PositiveInteger(samplesPerChannel), 0, channel);
}

There is much more metadata that can be stored; please see the Javadoc for loci.formats.meta.MetadataStore for a complete list.

Now that we have defined all of the metadata, we need to create a file writer:

```java
ImageWriter writer = new ImageWriter();
```

Now we must associate the ‘omexml’ object with the file writer:

```java
writer.setMetadataRetrieve(omexml);
```

The writer now knows to retrieve any metadata that it needs from ‘omexml’.

We now tell the writer which file it should write to:

```java
writer.setId("output-file.ome.tiff");
```

It is critical that the file name given to the writer ends with ".ome.tiff" or ".ome.tif", as it is the file name extension that determines which format will be written.

Now that everything is set up, we can save the image data. This is done plane by plane, and we assume that the pixel data is stored in a 2D byte array ‘pixelData’:

```java
int sizeC = omexml.getPixelsSizeC(0).getValue();
int sizeZ = omexml.getPixelsSizeZ(0).getValue();
int sizeT = omexml.getPixelsSizeT(0).getValue();
int samplesPerChannel = omexml.getChannelSamplesPerPixel(0).getValue();
sizeC /= samplesPerChannel;

int imageCount = sizeC * sizeZ * sizeT;
```

for (int image=0; image<imageCount; image++) { 13.2. Examples 67
Finally, we must tell the writer that we are finished, so that the output file can be properly closed:

```java
textwriter.close();
```

There should now be a complete OME-TIFF file at whichever path was specified above.

### 13.2.3 Converting files from FV1000 OIB/OIF to OME-TIFF

This document explains how to convert a file from FV1000 OIB/OIF to OME-TIFF using Bio-Formats version 4.2 and later. The first thing that must happen is we must create the object that stores OME-XML metadata. This is done as follows:

```java
ServiceFactory factory = new ServiceFactory();
OMEXMLService service = factory.getInstance(OMEXMLService.class);
IMetadata omexml = service.createOMEXMLMetadata();
```

The `omexml` object can now be used by both a file format reader and a file format writer for storing and retrieving OME-XML metadata.

Now that have somewhere to put metadata, we need to create a file reader and writer:

```java
ImageReader reader = new ImageReader();
ImageWriter writer = new ImageWriter();
```

Now we must associate the `omexml` object with the file reader and writer:

```java
reader.setMetadataStore(omexml);
writer.setMetadataRetrieve(omexml);
```

The reader now knows to store all of the metadata that it parses into `omexml`, and the writer knows to retrieve any metadata that it needs from `omexml`.

We now tell the reader and writer which files will be read from and written to, respectively:

```java
reader.setId("input-file.oib");
writer.setId("output-file.ome.tiff");
```

It is critical that the filename given to the writer ends with "".ome.tiff"" or "".ome.tif"", as it is the file name extension that determines which format will be written.

Now that everything is set up, we can convert the image data. This is done plane by plane:

```java
for (int series=0; series<reader.getSeriesCount(); series++) {
    reader.setSeries(series);
    writer.setSeries(series);
    byte[] plane = new byte[FormatTools.getPlaneSize(reader)];
    for (int image=0; image<reader.getImageCount(); image++) {
        reader.openBytes(image, plane);
        writer.saveBytes(image, plane);
    }
}
```
The body of the outer ‘for’ loop may also be replaced with the following:

```java
reader.setSeries(series);
writer.setSeries(series);
for (int image=0; image<reader.getImageCount(); image++) {
    byte[] plane = reader.openBytes(image);
    writer.saveBytes(image, plane);
}
```

But note that this will be a little slower.

Finally, we must tell the reader and writer that we are finished, so that the input and output files can be properly closed:

```java
reader.close();
writer.close();
```

There should now be a complete OME-TIFF file at whichever path was specified above.

### 13.2.4 Using Bio-Formats in MATLAB

This section assumes that you have installed the MATLAB toolbox as instructed in the [MATLAB user information page](#). Note the minimum supported MATLAB version is R2007b (7.5).

#### Increasing JVM memory settings

The default JVM settings in MATLAB can result in `java.lang.OutOfMemoryError: Java heap space` exceptions when opening large image files using Bio-Formats. Information about the Java heap space usage in MATLAB can be retrieved using:

```java
java.lang.Runtime.getRuntime.maxMemory
```

Default JVM settings can be increased by creating a `java.opts` file in the startup directory and overriding the default memory settings. We recommend using `-Xmx512m` in your `java.opts` file.

See also:

[http://www.mathworks.com/matlabcentral/answers/92813](http://www.mathworks.com/matlabcentral/answers/92813) How do I increase the heap space for the Java VM in MATLAB 6.0 (R12) and later versions?

#### Opening an image file

The first thing to do is initialize a file with the `bfopen` function:

```matlab
data = bfopen('/path/to/data/file');
```

This function returns an n-by-4 cell array, where n is the number of series in the dataset. If s is the series index between 1 and n:

- The `data{s, 1}` element is an m-by-2 cell array, where m is the number of planes in the s-th series. If t is the plane index between 1 and m:
  - The `data{s, 1}{t, 1}` element contains the pixel data for the t-th plane in the s-th series.
  - The `data{s, 1}{t, 2}` element contains the label for the t-th plane in the s-th series.
- The `data{s, 2}` element contains original metadata key/value pairs that apply to the s-th series.
- The `data{s, 3}` element contains color lookup tables for each plane in the s-th series.

34[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/bfopen.m](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/bfopen.m)
• The data{s, 4} element contains a standardized OME metadata structure, which is the same regardless of the input file format, and contains common metadata values such as physical pixel sizes - see OME metadata below for examples.

Accessing planes

Here is an example of how to unwrap specific image planes for easy access:

```java
data = bfopen('/path/to/data/file');
seriesCount = size(data, 1);
series1 = data{1, 1};
series2 = data{2, 1};
series3 = data{3, 1};
metadataList = data{1, 2};
% ...etc.
series1_planeCount = size(series1, 1);
series1_plane1 = series1{1, 1};
series1_label1 = series1{1, 2};
series1_plane2 = series1{2, 1};
series1_label2 = series1{2, 2};
series1_plane3 = series1{3, 1};
series1_label3 = series1{3, 2};
% ...etc.
```

Displaying images

If you want to display one of the images, you can do so as follows:

```java
data = bfopen('/path/to/data/file');
% plot the 1st series’s 1st image plane in a new figure
series1 = data{1, 1};
series1_plane1 = series1{1, 1};
series1_label1 = series1{1, 2};
series1_colorMaps = data{1, 3};
figure('Name', series1_label1);
if (isempty(series1_colorMaps{1}))
    colormap(gray);
else
    colormap(series1_colorMaps{1});
end
imagesc(series1_plane1);
```

This will display the first image of the first series with its associated color map (if present). If you would prefer not to apply the color maps associated with each image, simply comment out the calls to colormap.

If you have the image processing toolbox, you could instead use:

```java
imshow(series1_plane1, []);
```

You can also create an animated movie (assumes 8-bit unsigned data):

```java
v = linspace(0, 1, 256)';
cmap = [v v v];
for p = 1 : size(series1, 1)
    M(p) = im2frame(uint8(series1{p, 1}), cmap);
end
movie(M);
```
Retrieving metadata

There are two kinds of metadata:

- **Original metadata** is a set of key/value pairs specific to the input format of the data. It is stored in the data{s, 2} element of the data structure returned by `bfopen`.

- **OME metadata** is a standardized metadata structure, which is the same regardless of input file format. It is stored in the data{s, 4} element of the data structure returned by `bfopen`, and contains common metadata values such as physical pixel sizes, instrument settings, and much more. See the OME Model and Formats\textsuperscript{35} documentation for full details.

**Original metadata**  To retrieve the metadata value for specific keys:

```matlab
data = bfopen('/path/to/data/file');
% Query some metadata fields (keys are format-dependent)
metadata = data{1, 2};
subject = metadata.get('Subject');
title = metadata.get('Title');
```

To print out all of the metadata key/value pairs for the first series:

```matlab
data = bfopen('/path/to/data/file');
metadata = data{1, 2};
metadataKeys = metadata.keySet().iterator();
for i=1:metadata.size()
    key = metadataKeys.nextElement();
    value = metadata.get(key);
    fprintf('%s = %s
', key, value)
end
```

**OME metadata**  Conversion of metadata to the OME standard is one of Bio-Formats’ primary features. The OME metadata is always stored the same way, regardless of input file format.

To access physical voxel and stack sizes of the data:

```matlab
data = bfopen('/path/to/data/file');
omeMeta = data{1, 4};
stackSizeX = omeMeta.getPixelsSizeX(0).getValue(); % image width, pixels
stackSizeY = omeMeta.getPixelsSizeY(0).getValue(); % image height, pixels
stackSizeZ = omeMeta.getPixelsSizeZ(0).getValue(); % number of Z slices
voxelSizeX = omeMeta.getPixelsPhysicalSizeX(0).getValue(); % in µm
voxelSizeY = omeMeta.getPixelsPhysicalSizeY(0).getValue(); % in µm
voxelSizeZ = omeMeta.getPixelsPhysicalSizeZ(0).getValue(); % in µm
```

For more information about the methods to retrieve the metadata, see the MetadataRetrieve\textsuperscript{36} Javadoc page.

To convert the OME metadata into a string, use the `dumpXML()` method:

```matlab
omeXML = char(omeMeta.dumpXML());
```

**Reading from an image file**

The main inconvenience of the `bfopen.m`\textsuperscript{37} function is that it loads all the content of an image regardless of its size.

\textsuperscript{35}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{36}http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/meta/MetadataRetrieve.html
\textsuperscript{37}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/bfopen.m
To access the file reader without loading all the data, use the low-level `bfGetReader.m` function:

```matlab
reader = bfGetReader('path/to/data/file');
```

You can then access the OME metadata using the `getMetadataStore()` method:

```matlab
omeMeta = reader.getMetadataStore();
```

Individual planes can be queried using the `bfGetPlane.m` function:

```matlab
series1_plane1 = bfGetPlane(reader, 1);
```

### Saving files

The basic code for saving a 5D array into an OME-TIFF file is located in the `bfsave.m` function.

For instance, the following code will save a single image of 64 pixels by 64 pixels with 8 unsigned bits per pixels:

```matlab
plane = zeros(64, 64, 'uint8');
bfsave(plane, 'my-file.ome.tiff');
```

And the following code snippet will produce an image of 64 pixels by 64 pixels with 2 channels and 2 timepoints:

```matlab
plane = zeros(64, 64, 1, 2, 2, 'uint8');
bfsave(plane, 'my-file.ome.tiff');
```

By default, `bfsave` will create a minimal OME-XML metadata object containing basic information such as the pixel dimensions, the dimension order and the pixel type. To customize the OME metadata, it is possible to create a metadata object from the input array using `createMinimalOMEXMLMetadata.m`, add custom metadata and pass this object directly to `bfsave`:

```matlab
plane = zeros(64, 64, 1, 2, 2, 'uint8');
metadata = createMinimalOMEXMLMetadata(plane);
pixelSize = ome.xml.model.primitives.PositiveFloat(java.lang.Double(.05));
metadata.setPixelsPhysicalSizeX(pixelSize, 0);
metadata.setPixelsPhysicalSizeY(pixelSize, 0);
pixelSizeZ = ome.xml.model.primitives.PositiveFloat(java.lang.Double(.2));
metadata.setPixelsPhysicalSizeZ(pixelSizeZ, 0);
...
bfsave(plane, 'my-file.ome.tif', 'metadata', metadata);
```

For more information about the methods to store the metadata, see the `MetadataStore` Javadoc page.

---

38https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/bfGetReader.m
39https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/bfGetPlane.m
40https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/bfsave.m
41https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/matlab/createMinimalOMEXMLMetadata.m
42http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/meta/MetadataStore.html
CHAPTER
FOURTEEN

INTERFACING FROM NON-JAVA CODE

14.1 Interfacing with Bio-Formats from non-Java code

Bio-Formats is written in Java, and is easiest to use with other Java code. However, it is possible to call Bio-Formats from a program written in another language. But how to do so depends on your program’s needs.

Technologically, there are two broad categories of solutions: **in-process** approaches, and **inter-process** communication.

For details, see LOCI’s article [Interfacing from non-Java code](http://loci.wisc.edu/software/interfacing-non-java-code).

Recommended **in-process solution**: Bio-Formats C++ bindings

Recommended **inter-process solution**: Subimager

14.2 Bio- Formats C++ bindings

To make Bio-Formats accessible to software written in C++, we have created a Bio-Formats C++ interface (BF-CPP for short). It uses LOCI’s [jar2lib](http://loci.wisc.edu/software/jar2lib) program to generate a C++ proxy class for each equivalent Bio-Formats Java class. The resulting proxies are then compiled into a library, which represents the actual interface from C++ to Bio-Formats. Using this library in your projects gives you access to the image support of Bio-Formats.

BF-CPP comes with some standalone examples which you can use as a starting point in your own project:

- showinf
- minimum_writer

Other projects using BF-CPP include:

- [WiscScan](http://loci.wisc.edu/software/wiscscan) which uses BF-CPP to write OME-TIFF files.
- [XuvTools](http://www.xuvtools.org/devel:libblitzbioformats) which uses an adapted version of BF-CPP called BlitzBioFormats.

See the [build instructions](http://loci.wisc.edu/software/interfacing-non-java-code) for details on compiling BF-CPP from source. Once this is done, simply include it in your project as you would any other external library.

14.3 Build instructions for C++ bindings

This package provides language bindings for calling into the Bio-Formats Java library from C++ in a cross-platform manner. As of this writing the bindings are functional with GCC on Linux and Mac OS X systems, as well as with Visual C++ 2005 and Visual C++ 2008 on Windows.

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1. [http://loci.wisc.edu/software/interfacing-non-java-code](http://loci.wisc.edu/software/interfacing-non-java-code)
2. [http://loci.wisc.edu/software/jar2lib](http://loci.wisc.edu/software/jar2lib)
5. [http://loci.wisc.edu/software/wiscscan](http://loci.wisc.edu/software/wiscscan)
14.3.1 Compile-time dependencies

To build the Bio-Formats C++ bindings from source, the following modules are required:

- **Apache Maven** is a software project management and comprehension tool. Along with Ant, it is one of the supported build systems for the Bio-Formats Java library, and is used to generate the Bio-Formats C++ bindings.

- **CMake** is a cross-platform, open source build system generator, commonly used to build C++ projects in a platform-independent manner. CMake supports GNU make as well as Microsoft Visual Studio, allowing the Bio-Formats C++ bindings to be compiled on Windows, Mac OS X, Linux and potentially other platforms.

- **Boost Thread** is a project providing open source portable C++ source libraries. It has become a suite of de facto standard libraries for C++. The Bio-Formats C++ bindings require the Boost Thread module in order to handle C++ threads in a platform independent way.

- **Java Development Kit** At runtime, only the Java Runtime Environment (JRE) is necessary to execute the Bio-Formats code. However, the full J2SE development kit is required at compile time on some platforms (Windows in particular), since it comes bundled with the JVM shared library (jvm.lib) necessary to link with Java.

For information on installing these dependencies, refer to the page for your specific platform: *Windows*, *Mac OS X*, *Linux*.

14.3.2 How to build

The process of building the Bio-Formats C++ bindings is divided into two steps:

1. Generate a C++ project consisting of “proxies” which wrap the Java code. This step utilizes the Maven project management tool, specifically a Maven plugin called cppwrap.

2. Compile this generated C++ project. This step utilizes the cross-platform CMake build system.

For details on executing these build steps, refer to the page for your specific platform: *Windows*, *Mac OS X*, *Linux*.

14.3.3 Build results

If all goes well, the build system will:

1. Generate the Bio-Formats C++ proxy classes;

2. Build the Jace C++ library;

3. Build the Java Tools C++ library;

4. Build the Bio-Formats C++ shared library;

5. Build the showinf and minimum_writer command line tools, for testing the functionality.

Please be patient, as the build may require several minutes to complete.

Afterwards, the dist/formats-bsd subdirectory will contain the following files:

1. libjace.so / libjace.jnilib / jace.dll : Jace shared library

2. libformats-bsd.so / libformats-bsd.dylib / formats-bsd.dll : C++ shared library for BSD-licensed readers and writers

3. jace-runtime.jar : Jace Java classes needed at runtime

4. bioformats_package.jar : Bio-Formats Java library needed at runtime

5. libjtools.so / libjtools.jnilib / jtools.dll : Java Tools shared library

6. showinf / showinf.exe : Example command line application

7. minimum_writer / minimum_writer.exe : Example command line application

Items 1-4 are necessary and required to deploy Bio-Formats with your C++ application. Item 5 (jtools) is a useful helper library for managing the Java virtual machine from C++, but is not strictly necessary to use Bio-Formats. All other files, including the example programs and various build files generated by CMake, are not needed.

If you prefer, instead of using the bioformats_package.jar bundle, you can provide individual JAR files as appropriate for your application. For details, see *using Bio-Formats as a Java library*. 
Please direct any questions to the OME team on the forums or mailing lists.

14.4 Building C++ bindings in Windows

14.4.1 Compile-time dependencies – Windows

Windows users will need to visit the appropriate web sites and download and install the relevant binaries for all the dependencies. To configure the tools, you will need to edit or create several environment variables on your system. Access them by clicking the “Environment Variables” button from Control Panel, System, Advanced tab. Use semicolons to separate multiple directories in the PATH variable.

14.4.2 Compile-time dependencies – Windows – Maven

Download Maven. Unpack the Maven archive into your Program Files, then add the folder’s bin subdirectory to your PATH environment variable; e.g.:

C:\Program Files\apache-maven-3.0.4\bin

Once set, new Command Prompts will recognize “mvn” as a valid command.

14.4.3 Compile-time dependencies – Windows – CMake

Download and run the CMake installer. During installation, select the “Add CMake to the system PATH for all users” option to ensure that Bio-Formats build system can find your CMake executable. Once installed, new Command Prompts will recognize “cmake” and “cmake-gui” as valid commands.

14.4.4 Compile-time dependencies – Windows – Boost

The easiest way to install the Boost Thread library on Windows is to use the free installer from BoostPro. When running the installer:

- Under “Compilers,” check the version of Visual C++ matching your system.
- Under “Variants,” check all eight boxes.
- When choosing components, check “Boost DateTime” and “Boost Thread.”

14.4.5 Compile-time dependencies – Windows – Java Development Kit

Download and install the JDK. After the installation is complete, create a new environment variable called JAVA_HOME pointing to your Java installation; e.g.:

C:\Program Files\Java\jdk1.6.0_25

Setting JAVA_HOME is the easiest way to ensure that Maven can locate Java. You will also need to append your JDK’s client or server VM folder to the PATH; e.g.:

%JAVA_HOME%\jre\bin\client

---

12http://www.openmicroscopy.org/community/
13http://lists.openmicroscopy.org.uk/mailman/listinfo/
14http://maven.apache.org/
15http://cmake.org/
16http://www.boostpro.com/download/
17http://www.oracle.com/technetwork/java/javase/downloads/
This step ensures that a directory containing jvm.dll is present in the PATH. If you do not perform this step, you will receive a runtime error when attempting to initialize a JVM from native code.

Optionally, you can add the bin subdirectory to the PATH; e.g.:

%JAVA_HOME%\bin

Once set, new Command Prompts will recognize (e.g.) “javac” as a valid command.

### 14.4.6 Compile-time dependencies – Windows – Visual C++

In addition to the other prerequisites, you will also need a working copy of Visual C++. We have tested compilation with Visual C++ 2005 Professional and Visual C++ 2008 Express; other versions may or may not work.

You can download Visual C++ Express for free[^18]. You must launch the environment at least once before you will be able to compile the Bio-Formats C++ bindings.

### 14.4.7 How to build - Windows

Run Command Prompt and change to your Bio-Formats working copy. Then run:

```bash
# generate the Bio-Formats C++ bindings
cd components\formats-bsd
mvn -DskipTests package dependency:copy-dependencies cppwrap:wrap

# build the Bio-Formats C++ bindings
cd target\cppwrap
mkdir build
cd build
cmake-gui ..
```

The CMake GUI will open. Click the Configure button, and a dialog will appear. Select your installed version of Visual Studio, and click Finish.

When configuring, you can use the J2L_WIN_BUILD_DEBUG flag to indicate if this will be a Debug or Release build. If the flag is checked it will build as Debug, unchecked will build as Release.

Once configuration is complete, click Configure again, repeating as necessary until the Generate button becomes available. Then click Generate. Once generation is complete, close the CMake window.

Back at the Command Prompt, type:

```bash
start jace.sln
```

The solution will then open in Visual Studio. Select Release or Debug as appropriate from the drop-down menu. Press F7 to compile (or select Build Solution from the Build menu).

### 14.5 Building C++ bindings in Mac OS X

#### 14.5.1 Compile-time dependencies – Mac OS X

To install dependencies on Mac OS X, we advise using Homebrew[^19]:

```bash
brew install maven cmake boost
```

Unless otherwise configured, this will install binaries into /usr/local/.

[^18]: http://www.microsoft.com/express/
[^19]: https://github.com/mxcl/homebrew/
14.5.2 How to build – Mac OS X

The following commands will generate and build the Bio-Formats C++ bindings:

```
# generate the C++ bindings
cd components/formats-bsd
mvn -DskipTests package dependency:copy-dependencies cppwrap:wrap

# compile the C++ bindings
cd target/cppwrap
mkdir build
cd build
cmake ..
make
```

14.6 Building C++ bindings in Linux

14.6.1 Compile-time dependencies – Linux

The following directions are specific to Ubuntu Linux. Other Linux distributions may have similar packages available; check your package manager.

To install dependencies on Ubuntu Linux, execute:

```
# install code generation prerequisites
sudo aptitude install maven2

# install build prerequisites
sudo aptitude install build-essential cmake libboost-thread-dev

# install Java Development Kit
sudo aptitude install sun-java6-jdk
sudo update-alternatives --config java
```

Then select Sun’s Java implementation as the system default.

It may be possible to use a different Java compiler (i.e., omit the sun-java6-jdk package and update-alternatives step), but we have only tested the compilation process with Sun’s Java compiler.

14.6.2 How to build – Linux

The following commands will generate and build the Bio-Formats C++ bindings:

```
# generate the Bio-Formats C++ bindings
cd components/formats-bsd
mvn -DskipTests package dependency:copy-dependencies cppwrap:wrap

# build the Bio-Formats C++ bindings
cd target/cppwrap
mkdir build
cd build
cmake ..
make
```
15.1 Bio-Formats file format reader guide

This document is a brief guide to writing new Bio-Formats file format readers.

All format readers should extend either `loci.formats.FormatReader` or a reader in `loci.formats.in`.

15.1.1 Methods to override

- **boolean isSingleFile(String id)**
  Whether or not the named file is expected to be the only file in the dataset. This only needs to be overridden for formats whose datasets can contain more than one file.

- **boolean isThisType(RandomAccessInputStream)**
  Check the first few bytes of a file to determine if the file can be read by this reader. You can assume that index 0 in the stream corresponds to the index 0 in the file. Return true if the file can be read; false if not (or if there is no way of checking).

- **int fileGroupOption(String id)**
  Returns an indication of whether or not the files in a multi-file dataset can be handled individually. The return value should be one of the following:
  - FormatTools.MUST_GROUP: the files cannot be handled separately
  - FormatTools.CAN_GROUP: the files may be handled separately or as a single unit
  - FormatTools.CANNOT_GROUP: the files must be handled separately
  This method only needs to be overridden for formats whose datasets can contain more than one file.

- **String[] getSeriesUsedFiles(boolean noPixels)**
  You only need to override this if your format uses multiple files in a single dataset. This method should return a list of all files associated with the given file name and the current series (i.e. every file needed to display the current series). If the noPixels flag is set, then none of the files returned should contain pixel data. For an example of how this works, see `loci.formats.in.PerkinElmerReader`. It is recommended that the first line of this method be FormatTools.assertId(currentId, true, 1) - this ensures that the file name is non-null.

- **byte[] openBytes(int, byte[], int, int, int, int)**
  Returns a byte array containing the pixel data for a subimage specified image from the given file. The dimensions of the subimage (upper left X coordinate, upper left Y coordinate, width, and height) are specified in the final four int parameters. This should throw a `FormatException` if the image number is invalid (less than 0 or >= the number of images). The ordering of the array returned by openBytes should correspond to the values returned by isLittleEndian() and isInterleaved(). Also, the length of the byte array should be [image width * image height * bytes per pixel]. Extra bytes will generally be truncated. It is recommended that the first line of this method be FormatTools.checkPlaneParameters(this, no, buf.length, x, y, w, h) - this ensures that all of the parameters are valid.

---

4. http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#isThisType(loci.common.RandomAccessInputStream)
8. http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#openBytes(int, byte[], int, int, int, int)
• **protected void initFile(String)**\(^9\) The majority of the file parsing logic should be placed in this method. The idea is to call this method once (and only once!) when the file is first opened. Generally, you will want to start by calling super.initFile(String). You will also need to set up the stream for reading the file, as well as initializing any dimension information and metadata. Most of this logic is up to you; however, you should populate the ‘core’ variable (see loci.formats.CoreMetadata\(^10\)).

Note that each variable is initialized to 0 or null when super.initFile(String) is called. Also, super.initFile(String) constructs a Hashtable called “metadata” where you should store any relevant metadata.

• **public void close(boolean fileOnly)**\(^11\) Cleans up any resources used by the reader. Global variables should be reset to their initial state, and any open files or delegate readers should be closed.

Note that if the new format is a variant of a format currently supported by Bio-Formats, it is more efficient to make the new reader a subclass of the existing reader (rather than subclassing FormatReader\(^12\)). In this case, it is usually sufficient to override initFile(String) and isThisType(byte[]).

Every reader also has an instance of loci.formats.CoreMetadata\(^13\). All readers should populate the fields in CoreMetadata, which are essential to reading image planes.

If you read from a file using something other than RandomAccessInputStream\(^14\) or Location\(^15\), you **must** use the file name returned by Location.getMappedId(String), not the file name passed to the reader. Thus, a stub for initFile(String) might look like this:

```java
protected void initFile(String id) throws FormatException, IOException {
    super.initFile(id);
    RandomAccessInputStream in = new RandomAccessInputStream(id);
    // alternatively,
    // FileInputStream in = new FileInputStream(Location.getMappedId(id));
    // read basic file structure and metadata from stream
}
```

For more details, see the **Bio-Formats Javadocs**\(^16\) for Location.mapId(String, String) and Location.getMappedId(String).

### 15.1.2 Variables to populate

There are a number of global variables defined in loci.formats.FormatReader\(^17\) that should be populated in the constructor of any implemented reader.

These variables are:

- **boolean suffixNecessary** Indicates whether or not a file name suffix is required; true by default
- **boolean suffixSufficient** Indicates whether or not a specific file name suffix guarantees that this reader can open a particular file; true by default
- **boolean hasCompanionFiles** Indicates whether or not there is at least one file in a dataset of this format that contains only metadata (no images); false by default
- **String datasetDescription** A brief description of the layout of files in datasets of this format; only necessary for multi-file datasets
- **String[] domains** An array of imaging domains for which this format is used. Domains are defined in loci.formats.FormatTools\(^18\).

---

\(^9\) [http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/FormatReader.html#initFile(java.lang.String)]

\(^10\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/CoreMetadata.java]

\(^11\) [http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/loci/formats/IFormatReader.html#close(boolean)]

\(^12\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/FormatReader.java]

\(^13\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/CoreMetadata.java]

\(^14\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-common/src/loci/common/RandomAccessInputStream.java]

\(^15\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-common/src/loci/common/Location.java]

\(^16\) [http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/]

\(^17\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/FormatReader.java]

\(^18\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/FormatTools.java]
15.1.3 Other useful things

- `loci.common.RandomAccessInputStream`\(^{19}\) is a hybrid RandomAccessFile/InputStream class that is generally more efficient than either RandomAccessFile or InputStream, and implements the DataInput interface. It is recommended that you use this for reading files.

- `loci.common.Location`\(^{20}\) provides an API similar to java.io.File, and supports File-like operations on URLs. It is highly recommended that you use this instead of File. See the Javadocs\(^{21}\) for additional information.

- `loci.common.DataTools`\(^{22}\) provides a number of methods for converting bytes to shorts, ints, longs, etc. It also supports reading most primitive types directly from a RandomAccessInputStream (or other DataInput implementation).

- `loci.formats.ImageTools`\(^{23}\) provides several methods for manipulating primitive type arrays that represent images. Consult the source or Javadocs for more information.

- If your reader relies on third-party code which may not be available to all users, it is strongly suggested that you make a corresponding service class that interfaces with the third-party code. Please see Bio-Formats service and dependency infrastructure for a description of the service infrastructure, as well as the `loci.formats.services` package\(^{24}\).

- Several common image compression types are supported through subclasses of `loci.formats.codec.BaseCodec`\(^{25}\). These include JPEG, LZW, LZO, Base64, ZIP and RLE (PackBits).

- If you wish to convert a file’s metadata to OME-XML (strongly encouraged), please see Bio-Formats metadata processing for further information.

- Utility methods for reading and writing individual bits from a byte array can be found in `loci.formats.codec.BitBuffer`\(^{26}\) and `loci.formats.codec.BitWriter`\(^{27}\).

- Once you have written your file format reader, add a line to the `readers.txt`\(^{28}\) file with the fully qualified name of the reader, followed by a “#” and the file extensions associated with the file format. Note that `ImageReader`\(^{29}\), the master file format reader, tries to identify which format reader to use according to the order given in `readers.txt`\(^{30}\), so be sure to place your reader in an appropriate position within the list.

- The easiest way to test your new reader is by calling “java loci.formats.tools.ImageInfo <file name>”. If all goes well, you should see all of the metadata and dimension information, along with a window showing the images in the file. `ImageReader`\(^{31}\) can take additional parameters; a brief listing is provided below for reference, but it is recommended that you take a look at the contents of `loci.formats.tools.ImageInfo`\(^{32}\) to see exactly what each one does.

---

\(^{19}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-common/src/loci/common/RandomAccessInputStream.java

\(^{20}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-common/src/loci/common/Location.java

\(^{21}\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/

\(^{22}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-common/src/loci/common/DataTools.java

\(^{23}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/ImageTools.java

\(^{24}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/services/

\(^{25}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/codec/BaseCodec.java

\(^{26}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/codec/BitBuffer.java

\(^{27}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/codec/BitWriter.java

\(^{28}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/readers.txt

\(^{29}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/ImageReader.java

\(^{30}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/readers.txt

\(^{31}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-api/src/loci/formats/ImageReader.java

\(^{32}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/bio-formats-tools/src/loci/formats/tools/ImageInfo.java
Table of Argument and Action:

<table>
<thead>
<tr>
<th>Argument</th>
<th>Action</th>
</tr>
</thead>
<tbody>
<tr>
<td>-version</td>
<td>print the library version and exit</td>
</tr>
<tr>
<td>file</td>
<td>the image file to read</td>
</tr>
<tr>
<td>-nopix</td>
<td>read metadata only, not pixels</td>
</tr>
<tr>
<td>-nocore</td>
<td>do not output core metadata</td>
</tr>
<tr>
<td>-nometa</td>
<td>do not parse format-specific metadata table</td>
</tr>
<tr>
<td>-nofilter</td>
<td>do not filter metadata fields</td>
</tr>
<tr>
<td>-thumbs</td>
<td>read thumbnails instead of normal pixels</td>
</tr>
<tr>
<td>-minmax</td>
<td>compute min/max statistics</td>
</tr>
<tr>
<td>-merge</td>
<td>combine separate channels into RGB image</td>
</tr>
<tr>
<td>-nogroup</td>
<td>force multi-file datasets to be read as individual files</td>
</tr>
<tr>
<td>-stitch</td>
<td>stitch files with similar names</td>
</tr>
<tr>
<td>-separate</td>
<td>split RGB image into separate channels</td>
</tr>
<tr>
<td>-expand</td>
<td>expand indexed color to RGB</td>
</tr>
<tr>
<td>-omexml</td>
<td>populate OME-XML metadata</td>
</tr>
<tr>
<td>-normalize</td>
<td>normalize floating point images*</td>
</tr>
<tr>
<td>-fast</td>
<td>paint RGB images as quickly as possible*</td>
</tr>
<tr>
<td>-debug</td>
<td>turn on debugging output</td>
</tr>
<tr>
<td>-range</td>
<td>specify range of planes to read (inclusive)</td>
</tr>
<tr>
<td>-series</td>
<td>specify which image series to read</td>
</tr>
<tr>
<td>-swap</td>
<td>override the default input dimension order</td>
</tr>
<tr>
<td>-shuffle</td>
<td>override the default output dimension order</td>
</tr>
<tr>
<td>-map</td>
<td>specify file on disk to which name should be mapped</td>
</tr>
<tr>
<td>-preload</td>
<td>pre-read entire file into a buffer; significantly reduces the time required to read the images, but requires more memory</td>
</tr>
<tr>
<td>-crop</td>
<td>crop images before displaying; argument is ‘x,y,w,h’</td>
</tr>
<tr>
<td>-autoscale</td>
<td>used in combination with ‘-fast’ to automatically adjust brightness and contrast</td>
</tr>
<tr>
<td>-novalid</td>
<td>do not perform validation of OME-XML</td>
</tr>
<tr>
<td>-omexml-only</td>
<td>only output the generated OME-XML</td>
</tr>
<tr>
<td>-format</td>
<td>read file with a particular reader (e.g., ZeissZVI)</td>
</tr>
</tbody>
</table>

* = may result in loss of precision

- If you wish to test using TestNG, `loci.tests.testng.FormatReaderTest`\(^{33}\) provides several basic tests that work with all Bio-Formats readers. See the `FormatReaderTest` source code for additional information.

- For more details, please look at the source code and Javadocs\(^{34}\). Studying existing readers is probably the best way to get a feel for the API; we would recommend first looking at `loci.formats.in.ImarisReader`\(^{35}\) (this is the most straightforward one). `loci.formats.in.LIFReader`\(^{36}\) and `InCellReader`\(^{37}\) are also good references that show off some of the nicer features of Bio-Formats.

If you have questions about Bio-Formats, please contact the OME team\(^{38}\).

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\(^{33}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/test-suite/src/loci/tests/testng/FormatReaderTest.java

\(^{34}\)http://downloads.openmicroscopy.org/latest/bio-formats5.0/api/

\(^{35}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImarisReader.java

\(^{36}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LIFReader.java

\(^{37}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/InCellReader.java

\(^{38}\)http://www.openmicroscopy.org/site/community
16.1 Testing individual commits (internal developers)

At the bottom of many commit messages in https://github.com/openmicroscopy/bioformats, you will find a few lines similar to this:

To test, please run:

$ ant -Dtestng.directory=$DATA/metamorph test-automated

This shows the command(s) necessary to run automated tests against the files likely to be affected by that commit. If you want to run these tests, you will need to do the following:

Clone bioformats.git and checkout the appropriate branch (by following the directions on the Git usage\(^1\) page). Run this command to build all of the JAR files:

$ ant clean jars

Switch to the test-suite component:

$ cd components/test-suite

Run the tests, where $DATA is the path to the full data repository:

$ ant -Dtestng.directory=$DATA/metamorph test-automated

By default, 512 MB of memory are allocated to the JVM. You can increase this by adding the ‘-Dtestng.memory=XXXm’ option. You should now see output similar to this:

Buildfile: build.xml

init-title:
    [echo] ---------------------- bio-formats-testing-framework ----------------------

init-timestamp:
release-version:
init-manifest-cp:
init:
copy-source:

\(^1\)http://www.openmicroscopy.org/site/support/contributing/using-git.html
compile:

test-automated:
  [testng] [Parser] Running:
  [testng] Bio-Formats software test suite
  [testng] Scanning for files...
  [testng] Building list of tests...
  [testng] Ready to test 490 files
  [testng] ........................................

and then eventually:

  [testng] ==================================================
  [testng] Bio-Formats software test suite
  [testng] Total tests run: 19110, Failures: 0, Skips: 0
  [testng] ==============================================================

BUILD SUCCESSFUL
Total time: 16 minutes 42 seconds

Each of the dots represents a single passed test; a ‘-‘ is a skipped test, and an ‘F‘ is a failed test. This is mostly just for your amusement if you happen to be staring at the console while the tests run, as a more detailed report is logged to bio-formats-software-test-$DATE.log (where “$DATE” is the date on which the tests started in “yyyy-MM-dd_hh-mm-ss” format).

If Ant reports that the build was successful, then there is nothing that you need to do. Otherwise, it is helpful if you can provide the command, branch name, number of failures at the bottom of the Ant output, and the bio-formats-software-test-*log file.

### 16.2 Public test data

Most of the data-driven tests would benefit from having a comprehensive set of public sample data (see also #4086²).

Formats for which we already have public sample data:

- ICS (*)
- Leica LEI
- IPLab
- BMP (*)
- Image-Pro SEQ
- QuickTime (*)
- Bio-Rad PIC
- Image-Pro Workspace
- Fluoview/ABD TIFF (*)
- Perkin Elmer Ultraview
- Gatan DM3
- Zeiss LSM
- Openlab LIFF (*)
- Leica LIF (*)

²http://trac.openmicroscopy.org.uk/ome/ticket/4086
• TIFF (*)
• Khoros (http://netghost.narod.ru/gff/sample/images/viff/index.htm)
• MNG (Download3) (*)

Formats for which we can definitely generate public sample data:
• PNG/APNG
• JPEG
• PGM
• FITS
• PCX
• GIF
• Openlab Raw
• OME-XML
• OME-TIFF
• AVI
• PICT
• LIM
• PSD
• Targa
• Bio-Rad Gel
• Fake
• ECAT-7 (minctoecat)
• NRRD
• JPEG-2000
• Micromanager
• Text
• DICOM
• MINC (rawtominc)
• NIFTI (dicomnifti)
• Analyze 7.5 (medcon)
• SDT
• FV1000 .oib/.oif
• Zeiss ZVI
• Leica TCS
• Aperio SVS
• Imaris (raw)

Formats for which I need to check whether or not we can generate public sample data:
• IPLab Mac (Ivision)
• Delatavision
• MRC
• Gatan DM2

- Imaris (HDF)
- EPS
- Alicona AL3D
- Visitech
- InCell
- L2D
- FEI
- NAF
- MRW
- ARF
- LI-FLIM
- Oxford Instruments
- VG-SAM
- Hamamatsu HIS
- WA-TOP
- Seiko
- TopoMetrix
- UBM
- Quesant
- RHK
- Molecular Imaging
- JEOL
- Amira
- Unisoku
- Perkin Elmer Densitometer
- Nikon ND2
- SimplePCI .cxd
- Imaris (TIFF)
- Molecular Devices Gel
- Imacon .fff
- LEO
- JPK
- Nikon NEF
- Nikon TIFF
- Prairie
- Metamorph TIFF/STK/ND
- Improvision TIFF
- Photoshop TIFF
- FEI TIFF
- SimplePCI TIFF
- Burleigh
Formats for which we definitely cannot generate public sample data:

• SM-Camera
• SBIG

16.3 Bio-Formats service and dependency infrastructure

16.3.1 Description

The Bio-Formats service infrastructure is an interface driven pattern for dealing with external and internal dependencies. The design goal was mainly to avoid the cumbersome usage of ReflectedUniverse where possible and to clearly define both service dependency and interface between components. This is generally referred to as dependency injection\(^4\), dependency inversion\(^5\) or component based design\(^6\).

It was decided, at this point, to forgo the usage of potentially more powerful but also more complicated solutions such as:

• Spring (http://spring.io)
• Guice (http://code.google.com/p/google-guice/)
• ...

The Wikipedia page for dependency injection\(^7\) contains many other implementations in many languages.

An added benefit is the potential code reuse possibilities as a result of decoupling of dependency and usage in Bio-Formats readers. Implementations of the initial Bio-Formats services were completed as part of BioFormatsCleanup and tickets \#463\(^8\) and \#464\(^9\).

16.3.2 Writing a service

• Interface – The basic form of a service is an interface which inherits from loci.common.services.Service\(^{10}\). Here is a very basic example using the (now removed) OMENotesService

```java
public interface OMENotesService extends Service {

    /**
     * Creates a new OME Notes instance.
     * @param filename Path to the file to create a Notes instance for.
     */
    public void newNotes(String filename);
}
```

\(^4\)http://en.wikipedia.org/wiki/Dependency_injection
\(^6\)http://en.wikipedia.org/wiki/Component-based_software_engineering
\(^7\)http://en.wikipedia.org/wiki/Dependency_injection
\(^8\)http://trac.openmicroscopy.org.uk/ome/ticket/463
\(^9\)http://trac.openmicroscopy.org.uk/ome/ticket/464
\(^{10}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-common/src/loci/common/services/Service.java
• **Implementation** – This service then has an implementation, which is usually located in the Bio-Formats component or package which imports classes from an external, dynamic or other dependency. Again looking at the `OMENotesService`:

```java
public class OMENotesServiceImpl extends AbstractService
    implements OMENotesService {
    /**
     * Default constructor.
     */
    public OMENotesServiceImpl() {
        checkClassDependency(Notes.class);
    }

    /* (non-Javadoc)
     * @see loci.formats.dependency.OMENotesService#newNotes()
     */
    public void newNotes(String filename) {
        new Notes(null, filename);
    }
}
```

• **Style**

  – Extension of AbstractService to enable uniform runtime dependency checking is recommended. Java does not check class dependencies until classes are first instantiated so if you do not do this, you may end up with Class-NotFound or the like exceptions being emitted from your service methods. This is to be strongly discouraged. If a service has unresolvable classes on its CLASSPATH instantiation should fail, not service method invocation.

  – Service methods should not burden the implementer with numerous checked exceptions. Also external dependency exception instances should not be allowed to directly leak from a service interface. Please wrap these using a ServiceException.

  – By convention both the interface and implementation are expected to be in a package named loci.*.services. This is not a hard requirement but should be followed where possible.

• **Registration** – A service’s interface and implementation must finally be registered with the `loci.common.services.ServiceFactory` via the `services.properties` file. Following the `OMENotesService` again, here is an example registration:

```properties
# OME notes service (implementation in legacy ome-notes component)
loci.common.services.OMENotesService=loci.ome.notes.services.OMENotesServiceImpl
```

### 16.3.3 Using a service

```java
OMENotesService service = null;
try {
    ServiceFactory factory = new ServiceFactory();
    service = factory.getInstance(OMENotesService.class);
} catch (DependencyException de) {
    LOGGER.info("", de);
}
```

---

16.4 Code generation with xsd-fu

xsd-fu is a Python application designed to digest OME XML schema and produce an object oriented Java infrastructure to ease work with an XML DOM tree. It is usually run automatically when building from source (see Building from source) and so running it by hand should not be needed. xsd-fu is primarily used to generate the OME-XML model objects, enums and enum handlers, plus the MetadataStore and MetadataRetrieve interfaces and implementations.

16.4.1 Running the code generator

$ cd components/xsd-fu
$ ./xsd-fu
Output directory must be specified!
Usage: ./xsd-fu [-n xsd_namespace] [-p package] -o <output_dir> -l <lang> <path/to/ome.xsd>...
Generates Java classes from an OME XML Schema definition.

Default package: "ome.xml.model"
Default namespace: "xsd:",

Examples:
./xsd-fu -n 'xs:' -p ome.xml -o ome/xml schemas/ome.xsd

Report bugs to OME Devel <ome-devel@lists.openmicroscopy.org.uk>

$ cd components/xsd-fu
$ ./xsd-fu -o ../../
Missing subcommand!
Usage: ./xsd-fu <subcommand> ...
Executes an OME-XML Schema definition parsing and code generation subcommand.

Available subcommands:
  java_classes
  omexml_metadata
  omero_metadata
  omero_model
  metadata_store
  metadata_retrieve
  metadata_aggregate
  dummy_metadata
  filter_metadata
  enum_types
  enum_handlers
  doc_gen
  tab_gen
debug

Report bugs to OME Devel <ome-devel@lists.openmicroscopy.org.uk>

16.4.2 Generating the OME-XML Java classes

The following sections outline how to generate parts of the OME-XML Java toolchain which are composed of:

- OME model objects
• Enumerations for OME model properties
• Enumeration handlers for regular expression matching of enumeration strings
• Metadata store and Metadata retrieve interfaces for all OME model properties
• Various implementations of Metadata store and/or Metadata retrieve interfaces

All of the above can be generated by this Ant command:

```bash
$ cd components/ome-xml
$ ant generate-source
```

Run:

```bash
$ ant generate-source -v
```

to see the command-line options used.

### 16.4.3 Working with Enumerations and Enumeration Handlers

XsdFu code generates enumeration regular expressions using a flexible configuration file\(^{13}\). Each enumeration has a key-value listing of regular expression to exact enumeration value matches. For example:

```plaintext
[Correction]
".*Pl.*Apo.*" = "PlanApo"
".*Pl.*Flu.*" = "PlanFluor"
"s*Vio.*Corr.*" = "VioletCorrected"
"S.*Flu.*" = "SuperFluor"
"Neo.*flu.*" = "NeoFluar"
".*Flu.*tar.*" = "Fluotar"
"Fluo.*" = "Fluor"
".*Flua.*" = "Fluar"
"s*Apo.*" = "Apo"
```

### 16.4.4 Generate OMERO model specification files

Run `xsd-fu` with the `omero_model` subcommand.

### 16.4.5 Special Thanks

A special thanks goes out to Dave Kuhlman\(^{14}\) for his fabulous work on `generateDS`\(^{15}\) which `xsd-fu` makes heavy use of internally. See open Trac tickets for Bio-Formats\(^{16}\) for information on work currently planned or in progress.

For more general guidance about how to contribute to OME projects, see the Contributing developers documentation\(^{17}\).

---

\(^{13}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/xsd-fu/cfg/enum_handler.cfg

\(^{14}\)http://www.davekuhlman.org/

\(^{15}\)http://www.davekuhlman.org/generateDS.html

\(^{16}\)https://trac.openmicroscopy.org.uk/ome/report/44

\(^{17}\)http://www.openmicroscopy.org/site/support/contributing/index.html
Part IV

Formats
Bio-Formats supports over 120 different file formats. The Dataset Structure Table explains the file extension you should choose to open/import a dataset in any of these formats, while the Supported Formats table lists all of the formats and gives an indication of how well they are supported and whether Bio-Formats can write, as well as read, each format. The Summary of supported metadata fields table shows an overview of the OME data model fields populated for each format.

We are always looking for examples of files to help us provide better support for different formats. If you would like to help, you can upload files using our QA system uploader18. If you have any questions, or would prefer not to use QA, please email the ome-users mailing list19. If your format is already supported, please refer to the ‘we would like to have’ section on the individual page for that format, to see if your dataset would be useful to us.

18http://qa.openmicroscopy.org.uk/qa/upload/
19http://www.openmicroscopy.org/site/community/mailing-lists
This table shows the extension of the file that you should choose if you want to open/import a dataset in a particular format.

<table>
<thead>
<tr>
<th>Format name</th>
<th>File to choose</th>
<th>Structure of files</th>
</tr>
</thead>
<tbody>
<tr>
<td>AIM</td>
<td>.aim</td>
<td>Single file</td>
</tr>
<tr>
<td>ARF</td>
<td>.arf</td>
<td>Single file</td>
</tr>
<tr>
<td>Adobe Photoshop</td>
<td>.psd</td>
<td>Single file</td>
</tr>
<tr>
<td>Adobe Photoshop TIFF</td>
<td>.tif, .tiff</td>
<td>Single file</td>
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<tr>
<td>Alicona AL3D</td>
<td>.al3d</td>
<td>Single file</td>
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<tr>
<td>Amersham Biosciences GEL</td>
<td>.gel</td>
<td>Single file</td>
</tr>
<tr>
<td>Amira</td>
<td>.am, .amiramesh, .grey, .hx, .labels</td>
<td>Single file</td>
</tr>
<tr>
<td>Analyze 7.5</td>
<td>.img, .hdr</td>
<td>One .img file and one similarly-named .hdr file</td>
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<tr>
<td>Andor SIF</td>
<td>.sif</td>
<td>Single file</td>
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<td>Single file</td>
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<tr>
<td>Aperio SVS</td>
<td>.svs</td>
<td>Single file</td>
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<tr>
<td>Audio Video Interleave</td>
<td>.avi</td>
<td>Single file</td>
</tr>
<tr>
<td>BD Pathway</td>
<td>.exp, .tif</td>
<td>Multiple files (.exp, .dye, .ltp, …) plus one or more directories containing .tif and .bmp files</td>
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<td>Bio-Rad GEL</td>
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<td>Single file</td>
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<tr>
<td>Bio-Rad PIC</td>
<td>.pic,.xml,.raw</td>
<td>One or more .pic files and an optional lse.xml file</td>
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<tr>
<td>Bitplane Imaris 5.5 (HDF)</td>
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<td>Canon RAW</td>
<td>.cr2, .crw, .jpg, .thm, .wav</td>
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<td>One .vsi file and an optional directory with a similar name that contains at least one subdirectory with .ets files</td>
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<td>.c01,.dib</td>
<td>One or more .c01 files</td>
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<tr>
<td>Compix Simple-PCI</td>
<td>.cxd</td>
<td>Single file</td>
</tr>
<tr>
<td>DICOM</td>
<td>.dic, .dcm, .dicom, .jp2, .j2ki, .j2kr, .raw, .ima</td>
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<td>Deltavision</td>
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<td>One .dv, .r3d, or .d3d file and up to two optional .log files</td>
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<td>Single file</td>
</tr>
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<td>Single file</td>
</tr>
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<td>Format name</td>
<td>File to choose</td>
<td>Structure of files</td>
</tr>
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<td>----------------</td>
<td>---------------------------------------------------------------------------------------------------------------------------------------------------</td>
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<td>FEI/Philips</td>
<td>.img</td>
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<td>Flexible Image Transport System</td>
<td>.fits, .fts</td>
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<td>Gatan Digital Micrograph</td>
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<td>One .txt file plus one similarly-named .tif, .bmp, or .jpg file</td>
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<td>One .hed file plus one similarly-named .img file</td>
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<td>IPI Lab</td>
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<td>IVision</td>
<td>.ipm</td>
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<td>Image Cytometry Standard</td>
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<td>Image-Pro Sequence</td>
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<td>Single file</td>
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<tr>
<td>Image-Pro Workspace</td>
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<td>Improvision TIFF</td>
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<td>.xdce, .xml, .tiff, .tif, .xlog</td>
<td>One .xdce file with at least one .tif/.tiff or .im file</td>
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<td>.jp2, .j2k, .jpf</td>
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<td>JPX Instruments</td>
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<td>JPX</td>
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<td>Kodak Molecular Imaging</td>
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<td>LEO</td>
<td>.sxm, .tif, .tiff</td>
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<td>LI-FLIM</td>
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<td>Single file</td>
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<tr>
<td>Laboratory Imaging</td>
<td>.lim</td>
<td>Single file</td>
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<tr>
<td>Leica</td>
<td>.lei, .tif, .tiff, .raw</td>
<td>One .lei file with at least one .tif/.tiff file and an optional .txt file</td>
</tr>
<tr>
<td>Leica Image File Format</td>
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<td>Single file</td>
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<td>Leica SCN</td>
<td>.scn</td>
<td>Single file</td>
</tr>
<tr>
<td>Leica TCS TIFF</td>
<td>.tif, .tiff, .xml</td>
<td>Single file</td>
</tr>
<tr>
<td>Li-Cor L2D</td>
<td>.l2d, .scn, .tif</td>
<td>One .l2d file with one or more directories containing .tif/.tiff files</td>
</tr>
<tr>
<td>MIAS</td>
<td>.tif, .tiff, .txt</td>
<td>One directory per plate containing one directory per well, each with one or more .tif/.tiff files</td>
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<td>MINC MRI</td>
<td>.mnc</td>
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<tr>
<td>Medical Research Council</td>
<td>.mrc, .st, .ali, .map, .rec</td>
<td>Single file</td>
</tr>
<tr>
<td>Metamorph STK</td>
<td>.stk, .nd, .tif, .tiff</td>
<td>One or more .stk or .tif/.tiff files plus an optional .nd file</td>
</tr>
<tr>
<td>Metamorph TIFF</td>
<td>.tif, .tiff</td>
<td>One or more .tif/.tiff files</td>
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Continued on next page
<table>
<thead>
<tr>
<th>Format name</th>
<th>File to choose</th>
<th>Structure of files</th>
</tr>
</thead>
<tbody>
<tr>
<td>Micro-Manager</td>
<td>.tif, .tiff, .txt, .xml</td>
<td>A ‘metadata.txt’ file plus or more .tif files</td>
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<td>Minolta MRW</td>
<td>.mrw</td>
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<td>Molecular Imaging</td>
<td>.stc</td>
<td>Single file</td>
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<tr>
<td>Multiple Network Graphics</td>
<td>.mng</td>
<td>Single file</td>
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<tr>
<td>NITF</td>
<td>.nii, .img, .hdr</td>
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<td>NRRD</td>
<td>.nrrd, .nhdr</td>
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<td>Single file</td>
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<td>Nikon ND2</td>
<td>.nd2</td>
<td>Single file</td>
</tr>
<tr>
<td>Nikon NEF</td>
<td>.nef, .tiff</td>
<td>Single file</td>
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<tr>
<td>Nikon TIFF</td>
<td>.tiff, .tiff</td>
<td>Single file</td>
</tr>
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<td>OME-TIFF</td>
<td>.ome, .tiff, .ome.tiff</td>
<td>One or more .ome.tiff files</td>
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<td>OME-XML</td>
<td>.ome</td>
<td>Single file</td>
</tr>
<tr>
<td>Olympus APL</td>
<td>.apl, .tnb, .mtb, .tif</td>
<td>One .apl file, one .mtb file, one .tab file and a directory containing one or more .tif files</td>
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<tr>
<td>Olympus FV1000</td>
<td>.oib, .oif, .ptf, .lutf</td>
<td>Single .oib file or one .oif file and a similarly-named directory containing .tif/.tif files</td>
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<td>Olympus Fluoview/ABD TIFF</td>
<td>.tif, .tiff</td>
<td>One or more .tif/.tif files, and an optional .txt file</td>
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<td>Olympus SIS TIFF</td>
<td>.tif, .tiff</td>
<td>Single file</td>
</tr>
<tr>
<td>Olympus ScanR</td>
<td>.dat, .xml, .tiff</td>
<td>One .xml file, one ‘data’ directory containing .tif/.tif files, and optionally two .dat files</td>
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<tr>
<td>Oxford Instruments</td>
<td>.top</td>
<td>Single file</td>
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<tr>
<td>PCX</td>
<td>.pcx</td>
<td>Single file</td>
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<tr>
<td>PICT</td>
<td>.pict, .pct</td>
<td>Single file</td>
</tr>
<tr>
<td>POV-Ray</td>
<td>.df3</td>
<td>Single file</td>
</tr>
<tr>
<td>Perkin Elmer Densitometer</td>
<td>.hdr, .img</td>
<td>One .hdr file and a similarly-named .img file</td>
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<tr>
<td>PerkinElmer</td>
<td>.ano, .cfg, .csv, .htm, .rec, .tim, .zpo, .tiff</td>
<td>One .htm file, several other metadata files (.tim, .ano, .csv, …) and either .tif files or .2, .3, .4, etc. files</td>
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<tr>
<td>PerkinElmer Operetta</td>
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<td>Directory with XML file and one .tif/.tif file per plane</td>
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<td>Portable Gray Map</td>
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<td>One .xml file, one .cfg file, and one or more .tif/.tif files</td>
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<td>Pyramid TIFF</td>
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<td>QuickTime</td>
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<td>Single file</td>
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<td>RHK Technologies</td>
<td>.sm2, .sm3</td>
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<tr>
<td>SBIG</td>
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<td>SM Camera</td>
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<tr>
<td>SPC Image Data</td>
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<td>SPIDER</td>
<td>.spi</td>
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<tr>
<td>Seiko</td>
<td>.xqd, .xqf</td>
<td>Single file</td>
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<tr>
<td>SimplePCI TIFF</td>
<td>.tif, .tiff</td>
<td>Single file</td>
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<td>Simulated data</td>
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<td>Tagged Image File Format</td>
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<td>Single file</td>
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<tr>
<td>Text</td>
<td>.txt, .csv</td>
<td>Single file</td>
</tr>
<tr>
<td>TillVision</td>
<td>.vws, .psl, .inf</td>
<td>One .vws file and possibly one similarly-named directory</td>
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<td>TopoMetrix</td>
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<td>Single file</td>
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<tr>
<td>UBM</td>
<td>.pr3</td>
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<tr>
<td>Unisoku STM</td>
<td>.hdr, .dat</td>
<td>One .HDR file plus one similarly-named .DAT file</td>
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</table>

Continued on next page
Table 17.1 – continued from previous page

<table>
<thead>
<tr>
<th>Format name</th>
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<th>Structure of files</th>
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<td>Varian FDF</td>
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<td>Single file</td>
</tr>
<tr>
<td>Visitech XYS</td>
<td>.xys, .html</td>
<td>One .html file plus one or more .xys files</td>
</tr>
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<td>.mvd2, .aifs, .aix, .dat, .atsf</td>
<td>One .mvd2 file plus a ‘Data’ directory</td>
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<td>Volocity Library Clipping</td>
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<td>Single file</td>
</tr>
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<td>WA Technology TOP</td>
<td>.wat</td>
<td>Single file</td>
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<tr>
<td>Windows Bitmap</td>
<td>.bmp</td>
<td>Single file</td>
</tr>
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<td>Zeiss AxiophyVision TIFF</td>
<td>.tif, .xml</td>
<td>Single file</td>
</tr>
<tr>
<td>Zeiss CZI</td>
<td>.czi</td>
<td>Single file</td>
</tr>
<tr>
<td>Zeiss Laser-Scanning Microscopy</td>
<td>.ls, .mdb</td>
<td>One or more .ls files; if multiple .ls files are present, an .mdb file should also be present</td>
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<td>Zeiss Vision Image (ZVI)</td>
<td>.zvi</td>
<td>Single file</td>
</tr>
<tr>
<td>Zip</td>
<td>.zip</td>
<td>Single file</td>
</tr>
</tbody>
</table>

17.1 Flex Support

OMERO importers supports importing analyzed Flex files from an Opera system.

Basic configuration is done via the importer.ini. Once the user has run the Importer once, this file will be in the following location:

- C:\Documents and Settings\<username>\omero\importer.ini

The user will need to modify or add the [FlexReaderServerMaps] section of the INI file as follows:

```ini
... [FlexReaderServerMaps]
CIA-1 = \\hostname1\mount;\\archivehost1\mount
CIA-2 = \\hostname2\mount;\\archivehost2\mount
```

where the key of the INI file line is the value of the “Host” tag in the .mea measurement XML file (here: <Host name="CIA-1">) and the value is a semicolon-separated list of escaped UNC path names to the Opera workstations where the Flex files reside.

Once this resolution has been encoded in the configuration file and you have restarted the importer, you will be able to select the .mea measurement XML file from the Importer user interface as the import target.
# SUPPORTED FORMATS

## Ratings legend and definitions

<table>
<thead>
<tr>
<th>Format</th>
<th>Extensions</th>
<th>Pixels</th>
<th>Metadata</th>
<th>Openness</th>
<th>Presence</th>
<th>Utility</th>
<th>Export</th>
<th>BSD</th>
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<td>3i SlideBook</td>
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<td></td>
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<td>Andor Bio-Imaging</td>
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<tr>
<td>Division (ABD) TIFF</td>
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<td>OME-TIFF</td>
<td>.ome.tif³</td>
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<td>.pcraw, .rec</td>
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<th>Pixels</th>
<th>Metadata</th>
<th>Openness</th>
<th>Presence</th>
<th>Utility</th>
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<td>WA-TOP</td>
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<td>Windows Bitmap</td>
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<td>Zeiss AxioVision</td>
<td>.xml, .tiff</td>
<td>🔺</td>
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<tr>
<td>Zeiss AxiosVision ZVI (Zeiss Vision Image)</td>
<td>.czi³</td>
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<td>.lsm, .mdb</td>
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</table>

Bio-Formats currently supports **136** formats

<table>
<thead>
<tr>
<th>Ratings legend and definitions</th>
</tr>
</thead>
<tbody>
<tr>
<td>🔺</td>
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</tbody>
</table>

³http://www.zeiss.com/czi
Pixels  Our estimation of Bio-Formats’ ability to reliably extract complete and accurate pixel values from files in that format. The better this score, the more confident we are that Bio-Formats will successfully read your file without displaying an error message or displaying an erroneous image.

Metadata  Our certainty in the thoroughness and correctness of Bio-Formats’ metadata extraction and conversion from files of that format into standard OME-XML. The better this score, the more confident we are that all meaningful metadata will be parsed and populated as OME-XML.

Openness  This is not a direct expression of Bio-Formats’ performance, but rather indicates the level of cooperation the format’s controlling interest has demonstrated toward the scientific community with respect to the format. The better this score, the more tools (specification documents, source code, sample files, etc.) have been made available.

Presence  This is also not directly related to Bio-Formats, but instead represents our understanding of the format’s popularity, and is also as a measure of compatibility between applications. The better this score, the more common the format and the more software packages include support for it.

Utility  Our opinion of the format’s suitability for storing metadata-rich microscopy image data. The better this score, the wider the variety of information that can be effectively stored in the format.

Export  This indicates whether Bio-Formats is capable of writing the format (Bio-Formats can read every format on this list).

BSD  This indicates whether format is BSD-licensed. By default, format readers and writers are GPL-licensed.

### 18.1 3i SlideBook

Extensions: .sld

Developer: Intelligent Imaging Innovations

Owner: Intelligent Imaging Innovations

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions: 4.1, 4.2

Supported Metadata Fields: 3i SlideBook

We currently have:

• Numerous SlideBook datasets

We would like to have:

• A SlideBook specification document

• More SlideBook datasets (preferably acquired with the most recent SlideBook software)

Ratings

Pixels: 🔺

Metadata: 🔺

Openness: 🔺

Presence: 🔺

Utility: 🔺

Additional Information

Source Code: SlidebookReader.java

Notes:

4http://www.intelligent-imaging.com/

5http://www.intelligent-imaging.com/

6https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/SlidebookReader.java
We strongly encourage users to export their .sld files to OME-TIFF using the SlideBook software. Bio-Formats is not likely to support the full range of metadata that is included in .sld files, and so exporting to OME-TIFF from SlideBook is the best way to ensure that all metadata is preserved.

See also:
Slidebook software overview

### 18.2 Andor Bio-Imaging Division (ABD) TIFF

**Extensions:** .tif

**Developer:** Andor Bioimaging Department

**Owner:** Andor Technology

**Support**

- **BSD-licensed:** ✗
- **Export:** ✗

**Officially Supported Versions:**

**Supported Metadata Fields:** *Andor Bio-Imaging Division (ABD) TIFF*

We currently have:

- an ABD-TIFF specification document (from 2005 November, in PDF)
- a few ABD-TIFF datasets

We would like to have:

**Ratings**

- **Pixels:** 🔺
- **Metadata:** 🔺
- **Openness:** 🔺
- **Presence:** 🔻
- **Utility:** 🔺

**Additional Information**

**Source Code:** FluoviewReader.java

**Notes:**

Please note that while we have specification documents for this format, we are not able to distribute them to third parties. With a few minor exceptions, the ABD-TIFF format is identical to the Fluoview TIFF format.

### 18.3 AIM

**Extensions:** .aim

**Developer:** SCANCO Medical AG

**Support**

- **BSD-licensed:** ✗
Export: ✗
Officially Supported Versions:
Supported Metadata Fields: **AIM**
We currently have:
  • one .aim file
We would like to have:
  • an .aim specification document
  • more .aim files

**Ratings**

<table>
<thead>
<tr>
<th>Pixels:</th>
<th>Metadata:</th>
<th>Openness:</th>
<th>Presence:</th>
<th>Utility:</th>
</tr>
</thead>
</table>

**Additional Information**

Source Code: [AIMReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/AIMReader.java)

Notes:

### 18.4 Alicona 3D

Extensions: .al3d

Owner: [Alicona Imaging](http://www.alicona.com/)

**Support**

BSD-licensed: ✗

Export: ✗

Officially Supported Versions: 1.0

Supported Metadata Fields: **Alicona 3D**

We currently have:
  • an *AL3D specification document* (v1.0, from 2003, in PDF)
  • a few AL3D datasets
We would like to have:
  • more AL3D datasets (Z series, T series, 16-bit)

**Ratings**

<table>
<thead>
<tr>
<th>Pixels:</th>
<th>Metadata:</th>
<th>Openness:</th>
<th>Presence:</th>
</tr>
</thead>
</table>

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Utility:

Additional Information

Source Code: AliconaReader.java

Notes:

Known deficiencies:

- Support for 16-bit AL3D images is present, but has never been tested.
- Texture data is currently ignored.

18.5 Amersham Biosciences Gel

Extensions: .gel

Developer: Molecular Dynamics

Owner: GE Healthcare Life Sciences

Support

BSD-licensed: ✗

Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Amersham Biosciences Gel

We currently have:

- a GEL specification document (Revision 2, from 2001 Mar 15, in PDF)
- a few GEL datasets

We would like to have:

Ratings

Pixels: ▲

Metadata: ▲

Openness: ▼

Presence: ▼

Utility: ▼

Additional Information

Source Code: GelReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

See also:

GEL Technical Overview

---

14 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/AliconaReader.java
15 http://www.gelifesciences.com/
16 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/GelReader.java
17 http://www.awaresystems.be/imaging/tiff/tifftags/docs/gel.html
18.6 Amira Mesh

Extensions: .am, .amiramesh, .grey, .hx, .labels
Developer: Visage Imaging\textsuperscript{18}

Support
BSD-licensed: \xmark
Export: \xmark

Officially Supported Versions:
Supported Metadata Fields: \textit{Amira Mesh}

We currently have:
- a few Amira Mesh datasets

We would like to have:
- more Amira Mesh datasets

Ratings

Pixels: \▲
Metadata: \square
Openness: \▼
Presence: \▼
Utility: \▼

Additional Information
Source Code: \texttt{AmiraReader.java}\textsuperscript{19}

Notes:

18.7 Analyze 7.5

Extensions: .img, .hdr
Developer: Mayo Foundation Biomedical Imaging Resource\textsuperscript{20}

Support
BSD-licensed: \xmark
Export: \xmark

Officially Supported Versions:
Supported Metadata Fields: \textit{Analyze 7.5}

We currently have:
- an Analyze 7.5 specification document\textsuperscript{21}
- several Analyze 7.5 datasets

\textsuperscript{18}\url{http://www.amiravis.com/}
\textsuperscript{19}\url{https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/AmiraReader.java}
\textsuperscript{20}\url{http://www.mayo.edu/bir}
\textsuperscript{21}\url{http://analyzedirect.com/support/10.0/Documents/Analyze_Resource_01.pdf}
We would like to have:

**Ratings**

Pixels: 🔹
Metadata: 🔹
Openness: 🔹
Presence: 🔹
Utility: ◼

**Additional Information**

Source Code: [AnalyzeReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/AnalyzeReader.java)

Notes:

### 18.8 Animated PNG

Extensions: .png

Developer: [The Animated PNG Project](http://www.animatedpng.com/)

**Support**

BSD-licensed: ✔

Export: ✔

**Officially Supported Versions:**

Supported Metadata Fields: *Animated PNG*

Freely Available Software:

- Firefox 3+²⁴
- Opera 9.5+²⁵
- KSquirrel²⁶

We currently have:

- a specification document²⁷
- several APNG files

We would like to have:

**Ratings**

Pixels: 🔹
Metadata: 🔹
Openness: 🔹
Presence: 🔹
Utility: ◼

**Additional Information**

²²[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/AnalyzeReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/AnalyzeReader.java)


²⁴[http://www.mozilla.com/firefox](http://www.mozilla.com/firefox)

²⁵[http://www.opera.com/download](http://www.opera.com/download)


18.9 Aperio AFI

Extensions: .afi, .svs
Owner: Aperio

Support
BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: Aperio AFI
We currently have:
• several AFI datasets

We would like to have:

Ratings
Pixels: 
Metadata: 
Openness: 

Presence: 
Utility: 

Additional Information
Source Code: AFIReader.java

Notes:

See also:
Aperio ImageScope

18.10 Aperio SVS TIFF

Extensions: .svs
Owner: Aperio

Support
BSD-licensed: 
Export: 

Officially Supported Versions: 8.0, 8.2, 9.0
Supported Metadata Fields: Aperio SVS TIFF
We currently have:

Source Code: APNGReader.java
Notes:

See also:
Aperio ImageScope
• many SVS datasets
• an SVS specification document
• the ability to generate additional SVS datasets

We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: SVSReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

See also:

Aperio ImageScope

### 18.11 Applied Precision CellWorX

Extensions: .htd, .pnl

Developer: Applied Precision

**Support**

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Applied Precision CellWorX

We currently have:

• a few CellWorX datasets

We would like to have:

• a CellWorX specification document
• more CellWorX datasets

**Ratings**

Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

---

35 [http://www.api.com](http://www.api.com)
Additional Information

Source Code: CellWorxReader.java

Notes:

18.12 AVI (Audio Video Interleave)

Extensions: .avi

Developer: Microsoft

Support

BSD-licensed: ✔

Export: ✔

Officially Supported Versions:

Supported Metadata Fields: AVI (Audio Video Interleave)

Freely Available Software:

- AVI Reader plugin for ImageJ
- AVI Writer plugin for ImageJ

We currently have:

- several AVI datasets

We would like to have:

- more AVI datasets, including:

  - files with audio tracks and/or multiple video tracks
  - files compressed with a common unsupported codec
  - 2+ GB files

Ratings

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

Additional Information

Source Code: AVIReader.java

Notes:

- Bio-Formats can save image stacks as AVI (uncompressed).
- The following codecs are supported for reading:
  - Microsoft Run-Length Encoding (MSRLE)
  - Microsoft Video (MSV1)
  - Raw (uncompressed)

36 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/CellWorxReader.java
37 http://www.microsoft.com/
40 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/AVIReader.java
– JPEG

See also:
AVI RIFF File Reference\[^{41}\] AVI on Wikipedia\[^{42}\]

### 18.13 Axon Raw Format

**Extensions:** .arf

**Owner:** INDECBioSystems\[^{43}\]

**Support**

BSD-licensed: ✗

Export: ✗

**Officially Supported Versions:**

**Supported Metadata Fields:** Axon Raw Format

We currently have:

- one ARF dataset
- a specification document\[^{44}\]

We would like to have:

- more ARF datasets

**Ratings**

- Pixels: ▲
- Metadata: ▼
- Openness: ▲
- Presence: ▼
- Utility: ▼

**Additional Information**

Source Code: ARFReader.java\[^{45}\]

**Notes:**

### 18.14 BD Pathway

**Extensions:** .exp, .tif

**Owner:** BD Biosciences\[^{46}\]

**Support**

BSD-licensed: ✗

Export: ✗

**Officially Supported Versions:**


\[^{42}\]http://en.wikipedia.org/wiki/Audio_Video_Interleave

\[^{43}\]http://www.indecbiosystems.com/

\[^{44}\]http://www.indecbiosystems.com/imagingworkbench/ApplicationNotes/IWAppNote11-ARF_File_Format.pdf

\[^{45}\]https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ARFReader.java

\[^{46}\]http://www.bdbiosciences.com
Supported Metadata Fields: *BD Pathway*

We currently have:

- a few BD Pathway datasets

We would like to have:

- more BD Pathway datasets

**Ratings**

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

**Additional Information**

Source Code: BDReader.java

Notes:

18.15 Becker & Hickl SPCImage

Extensions: .sdt

Owner: Becker-Hickl

**Support**

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: *Becker & Hickl SPCImage*

We currently have:

- an SDT specification document (from 2008 April, in PDF)
- an SDT specification document (from 2006 June, in PDF)
- Becker & Hickl’s SPCImage software
- a large number of SDT datasets
- the ability to produce new datasets

We would like to have:

**Ratings**

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

---

47 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/BDReader.java

48 http://www.becker-hickl.de/

49 http://www.becker-hickl.de/software/tcspc/softwaretcpspecial.htm
Additional Information

Source Code: SDTReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

18.16 Bio-Rad Gel

Extensions: .1sc

Owner: Bio-Rad

Support

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: Bio-Rad Gel

We currently have:

• software that can read Bio-Rad Gel files
• several Bio-Rad Gel files

We would like to have:

• a Bio-Rad Gel specification
• more Bio-Rad Gel files

Ratings

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

Additional Information

Source Code: BioRadGelReader.java

Notes:

18.17 Bio-Rad PIC

Extensions: .pic, .raw, .xml

Developer: Bio-Rad

Owner: Carl Zeiss, Inc.

Support

BSD-licensed: 

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Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Bio-Rad PIC

Freely Available Software:
  • Bio-Rad PIC reader plugin for ImageJ

We currently have:
  • a PIC specification document (v4.5, in PDF)
  • an older PIC specification document (v4.2, from 1996 December 16, in DOC)
  • a large number of PIC datasets
  • the ability to produce new datasets

We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▲

Openness: ▲

Presence: ▲

Utility: ▲

**Additional Information**

Source Code: BioRadReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

  • Commercial applications that support this format include:
    – Bitplane Imaris
    – SVI Huygens

18.18 Bio-Rad SCN

Extensions: .scn

Developer: Bio-Rad

Owner: Bio-Rad

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Bio-Rad SCN

We currently have:
  • a few Bio-Rad .scn files

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54http://rsb.info.nih.gov/ij/plugins/biorad.html
56http://www.bitplane.com/
57http://svi.nl/
58http://www.bio-rad.com
We would like to have:

**Ratings**

- Pixels: ▲
- Metadata: ▼
- Openness: ▼
- Presence: ▼
- Utility: ▼

**Additional Information**

Source Code: BioRadSCNReader.java

Notes:

## 18.19 Bitplane Imaris

Extensions: .ims

Owner: Bitplane

**Support**

- BSD-licensed: ✗
- Export: ✗

Officially Supported Versions: 2.7, 3.0, 5.5

Supported Metadata Fields: Bitplane Imaris

We currently have:

- an Imaris (RAW) specification document (from no later than 1997 November 11, in HTML)
- an Imaris 5.5 (HDF) specification document
- Bitplane’s bfFileReaderImaris3N code (from no later than 2005, in C++)
- several older Imaris (RAW) datasets
- one Imaris 3 (TIFF) dataset
- several Imaris 5.5 (HDF) datasets

We would like to have:

- an Imaris 3 (TIFF) specification document
- more Imaris 3 (TIFF) datasets

**Ratings**

- Pixels: ▲
- Metadata: ▲
- Openness: ▼
- Presence: ▼
- Utility: ▼

**Additional Information**

[59]https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/BioRadSCNReader.java


Source Code: ImarisHDFReader.java\textsuperscript{62}, ImarisTiffReader.java\textsuperscript{63}, ImarisReader.java\textsuperscript{64}

Notes:

- **There are three distinct Imaris formats:**
  1. the old binary format (introduced in Imaris version 2.7)
  2. Imaris 3, a TIFF variant (introduced in Imaris version 3.0)
  3. Imaris 5.5, an HDF variant (introduced in Imaris version 5.5)

### 18.20 Bruker MRI

**Developer:** Bruker\textsuperscript{65}

**Support**

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: *Bruker MRI*

**Freely Available Software:**

- Bruker plugin for ImageJ\textsuperscript{66}

We currently have:

- a few Bruker MRI datasets

We would like to have:

- an official specification document

**Ratings**

Pixels: 

Metadata: ▲

Openness: ▼

Presence: 

Utility: ▼

**Additional Information**

Source Code: BrukerReader.java\textsuperscript{67}

Notes:

### 18.21 Burleigh

**Extensions:** .img

Owner: Burleigh Instruments

**Support**

\textsuperscript{62}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImarisHDFReader.java

\textsuperscript{63}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImarisTiffReader.java

\textsuperscript{64}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImarisReader.java

\textsuperscript{65}http://www.bruker.com/

\textsuperscript{66}http://rsbweb.nih.gov/ij/plugins/bruker.html

\textsuperscript{67}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/BrukerReader.java
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Burleigh

We currently have:

• Pascal code that can read Burleigh files (from ImageSXM)
• a few Burleigh files

We would like to have:

• a Burleigh file format specification
• more Burleigh files

Ratings

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information

Source Code: BurleighReader.java

Notes:

18.22 Canon DNG

Extensions: .cr2, .crw

Developer: Canon

Support

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Canon DNG

Freely Available Software:

• IrfanView

We currently have:

• a few example datasets

We would like to have:

• an official specification document

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68 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/BurleighReader.java
69 http://canon.com
70 http://www.irfanview.com/
Ratings
Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: DNGReader.java

18.23 Cellomics

Extensions: .c01
Developer: Thermo Fisher Scientific

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: Cellomics
We currently have:
• a few Cellomics .c01 datasets

We would like to have:
• a Cellomics .c01 specification document
• more Cellomics .c01 datasets

Ratings
Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: CellomicsReader.java

Notes:

71 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/DNGReader.java
72 http://www.thermofisher.com/
73 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/CellomicsReader.java
18.24 cellSens VSI

Extensions: .vsi
Developer: Olympus

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: cellSens VSI

We currently have:
• a few example datasets

We would like to have:
• an official specification document

Ratings
Pixels: ▼
Metadata: ▲
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: CellSensReader.java

Notes:

18.25 CellVoyager

Extensions: .xml,.tif
Owner: Yokogawa

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: CellVoyager

We currently have:
• a few example datasets

We would like to have:

Ratings
Pixels: ▲
Metadata: ▲

http://www.olympus.com/
http://www.yokogawa.com/
18.26 DeltaVision

Extensions: .dv, .r3d

Owner: Applied Precision

Support

BSD-licensed: X
Export: X

Officially Supported Versions:

Supported Metadata Fields: DeltaVision

Freely Available Software:

• DeltaVision Opener plugin for ImageJ

Sample Datasets:

• Applied Precision Datasets

We currently have:

• a DV specification document (v2.10 or newer, in HTML)
• numerous DV datasets

We would like to have:

Ratings

Pixels:
Metadata:
Openness:
Presence:
Utility:

Additional Information

Source Code: DeltavisionReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

• The Deltavision format is based on the Medical Research Council (MRC) file format.
• Commercial applications that support DeltaVision include:

77 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/CellVoyagerReader.java
78 http://www.api.com/
80 http://www.api.com/downloads/software/softworxexplorer2.0/SampleImages.zip
81 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/DeltavisionReader.java
See also:
DeltaVision system description

18.27 DICOM

Extensions: .dcm, .dicom

Developer: National Electrical Manufacturers Association

Support

BSD-licensed: ✓

Export: ✗

Officially Supported Versions:

Supported Metadata Fields: DICOM

Freely Available Software:

• OsiriX Medical Imaging Software
• ezDICOM
• Wikipedia’s list of freeware health software

Sample Datasets:

• MRI Chest from FreeVol-3D web site
• Medical Image Samples from Sebastien Barre’s Medical Imaging page
• DICOM sample image sets from OsiriX web site

We currently have:

• DICOM specification documents (PS 3 - 2007, from 2006 December 28, in DOC and PDF)
• numerous DICOM datasets

We would like to have:

Ratings

Pixels: ▲

Metadata: ▲

Openness: ▲

Presence: ◻

Utility: ◻
Additional Information

Source Code: DicomReader.java

Notes:

- DICOM stands for “Digital Imaging and Communication in Medicine”.
- Bio-Formats supports both compressed and uncompressed DICOM files.

See also:
DICOM homepage

18.28 ECAT7

Extensions: .v
Developer: Siemens

Support

BSD-licensed: 
Export: 

Officially Supported Versions:

Supported Metadata Fields: ECAT7
We currently have:
• a few ECAT7 files

We would like to have:
• an ECAT7 specification document
• more ECAT7 files

Ratings

Pixels: 
Metadata: 
Openness: 
Presence: 
Utility: 

Additional Information

Source Code: Ecat7Reader.java

Notes:

18.29 EPS (Encapsulated PostScript)

Extensions: .eps, .epsi, .ps
Developer: Adobe

Support

94https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/DicomReader.java
95http://medical.nema.org/
96http://www.siemens.com
97https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/Ecat7Reader.java
98http://www.adobe.com/
BSD-licensed:

Export:

Officially Supported Versions:

Supported Metadata Fields: EPS (Encapsulated PostScript)

Freely Available Software:

- EPS Writer plugin for ImageJ

We currently have:

- a few EPS datasets
  - the ability to produce new datasets

We would like to have:

Ratings

Pixels:

Metadata:

Openness:

Presence:

Utility:

Additional Information

Source Code: EPSReader.java Source Code: EPSWriter.java

Notes:

- Bio-Formats can save individual planes as EPS.
- Certain types of compressed EPS files are not supported.

18.30 Evotec/PerkinElmer Opera Flex

Extensions: .flex, .mea, .res

Developer: Evotec Technologies, now PerkinElmer

Support

BSD-licensed:

Export:

Officially Supported Versions:

Supported Metadata Fields: Evotec/PerkinElmer Opera Flex

We currently have:

- many Flex datasets

We would like to have:

- a freely redistributable LuraWave LWF decoder

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100 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/EPSReader.java
101 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/EPSWriter.java
102 http://www.perkinelmer.com/
Ratings

Pixels: ⬆
Metadata: ⬆
Openness: ⬇️
Presence: ⬇️
Utility: ⬇️

Additional Information

Source Code: FlexReader.java

Notes:

The LuraWave LWF decoder library (i.e. lwf_jsdk2.6.jar) with license code is required to decode wavelet-compressed Flex files.

See also:
LuraTech (developers of the proprietary LuraWave LWF compression used for Flex image planes)

18.31 FEI

Extensions: .img
Developer: FEI

Support

BSD-licensed: ☝️
Export: ☝️

Officially Supported Versions:

Supported Metadata Fields: FEI

We currently have:

• a few FEI files

We would like to have:

• a specification document
• more FEI files

Ratings

Pixels: ⬇️
Metadata: ⬇️
Openness: ⬇️
Presence: ⬇️
Utility: ⬇️

Additional Information

Source Code: FEIReader.java

Notes:

https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/FlexReader.java
http://www.luratech.com/
http://www.fei.com/
https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/FEIReader.java

18.31. FEI
18.32 FEI TIFF

Extensions: .tiff
Developer: FEI

Support
BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: FEI TIFF
We currently have:
• a few FEI TIFF datasets
We would like to have:

Ratings
Pixels:▲
Metadata:□
Openness:□
Presence:▼
Utility:▼

Additional Information
Source Code: FEITiffReader.java
Notes:

18.33 FITS (Flexible Image Transport System)

Extensions: .fits
Developer: National Radio Astronomy Observatory

Support
BSD-licensed: ✔
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: FITS (Flexible Image Transport System)
We currently have:
• a FITS specification document (NOST 100-2.0, from 1999 March 29, in HTML)
• several FITS datasets
We would like to have:

Ratings
Pixels:▲

107http://www.fei.com
109http://www.nrao.edu/
110http://archive.stsci.edu/fits/fits_standard/
18.34 Gatan Digital Micrograph

Extensions: .dm3
Owner: Gatan

Support

BSD-licensed: 
Export: 

Officially Supported Versions: 3

Supported Metadata Fields: *Gatan Digital Micrograph* 

Freely Available Software:

- DM3 Reader plugin for ImageJ
- EMAN

We currently have:

- Gatan’s ImageReader2003 code (from 2003, in C++)
- numerous DM3 datasets

We would like to have:

- a DM3 specification document

Ratings

Pixels: 
Metadata: 
Openness: 
Presence: 
Utility: 

Additional Information

Source Code: *GatanReader.java* 

Notes:

See also:

MAST:FITS homepage, FITS Support Office

18.34. Gatan Digital Micrograph

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111 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/FitsReader.java

112 http://archive.stsci.edu/fits/

113 http://fits.gsfc.nasa.gov/

114 http://www.gatan.com/


116 http://blake.bcm.edu/EMAN/

117 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/GatanReader.java
Notes:
Commercial applications that support .dm3 files include DataSqueeze\(^\text{118}\).

18.35 Gatan Digital Micrograph 2

Extensions: .dm2
Developer: Gatan\(^\text{119}\)

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions: 2

Supported Metadata Fields: *Gatan Digital Micrograph 2*

We currently have:
- Pascal code that can read DM2 files (from ImageSXM)
- a few DM2 files

We would like to have:
- an official DM2 specification document
- more DM2 files

Ratings

- Pixels: 
- Metadata: ▼
- Openness: ▼
- Presence: ▼
- Utility: 

Additional Information

Source Code: *GatanDM2Reader.java*\(^\text{120}\)

Notes:

18.36 GIF (Graphics Interchange Format)

Extensions: .gif
Developer: CompuServe\(^\text{121}\)
Owner: Unisys\(^\text{122}\)

Support

- BSD-licensed: ✓
- Export: ❌

Officially Supported Versions:

\(^\text{18}\) http://www.datasqueezesoftware.com/
\(^\text{119}\) http://www.gatan.com
\(^\text{120}\) https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/GatanDM2Reader.java
\(^\text{121}\) http://www.compuserve.com/
\(^\text{122}\) http://www.unisys.com/
Supported Metadata Fields: *GIF (Graphics Interchange Format)*

Freely Available Software:

- Animated GIF Reader plugin for ImageJ\(^{123}\)
- GIF Stack Writer plugin for ImageJ\(^{124}\)

We currently have:

- a GIF specification document\(^{125}\) (Version 89a, from 1990, in HTML)
- numerous GIF datasets
- the ability to produce new datasets

We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▲

Openness: ▼

Presence: ▲

Utility: ▼

**Additional Information**

Source Code: GIFReader.java\(^{126}\)

Notes:

18.37 Hamamatsu Aquacosmos NAF

Extensions: .naf

Developer: Hamamatsu\(^{127}\)

**Support**

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: *Hamamatsu Aquacosmos NAF*

We currently have:

- a few NAF files

We would like to have:

- a specification document
- more NAF files

**Ratings**

Pixels: ▼

Metadata: ▼

Openness: ▼

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\(^{125}\) [http://tronche.com/computer-graphics/gif/](http://tronche.com/computer-graphics/gif/)

\(^{126}\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/GIFReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/GIFReader.java)

\(^{127}\) [http://www.hamamatsu.com/](http://www.hamamatsu.com/)
18.38 Hamamatsu HIS

Extensions: .his
Owner: Hamamatsu

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: Hamamatsu HIS
We currently have:
• Pascal code that can read HIS files (from ImageSXM)
• several HIS files
We would like to have:
• an HIS specification
• more HIS files

Ratings
Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: HISReader.java

Notes:

18.39 Hamamatsu ndpi

Extensions: .ndpi
Developer: Hamamatsu

Support
BSD-licensed: ❌

Notes:

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128 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NAFReader.java
129 http://www.hamamatsu.com
130 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/HISReader.java
131 http://www.hamamatsu.com
Export:❌
Officially Supported Versions:
Supported Metadata Fields: *Hamamatsu ndpi*
Freely Available Software:
  • NDP.view
Sample Datasets:
  • OpenSlide
We currently have:
  • many example datasets
We would like to have:
  • an official specification document

**Ratings**

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: NDPIReader.java

Notes:

18.40 Hamamatsu VMS

Extensions: .vms
Developer: Hamamatsu

**Support**

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: *Hamamatsu VMS*
Sample Datasets:
  • OpenSlide
We currently have:
  • a few example datasets
  • developer documentation from the OpenSlide project
We would like to have:

---

133 http://openslide.cs.cmu.edu/download/openslide-testdata/Hamamatsu/
134 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NDPIReader.java
135 http://www.hamamatsu.com
136 http://openslide.cs.cmu.edu/download/openslide-testdata/Hamamatsu-vms/
137 http://openslide.org/Hamamatsu%20format/
• an official specification document
• more example datasets

Ratings
Pixels: 
Metadata: 
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: HamamatsuVMSReader.java\textsuperscript{138}
Notes:

18.41 Hitachi S-4800

Extensions: .txt, .tif, .bmp, .jpg
Developer: Hitachi\textsuperscript{139}
Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: Hitachi S-4800
We currently have:
• several Hitachi S-4800 datasets
We would like to have:

Ratings
Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▼

Additional Information
Source Code: HitachiReader.java\textsuperscript{140}
Notes:
\begin{footnotesize}
\textsuperscript{138}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/HamamatsuVMSReader.java
\textsuperscript{139}http://www.hitachi-hta.com/sites/default/files/technotes/Hitachi_4800_STEM.pdf
\textsuperscript{140}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/HitachiReader.java
\end{footnotesize}
18.42 ICS (Image Cytometry Standard)

Extensions: .ics, .ids

Developer: P. Dean et al.

Support

BSD-licensed: ✓

Export: ✓

Officially Supported Versions: 1.0, 2.0

Supported Metadata Fields: ICS (Image Cytometry Standard)

Freely Available Software:

• Libics (ICS reference library)\(^{141}\)
• ICS Opener plugin for Image\(^{142}\)
• IrfanView\(^{143}\)

We currently have:

• numerous ICS datasets

We would like to have:

Ratings

Pixels: ▲

Metadata: ▲

Openness: ▲

Presence: ▲

Utility: ▲

Additional Information

Source Code: ICSReader.java\(^{144}\) Source Code: ICSWriter.java\(^{145}\)

Notes:

• ICS version 1.0 datasets have two files - an .ics file that contains all of the metadata in plain-text format, and an .ids file that contains all of the pixel data.

• ICS version 2.0 datasets are a single .ics file that contains both pixels and metadata.

Commercial applications that can support ICS include:

• Bitplane Imaris\(^{146}\)
• SVI Huygens\(^{147}\)

18.43 Imacon

Extensions: .fff

Owner: Hasselblad\(^{148}\)

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\(^{141}\)http://libics.sourceforge.net/

\(^{142}\)http://valelab.ucsf.edu/%7Enstuurman/IJplugins/Ics_Opener.html

\(^{143}\)http://www.irfanview.com/

\(^{144}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/ICSReader.java

\(^{145}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/ICSWriter.java

\(^{146}\)http://www.bitplane.com/

\(^{147}\)http://svi.nl/

\(^{148}\)http://www.hasselbladusa.com/
Support

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: Imacon

We currently have:

• one Imacon file

We would like to have:

• more Imacon files

Ratings

Pixels:

Metadata:

Openness:

Presence:

Utility:

Additional Information

Source Code: ImaconReader.java

Notes:

18.44 ImagePro Sequence

Extensions: .seq

Owner: Media Cybernetics

Support

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: ImagePro Sequence

We currently have:

• the Image-Pro Plus software

• a few SEQ datasets

• the ability to produce more datasets

We would like to have:

• an official SEQ specification document

Ratings

Pixels:

Metadata:
18.45 ImagePro Workspace

Extensions: .ipw

Owner: Media Cybernetics

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: ImagePro Workspace

We currently have:

• the Image-Pro Plus software
• a few IPW datasets
• the ability to produce more datasets

We would like to have:

• an official IPW specification document
• more IPW datasets:
  – multiple datasets in one file
  – 2+ GB files

Ratings

Pixels: 🔺

Metadata: 🔺

Openness: ▼

Presence: ▼

Utility: ▼

Additional Information

Source Code: IPWReader.java

Notes:

Bio-Formats uses a modified version of the Apache Jakarta POI library to read IPW files.

152https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/SEQReader.java
153http://www.mediacy.com/
155https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/IPWReader.java
156http://jakarta.apache.org/poi/
18.46 IMAGIC

Extensions: .hed, .img
Developer: Image Science

Support

BSD-licensed: X
Export: X

Officially Supported Versions:
Supported Metadata Fields: IMAGIC

Freely Available Software:
  • em2em

We currently have:
  • one example dataset
  • official file format documentation

We would like to have:
  • more example datasets

Ratings

Pixels:▲
Metadata:▲
Openness:▲
Presence:
Utility:

Additional Information

Source Code: ImagicReader.java

Notes:

See also:
IMAGIC specification

18.47 IMOD

Extensions: .mod
Developer: Boulder Laboratory for 3-Dimensional Electron Microscopy of Cells
Owner: Boulder Laboratory for 3-Dimensional Electron Microscopy of Cells

Support

BSD-licensed: X
Export: X

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157 http://www.imagescience.de
158 http://www.imagescience.de/em2em.html
159 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImagicReader.java
160 http://www.imagescience.de/em2em.html
161 http://bio3d.colorado.edu
162 http://bio3d.colorado.edu
Officially Supported Versions:

Supported Metadata Fields: IMOD

Freely Available Software:
- IMOD

We currently have:
- a few sample datasets
- official documentation

We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: IMODReader.java

Notes:

### 18.48 Improvision Openlab LIFF

**Extensions:** .liff

**Developer:** Improvision

**Owner:** PerkinElmer

**Support**

BSD-licensed: ❌
Export: ❌

Officially Supported Versions: 2.0, 5.0

**Supported Metadata Fields:** Improvision Openlab LIFF

We currently have:
- an Openlab specification document (from 2000 February 8, in DOC)
- Improvision’s XLIFFFileImporter code for reading Openlab LIFF v5 files (from 2006, in C++)
- several Openlab datasets

We would like to have:
- more Openlab datasets (preferably with 32-bit integer data)

**Ratings**

Pixels: ▲
Metadata: ▲

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[163] http://bio3d.colorado.edu/imod/
[165] https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/IMODReader.java
Openness: ▲
Presence: ▲
Utility: ▼

Additional Information

Source Code: OpenlabReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

See also:
Openlab software review

18.49 Improvision Openlab Raw

Extensions: .raw
Developer: Improvision
Owner: PerkinElmer

Support

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Improvision Openlab Raw

We currently have:

• an Openlab Raw specification document (from 2004 November 09, in HTML)
• a few Openlab Raw datasets

We would like to have:

Ratings

Pixels: ▲
Metadata: ▲
Openness: ▲

Presence: ▼
Utility: ▼

Additional Information

Source Code: OpenlabRawReader.java

Notes:

See also:

Openlab software review

168 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/OpenlabReader.java
169 http://www.improvision.com/products/openlab/
170 http://www.improvision.com/
171 http://www.perkinelmer.com/
172 http://cellularimaging.perkinelmer.com/support/technical_notes/detail.php?id=344
174 http://www.improvision.com/products/openlab/
18.50 Improvision TIFF

Extensions: .tif
Developer: Improvision
Owner: PerkinElmer

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Improvision TIFF

We currently have:
- an Improvision TIFF specification document
- a few Improvision TIFF datasets

We would like to have:

Ratings
Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▲

Additional Information
Source Code: ImprovisionTiffReader.java

Notes:
Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

See also:
Openlab software overview

18.51 Imspector OBF

Extensions: .obf, .msr
Developer: Department of NanoBiophotonics, MPI-BPC
Owner: MPI-BPC

Support
BSD-licensed: ✓
Export: ✗

Officially Supported Versions:

175 http://www.improvision.com/
176 http://www.perkinelmer.com/
177 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImprovisionTiffReader.java
178 http://www.improvision.com/products/openlab/
179 https://imspector.mpibpc.mpg.de/index.html
180 http://www.mpibpc.mpg.de/
Supported Metadata Fields: *Inspector OBF*

We currently have:

- a few `.msr` datasets
- a specification document\(^\text{181}\)

We would like to have:

**Ratings**

- Pixels: ▲
- Metadata: ▫
- Openness: ▲
- Presence: ▼
- Utility: ▼

**Additional Information**

Source Code: `OBFReader.java`\(^\text{182}\)

Notes:

### 18.52 InCell 1000

Extensions: `.xdce`, `.tif`

Developer: GE\(^\text{183}\)

**Support**

- BSD-licensed: ☒
- Export: ☒

Officially Supported Versions:

Supported Metadata Fields: *InCell 1000*

We currently have:

- a few InCell 1000 datasets

We would like to have:

- an InCell 1000 specification document
- more InCell 1000 datasets

**Ratings**

- Pixels: ▲
- Metadata: ▲
- Openness: ▫
- Presence: ▼
- Utility: ▼

**Additional Information**

Source Code: `InCellReader.java`\(^\text{184}\)

\(^\text{181}\) https://imspector.mpibpc.mpg.de/documentation/fileformat.html

\(^\text{182}\) https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/OBFReader.java

\(^\text{183}\) http://gelifesciences.com/

\(^\text{184}\) https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/InCellReader.java
Notes:

18.53 InCell 3000

Extensions: .frm
Developer: GE

Support

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: InCell 3000

Sample Datasets:

- Broad Bioimage Benchmark Collection

We currently have:

- a few example datasets

We would like to have:

- an official specification document

Ratings

Pixels:
Metadata:
Openness:
Presence:
Utility:

Additional Information

Source Code: InCell3000Reader.java

Notes:

18.54 INR

Extensions: .inr

Support

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: INR

We currently have:

- several sample .inr datasets

Notes:

185 http://gelifesciences.com/
186 http://www.broadinstitute.org/bbcb/BBBC013/
187 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/InCell3000Reader.java
We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▼

Openness: ▼

Presence: ▼

Utility: ▼

**Additional Information**

Source Code: INRReader.java

Notes:

**18.55 Inveon**

Extensions: .hdr

**Support**

BSD-licensed: ✗

Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Inveon

We currently have:

a few Inveon datasets

We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▲

Openness: ▼

Presence: ▼

Utility: ▼

**Additional Information**

Source Code: InveonReader.java

Notes:

**18.56 IPLab**

Extensions: .ipl

Developer: Scanalytics

Owner: was BD Biosystems, now BioVision Technologies

Notes:

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188 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/INRReader.java

189 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/InveonReader.java

190 http://wwwbdbiosciencescom/

191 http://wwwbioviscomiplabhtm
Support

BSD-licensed: ✗
Export: ✗
Officially Supported Versions:
Supported Metadata Fields: **IPLab**
Freely Available Software:
  - IPLab Reader plugin for ImageJ\(^{192}\)
We currently have:
  - an IPLab specification document (v3.6.5, from 2004 December 1, in PDF)
  - several IPLab datasets
We would like to have:
  - more IPLab datasets (preferably with 32-bit integer or floating point data)

Ratings

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▼

Additional Information

Source Code: **IPLabReader.java**\(^{193}\)
Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

Commercial applications that support IPLab include:
  - Bitplane Imaris\(^{194}\)
  - SVI Huygens\(^{195}\)

See also:
IPLab software review\(^{196}\)

18.57 IPLab-Mac

Extensions: .ipm
Owner: **BioVision Technologies**\(^{197}\)

Support

BSD-licensed: ✗
Export: ✗
Officially Supported Versions:

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\(^{192}\)[http://rsb.info.nih.gov/ij/plugins/iplab-reader.html]
\(^{193}\)[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/IPLabReader.java]
\(^{194}\)[http://www.bitplane.com/]
\(^{195}\)[http://svi.nl/]
\(^{196}\)[http://www.biovis.com/iplab.htm]
\(^{197}\)[http://biovis.com/]

18.57. IPLab-Mac 141
Supported Metadata Fields: *IPLab-Mac*

We currently have:

- a few IPLab-Mac datasets
- a specification document

We would like to have:

- more IPLab-Mac datasets

**Ratings**

- Pixels: 🔺
- Metadata: 🔻
- Openness: 🔺
- Presence: 🔻
- Utility: 🔻

**Additional Information**

Source Code: [IvisionReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/IvisionReader.java)

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

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**18.58 JEOL**

Extensions: .dat, .img, .par

Owner: JEOL

**Support**

- BSD-licensed: ✗
- Export: ✗

Officially Supported Versions:

Supported Metadata Fields: *JEOL*

We currently have:

- Pascal code that reads JEOL files (from ImageSXM)
- a few JEOL files

We would like to have:

- an official specification document
- more JEOL files

**Ratings**

- Pixels: 🔻
- Metadata: 🔻
- Openness: 🔻
- Presence: 🔻
- Utility: 🔻

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18.58. JEOL

198 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/IvisionReader.java

199 http://www.jeol.com
### Additional Information

Source Code: [JEOLReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/JEOLReader.java)

Notes:

#### 18.59 JPEG

Extensions: .jpg

Developer: [Independent JPEG Group](http://www.ijg.org/)

Support

BSD-licensed: ✓

Export: ✓

Officially Supported Versions:

Supported Metadata Fields: JPEG

We currently have:

- a JPEG specification document (v1.04, from 1992 September 1, in PDF)
- numerous JPEG datasets
- the ability to produce more datasets

We would like to have:

### Ratings

Pixels: 🔴

Metadata: 🔴

Openness: 🔵

Presence: 🔵

Utility: 🔴

### Additional Information

Source Code: [JPEGReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/JPEGReader.java)  
Source Code: [JPEGWriter.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/JPEGWriter.java)

Notes:

Bio-Formats can save individual planes as JPEG. Bio-Formats uses the [Java Image I/O](http://docs.oracle.com/javase/6/docs/technotes/guides/imageio/) API to read and write JPEG files. JPEG stands for “Joint Photographic Experts Group”.

See also:

JPEG homepage

#### 18.60 JPEG 2000

Extensions: .jp2

Developer: [Independent JPEG Group](http://www.ijg.org/)

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200[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/JEOLReader.java](#)
201[http://www.ijg.org/](#)
202[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/JPEGReader.java](#)
203[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/JPEGWriter.java](#)
204[http://docs.oracle.com/javase/6/docs/technotes/guides/imageio/](#)
205[http://www.jipeg.org/jpeg/index.html](#)
206[http://www.ijg.org/](#)
Support

BSD-licensed: ✔
Export: ✔

Officially Supported Versions:

Supported Metadata Fields: JPEG 2000

Freely Available Software:

- JJ2000 (JPEG 2000 library for Java)

We currently have:

- a JPEG 2000 specification document (final draft, from 2000, in PDF)
- a few .jp2 files

We would like to have:

Ratings

Pixels: 
Metadata: 
Openness: 
Presence: 
Utility: 

Additional Information


Notes:

Bio-Formats uses the JAI Image I/O Tools library to read JP2 files. JPEG stands for “Joint Photographic Experts Group”.

18.61 JPK

Extensions: .jpk
Developer: JPK Instruments

Support

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: JPK

We currently have:

- Pascal code that can read JPK files (from ImageSXM)
- a few JPK files

We would like to have:

- an official specification document

Notes:

Bio-Formats uses the JAI Image I/O Tools library to read JP2 files. JPEG stands for “Joint Photographic Experts Group”. 

18.61. JPK
• more JPK files

Ratings

Pixels:  
Metadata:  
Openness:  
Presence:  
Utility:  

Additional Information

Source Code: JPKReader.java

Notes:

18.62 JPX

Extensions: .jpx

Developer: JPEG Committee

Support

BSD-licensed:  
Export:  

Officially Supported Versions:

Supported Metadata Fields: *JPX*

We currently have:

• a few .jpx files

We would like to have:

Ratings

Pixels:  
Metadata:  
Openness:  
Presence:  
Utility:  

Additional Information

Source Code: JPXReader.java

Notes:

18.63 Khoros VIFF (Visualization Image File Format) Bitmap

Extensions: .xv

Developer: Khoral

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214 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/JPKReader.java
215 http://www.jpeg.org/jpeg2000/
216 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/JPXReader.java
217 http://www.khoral.com/company/
Owner: AccuSoft

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: *Khoros VIFF (Visualization Image File Format) Bitmap*

Sample Datasets:

- VIFF Images

We currently have:

- several VIFF datasets

We would like to have:

Ratings

Pixels: 🟢

Metadata: 🟢

Openness: 🟢

Presence: 🟢

Utility: 🟢

Additional Information

Source Code: KhorosReader.java

Notes:

See also:

VisiQuest software overview (formerly known as KhorosPro)

### 18.64 Kodak BIP

Extensions: .bip

Developer: Kodak/Carestream

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: *Kodak BIP*

We currently have:

- a few .bip datasets

We would like to have:

- an official specification document

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218 http://www.accusoft.com/company/
220 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/KhorosReader.java
221 http://www.accusoft.com/products/visiquest/
222 http://carestream.com
### Ratings

**Pixels:** ▲
**Metadata:** ▼
**Openness:** ▼
**Presence:** ▼
**Utility:** ▼

### Additional Information

**Source Code:** KodakReader.java²²³

**Notes:**

**See also:**

Information on Image Station systems²²⁴

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### 18.65 Lambert Instruments FLIM

**Extensions:** .fli

**Developer:** Lambert Instruments²²⁵

**Support**

- BSD-licensed: ✗
- Export: ✗

**Officially Supported Versions:**

**Supported Metadata Fields:** Lambert Instruments FLIM

We currently have:

- an LI-FLIM specification document
- several example LI-FLIM datasets

We would like to have:

**Ratings**

**Pixels:** ▲
**Metadata:** ▲
**Openness:** ▲
**Presence:** ▼
**Utility:** ▼

**Additional Information**

**Source Code:** LiFlimReader.java²²⁶

**Notes:**

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

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²²³: https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/KodakReader.java
²²⁴: http://carestream.com/PublicContent.aspx?langType=1033&id=448953
²²⁵: http://www.lambert-instruments.com
²²⁶: https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LiFlimReader.java

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18.65. Lambert Instruments FLIM 147
18.66 LaVision Imspector

Extensions: .msr
Developer: LaVision BioTec

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: LaVision Imspector
We currently have:
  • a few .msr files
We would like to have:

Ratings
Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: InspectorReader.java
Notes:

18.67 Leica LCS LEI

Extensions: .lei, .tif
Developer: Leica Microsystems CMS GmbH
Owner: Leica

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: Leica LCS LEI
Freely Available Software:
  • Leica LCS Lite
We currently have:
  • an LEI specification document (beta 2.000, from no later than 2004 February 17, in PDF)
  • many LEI datasets

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227 http://www.lavisionbiotec.com/
228 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ImspectorReader.java
229 http://www.leica-microsystems.com/
230 http://www.leica.com/
231 ftp://ftp.llt.de/softlib/LCSLite/LCSLite2611537.exe
We would like to have:

**Ratings**

- **Pixels:**
- **Metadata:**
- **Openness:**
- **Presence:**
- **Utility:**

**Additional Information**

**Source Code:** LeicaReader.java

**Notes:**

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

LCS stands for “Leica Confocal Software”. LEI presumably stands for “Leica Experimental Information”.

Commercial applications that support LEI include:

- Bitplane Imaris
- SVI Huygens
- Image-Pro Plus

### 18.68 Leica LAS AF LIF (Leica Image File Format)

**Extensions:** .lif

**Developer:** Leica Microsystems CMS GmbH

**Owner:** Leica

**Support**

- BSD-licensed: ✗
- Export: ✗

**Officially Supported Versions:** 1.0, 2.0

**Supported Metadata Fields:** Leica LAS AF LIF (Leica Image File Format)

**Freely Available Software:**

- Leica LAS AF Lite (links at bottom of page)

We currently have:

- a LIF specification document (version 2, from no later than 2007 July 26, in PDF)
- a LIF specification document (version 1, from no later than 206 April 3, in PDF)
- numerous LIF datasets

We would like to have:

**Ratings**

- **Pixels:**

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232 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LeicaReader.java

233 http://www.bitplane.com/

234 http://svi.nl/

235 http://www.mediacy.com/

236 http://www.leica-microsystems.com/

237 http://www.leica.com/

238 http://www.leica-microsystems.com/products/microscope-software/software-for-life-science-research/las-af-4-advanced-fluorescence/
Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties. LAS stands for “Leica Application Suite”. AF stands for “Advanced Fluorescence”.

Commercial applications that support LIF include:

- Bitplane Imaris
- SVI Huygens
- Amira

18.69 Leica SCN

Extensions: .scn

Developer: Leica Microsystems

Support

BSD-licensed:

Export:

Officially Supported Versions: 2012-03-10

Supported Metadata Fields: Leica SCN

We currently have:

- a few sample datasets

We would like to have:

- an official specification document
- sample datasets that cannot be opened

Ratings

Pixels:

Metadata:

Openness:

Presence:

Utility:

Additional Information

Source Code: LeicaSCNReader.java

---

239 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LIFReader.java
240 http://www.bitplane.com/
241 http://svi.nl/
242 http://www.amira.com/
243 http://www.leica-microsystems.com/
244 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LeicaSCNReader.java
18.70 LEO

Extensions: .sxm
Owner: Zeiss

Support

BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: LEO

We currently have:
• Pascal code that can read LEO files (from ImageSXM)
• a few LEO files

We would like to have:
• an official specification document
• more LEO files

Ratings

Pixels: 
Metadata: 
Openness: 
Presence: 
Utility: 

Additional Information

Source Code: LEOReader.java

Notes:

18.71 Li-Cor L2D

Extensions: .l2d, .tif, .scn
Owner: LiCor Biosciences

Support

BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: Li-Cor L2D

We currently have:
• a few L2D datasets

Notes:
We would like to have:

- an official specification document
- more L2D datasets

**Ratings**

- Pixels: ▲
- Metadata: ▼
- Openness: ▼
- Presence: ▼
- Utility: ▼

**Additional Information**

Source Code: L2DReader.java\(^{248}\)

Notes:

L2D datasets cannot be imported into OME using server-side import. They can, however, be imported from ImageJ, or using the omeul utility.

### 18.72 LIM (Laboratory Imaging/Nikon)

**Extensions:** .lim

**Owner:** Laboratory Imaging\(^{249}\)

**Support**

- BSD-licensed: ✗
- Export: ✗

**Officially Supported Versions:**

**Supported Metadata Fields:** LIM (Laboratory Imaging/Nikon)

We currently have:

- several LIM files
- the ability to produce more LIM files

We would like to have:

- an official specification document

**Ratings**

- Pixels: ▼
- Metadata: ▼
- Openness: ▼
- Presence: ▼
- Utility: ▼

**Additional Information**

Source Code: LIMReader.java\(^{250}\)

Notes:

\(^{248}\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/L2DReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/L2DReader.java)


\(^{250}\) [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LIMReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/LIMReader.java)
Bio-Formats only supports uncompressed LIM files. Commercial applications that support LIM include:

- NIS Elements

### 18.73 MetaMorph 7.5 TIFF

Extensions: .tiff

Owner: Molecular Devices

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: MetaMorph 7.5 TIFF

We currently have:

- a few Metamorph 7.5 TIFF datasets

We would like to have:

**Ratings**

Pixels: ⬆️

Metadata: ⬆️

Openness: ⬆️

Presence: ⬇️

Utility: 😕

**Additional Information**

Source Code: MetamorphTiffReader.java

Notes:

### 18.74 MetaMorph Stack (STK)

Extensions: .stk, .nd

Owner: Molecular Devices

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: MetaMorph Stack (STK)

We currently have:

- an STK specification document (from 2006 November 21, in DOC)

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251 http://www.nis-elements.com/
252 http://www.moleculardevices.com/
253 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/MetamorphTiffReader.java
254 http://www.moleculardevices.com/
• an older STK specification document (from 2005 March 25, in DOC)
• an ND specification document (from 2002 January 24, in PDF)
• a large number of datasets

We would like to have:

**Ratings**

Pixels: ▼
Metadata: ▲
Openness: ▲
Presence: ▲
Utility: ▲

**Additional Information**

Source Code: MetamorphReader.java

Notes:

*Please note that while we have specification documents for this format, we are not able to distribute them to third parties.*

Commercial applications that support STK include:

• Bitplane Imaris
• SVI Huygens
• DIMIN

**See also:**

Metamorph imaging system overview

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**18.75 MIAS (Maia Scientific)**

Extensions: .tif

Developer: Maia Scientific

**Support**

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: MIAS (Maia Scientific)

We currently have:

• several MIAS datasets

We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▼

---

255 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/MetamorphReader.java
256 http://www.bitplane.com/
257 http://svi.nl/
258 http://dimin.net/
259 http://www.metamorph.com/
260 http://www.selectscience.net/supplier/maia-scientific/?compID=6088
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: MIASReader.java\(^{261}\)
Notes:

### 18.76 Micro-Manager

Extensions: .tif, .txt, .xml
Developer: Vale Lab\(^{262}\)

**Support**
- BSD-licensed: ☑
- Export: ✗

Officially Supported Versions:
- Supported Metadata Fields: Micro-Manager

Freely Available Software:
- Micro-Manager\(^{263}\)

We currently have:
- many Micro-manager datasets

We would like to have:

**Ratings**
- Pixels: ▲
- Metadata: ▲
- Openness: ▲
- Presence: ▼
- Utility: ☐

Additional Information
Source Code: MicromanagerReader.java\(^{264}\)
Notes:

### 18.77 MINC MRI

Extensions: .mnc
Developer: McGill University\(^{265}\)

**Support**
- BSD-licensed: ✗

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\(^{261}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/MIASReader.java
\(^{262}\)http://valelab.ucsf.edu/
\(^{263}\)http://micro-manager.org/
\(^{264}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/MicromanagerReader.java
\(^{265}\)http://www.bic.mni.mcgill.ca/ServicesSoftware/MINC
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: MINC MRI

Freely Available Software:

• MINC

We currently have:

• a few MINC files

We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▼

Openness: ▼

Presence: ▼

Utility: ▼

**Additional Information**

Source Code: MINCReader.java

Notes:

**18.78 Minolta MRW**

Extensions: .mrw

Developer: Minolta

**Support**

BSD-licensed: ✗

Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Minolta MRW

Freely Available Software:

• dcraw

We currently have:

• several .mrw files

We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▼

Openness: ▼

Presence: ▼

---

266 http://bic.mni.mcgill.ca/ServicesSoftware/MINC

267 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/MINCReader.java

268 http://www.konicaminolta.com/

269 http://www.cybercom.net/%7Edcoffin/dcraw/
Utility: ▼

Additional Information

Source Code: MRWReader.java

Notes:

18.79 MNG (Multiple-image Network Graphics)

Extensions: .mng

Developer: MNG Development Group

Support

BSD-licensed: ✔

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: MNG (Multiple-image Network Graphics)

Freely Available Software:

- libmng (MNG reference library)

Sample Datasets:

- MNG sample files

We currently have:

- the libmng-testsuites package (from 2003 March 05, in C)
- a large number of MNG datasets

We would like to have:

Ratings

Pixels: □

Metadata: □

Openness: ▲

Presence: ▼

Utility: ▼

Additional Information

Source Code: MNGReader.java

Notes:

See also:

MNG homepage

MNG specification

http://www.libpng.org/pub/mng/mngsuite-1.0/MNGsuite.zip/download

http://downloads.sourceforge.net/libmng/MNGsuite-20030305.zip

http://www.libpng.org/pub/mng/spec
18.80 Molecular Imaging

Extensions: .stp
Owner: Molecular Imaging Corp, San Diego CA (closed)

Support

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: Molecular Imaging
We currently have:
  • Pascal code that reads Molecular Imaging files (from ImageX)
  • a few Molecular Imaging files

We would like to have:
  • an official specification document
  • more Molecular Imaging files

Ratings

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information

Source Code: MolecularImagingReader.java

Notes:

18.81 MRC (Medical Research Council)

Extensions: .mrc
Developer: MRC Laboratory of Molecular Biology

Support

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: MRC (Medical Research Council)
Sample Datasets:
  • golgi.mrc

We currently have:
  • an MRC specification document (in HTML)

Notes:

278 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/MolecularImagingReader.java
279 http://www2.mrc-lmb.cam.ac.uk/
280 http://bio3d.colorado.edu/imod/files/imod_data.tar.gz
281 http://ami.scripps.edu/software/mrctools/mrc_specification.php
• another MRC specification document\textsuperscript{282} (in TXT)
• a few MRC datasets

We would like to have:

Ratings
Pixels: 🔺
Metadata: 🔺
Openness: 🔺
Presence: ☐
Utility: ☐

Additional Information
Source Code: MRCReader.java\textsuperscript{283}

Notes:
Commercial applications that support MRC include:
• Bitplane Imaris\textsuperscript{284}

See also:
MRC on Wikipedia\textsuperscript{285}

\textbf{18.82 NEF (Nikon Electronic Format)}

Extensions: .nef, .tif
Developer: Nikon\textsuperscript{286}

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: \textit{NEF (Nikon Electronic Format)}

Sample Datasets:
• neffile1.zip\textsuperscript{287}
  • Sample NEF images\textsuperscript{288}

We currently have:
• a NEF specification document (v0.1, from 2003, in PDF)
• several NEF datasets

We would like to have:

Ratings
Pixels: 🔺

\begin{flushright}
\footnotesize
\textsuperscript{282}http://bio3d.colorado.edu/imod/doc/mrc_format.txt
\textsuperscript{283}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/MRCReader.java
\textsuperscript{284}http://www.bitplane.com/
\textsuperscript{285}http://en.wikipedia.org/wiki/MRC_%28file_format%29
\textsuperscript{286}http://www.nikon.com/
\textsuperscript{287}http://www.outbackphoto.com/workshop/NEF_conversion/neffile1.zip
\textsuperscript{288}http://www.nikondigital.org/articles/library/nikon_d2x_first_impressions.htm
\end{flushright}

18.82. NEF (Nikon Electronic Format) 159
Openness: ◼
Presence: ◼
Utility: ◼

Additional Information
Source Code: NikonReader.java

Notes:
Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

See also:
NEF Conversion

18.83 NIfTI

Extensions: .img, .hdr
Developer: National Institutes of Health

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: NIfTI
Sample Datasets:
• Official test data

We currently have:
• NIfTI specification documents
• several NIfTI datasets

We would like to have:

Ratings
Pixels: ◆
Metadata: ◼
Openness: ◆
Presence: ◼
Utility: ◼

Additional Information
Source Code: NiftiReader.java

Notes:

289https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NikonReader.java
290http://www.outbackphoto.com/workshop/NEF_conversion/nefconversion.html
291http://www.nih.gov/
293http://nifti.nimh.nih.gov/nifti-1/
294https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NiftiReader.java
18.84 Nikon Elements TIFF

Extensions: .tiff
Developer: Nikon

Support
BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: Nikon Elements TIFF

We currently have:
• a few Nikon Elements TIFF files

We would like to have:
• more Nikon Elements TIFF files

Ratings
Pixels: 
Metadata: 
Openness: 
Presence: 
Utility: 

Additional Information
Source Code: NikonElementsTiffReader.java

Notes:

18.85 Nikon EZ-C1 TIFF

Extensions: .tiff
Developer: Nikon

Support
BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: Nikon EZ-C1 TIFF

We currently have:
• a few Nikon EZ-C1 TIFF files

We would like to have:

Ratings
Pixels: 
Metadata: 

Notes:

295 http://www.nikon.com
296 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NikonElementsTiffReader.java
297 http://www.nikon.com/
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: NikonTiffReader.java

Notes:

18.86 Nikon NIS-Elements ND2

Extensions: .nd2
Developer: Nikon USA

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Nikon NIS-Elements ND2

Freely Available Software:
- NIS-Elements Viewer from Nikon

We currently have:
- many ND2 datasets

We would like to have:
- an official specification document

Ratings
Pixels: ▲
Metadata: ▲
Openness: ▼
Presence: ▲
Utility: ▲

Additional Information
Source Code: NativeND2Reader.java

Notes:

There are two distinct versions of ND2: an old version, which uses JPEG-2000 compression, and a new version which is either uncompressed or Zip-compressed. We are not aware of the version number or release date for either format.

Bio-Formats uses the JAI Image I/O Tools library to read ND2 files compressed with JPEG-2000.

There is also an ND2 reader that uses Nikon’s native libraries. To use it, you must be using Windows and have Nikon’s ND2 reader plugin for ImageJ installed. Additionally, you will need to download LegacyND2Reader.dll and place it in your ImageJ plugin folder.

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298 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NikonTiffReader.java
299 http://www.nikonusa.com/
300 http://www.nis-elements.com/resources-downloads.html
301 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/NativeND2Reader.java
302 http://java.net/projects/jai-imageio
304 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/lib/LegacyND2Reader.dll?raw=true
18.87 NRRD (Nearly Raw Raster Data)

Extensions: .nrrd, .nhdr, .raw, .txt

Developer: Teem developers\(^{305}\)

Support

BSD-licensed: ✔️

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: NRRD (Nearly Raw Raster Data)

Freely Available Software:

- nrrd (NRRD reference library)\(^{306}\)

Sample Datasets:

- Diffusion tensor MRI datasets\(^{307}\)

We currently have:

- an nrrd specification document\(^{308}\) (v1.9, from 2005 December 24, in HTML)
- a few nrrd datasets

We would like to have:

Ratings

Pixels: 🔺

Metadata: 🔺

Openness: 🔺

Presence: 🔻

Utility: 🔺

Additional Information

Source Code: NRRDReader.java\(^{309}\)

Notes:

18.88 Olympus CellR/APL

Extensions: .apl, .mtb, .tnb, .tif, .obsep

Owner: Olympus\(^{310}\)

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Olympus CellR/APL

http://teem.sourceforge.net/

http://teem.sourceforge.net/nrrd/

http://www.sci.utah.edu/~gk/DTI-data/

http://teem.sourceforge.net/nrrd/format.html

https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/NRRDReader.java

http://www.olympus.com/
We currently have:
  • a few CellR datasets

We would like to have:
  • more Cellr datasets
  • an official specification document

Ratings

Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ◯
Utility: ◯

Additional Information

Source Code: APLReader.java

Notes:

18.89 Olympus FluoView FV1000

Extensions: .oib, .oif

Owner: Olympus

Support

BSD-licensed: ✗
Export: ✗

Officially Supported Versions: 1.0, 2.0

Supported Metadata Fields: Olympus FluoView FV1000

Freely Available Software:
  • FV-Viewer from Olympus

We currently have:
  • an OIF specification document (v2.0.0.0, from 2008, in PDF)
  • an FV1000 specification document (v1.0.0.0, from 2004 June 22, in PDF)
  • older FV1000 specification documents (draft, in DOC and XLS)
  • many FV1000 datasets

We would like to have:
  • more OIB datasets (especially 2+ GB files)
  • more FV1000 version 2 datasets

Ratings

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▲

**Additional Information**

**Source Code:** FV1000Reader.java

**Notes:**

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

Bio-Formats uses a modified version of the [Apache Jakarta POI](http://jakarta.apache.org/poi/) library to read OIB files. OIF stands for “Original Imaging Format”. OIB stands for “Olympus Image Binary”. OIF is a multi-file format that includes an .oif file and a directory of .tif, .roi, .pty, .lut, and .bmp files. OIB is a single file format.

Commercial applications that support this format include:

- Bitplane Imaris
- SVI Huygens

**See also:**

Olympus FluoView Resource Center

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### 18.90 Olympus FluoView TIFF

**Extensions:** .tif

**Owner:** Olympus

**Support**

BSD-licensed: ❌

Export: ❌

**Officially Supported Versions:**

**Supported Metadata Fields:** *Olympus FluoView TIFF*

**Freely Available Software:**

- DIMIN

We currently have:

- a FluoView specification document (from 2002 November 14, in DOC)
- Olympus’ FluoView Image File Reference Suite (from 2002 March 1, in DOC)
- several FluoView datasets

We would like to have:

**Ratings**

Pixels: ▲

Metadata: ▲

Openness: ▲

Presence: ▼

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314 | https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/FV1000Reader.java
---

315 | http://jakarta.apache.org/poi/
---

316 | http://www.bitplane.com/
---

317 | http://svi.nl/
---

318 | http://www.olympusfluoview.com
---

319 | http://www.olympus.com/
---

320 | http://www.dimin.net/
---
Utility:

Additional Information

Source Code: FluoviewReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

Commercial applications that support this format include:

- Bitplane Imaris
- SVI Huygens

18.91 Olympus ScanR

Extensions: .xml, .dat, .tif

Developer: Olympus

Owner: Olympus

Support

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: Olympus ScanR

We currently have:

- several ScanR datasets

We would like to have:

Ratings

Pixels:

Metadata:

Openness:

Presence:

Utility:

Additional Information

Source Code: ScanrReader.java

Notes:

18.92 Olympus SIS TIFF

Extensions: .tiff

Developer: Olympus

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321 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/FluoviewReader.java

322 http://www.bitplane.com/

323 http://svi.nl/

324 http://www.olympus.com/

325 http://www.olympus-sis.com/

326 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ScanrReader.java

327 http://www.olympus-sis.com/
Support

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: *Olympus SIS TIFF*

We currently have:
- a few example SIS TIFF files

We would like to have:

**Ratings**

Pixels: 
Metadata: 
Openness: 
Presence: 🔻
Utility: 

**Additional Information**

Source Code: *SISReader.java*  
Notes:

**18.93 OME-TIFF**

Extensions: *.ome.tiff*  
Developer: *Open Microscopy Environment*  

**Support**

BSD-licensed: ✔
Export: ✔


Supported Metadata Fields: *OME-TIFF*

We currently have:
- an *OME-TIFF specification document* (from 2006 October 19, in HTML)
- many OME-TIFF datasets
- the ability to produce additional datasets

We would like to have:

**Ratings**

Pixels: 
Metadata: 
Openness: 
Presence: 🔻

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Utility: ▲

Additional Information

Source Code: OMETiffReader.java \[^{332}\] Source Code: OMETiffWriter.java \[^{333}\]

Notes:

Bio-Formats can save image stacks as OME-TIFF.

Commercial applications that support OME-TIFF include:

• Bitplane Imaris \[^{334}\]
• SVI Huygens \[^{335}\]

See also:

OME-TIFF technical overview \[^{336}\]

18.94 OME-XML

Extensions: .ome \[^{337}\]

Developer: Open Microscopy Environment \[^{338}\]

Support

BSD-licensed: ✔

Export: ✔


Supported Metadata Fields: OME-XML

We currently have:

• OME-XML specification documents \[^{339}\]
• many OME-XML datasets
• the ability to produce more datasets

We would like to have:

Ratings

Pixels: ▲

Metadata: ▲

Openness: ▲

Presence: ▼

Utility: ▲

Additional Information

Source Code: OMEXMLReader.java \[^{340}\] Source Code: OMEXMLWriter.java \[^{341}\]

Notes:

\[^{332}\] https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/OMETiffReader.java
\[^{333}\] https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/OMETiffWriter.java
\[^{334}\] http://www.bitplane.com/
\[^{335}\] http://svi.nl/
\[^{336}\] http://www.openmicroscopy.org/site/support/ome-model/ome-tiff/index.html
\[^{337}\] http://www.openmicroscopy.org/Schemas/
\[^{338}\] http://www.openmicroscopy.org/OME-XML/
\[^{339}\] http://www.openmicroscopy.org/Schemas/OME-XML/
\[^{340}\] https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/OMEXMLReader.java
\[^{341}\] https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/OMEXMLWriter.java
Bio-Formats uses the OME-XML Java library\(^{342}\) to read OME-XML files.

Commercial applications that support OME-XML include:

- Bitplane Imaris\(^{343}\)
- SVI Huygens\(^{344}\)

### 18.95 Oxford Instruments

**Extensions:** .top

**Owner:** Oxford Instruments\(^{345}\)

**Support**

- BSD-licensed: \(\times\)
- Export: \(\times\)

**Officially Supported Versions:**

**Supported Metadata Fields:** *Oxford Instruments*

We currently have:

- Pascal code that can read Oxford Instruments files (from ImageSXM)
- A few Oxford Instruments files

We would like to have:

- An official specification document
- More Oxford Instruments files

**Ratings**

- Pixels: ▼
- Metadata: ▼
- Openness: ▼
- Presence: ▼
- Utility: ▼

**Additional Information**

**Source Code:** OxfordInstrumentsReader.java\(^{346}\)

**Notes:**

### 18.96 PCORAW

**Extensions:** .pcoraw, .rec

**Developer:** PCO\(^{347}\)

**Support**

- BSD-licensed: \(\times\)

---

\(^{342}\)http://www.openmicroscopy.org/site/support/ome-model/ome-xml/java-library.html

\(^{343}\)http://www.bitplane.com/

\(^{344}\)http://svi.nl/

\(^{345}\)http://www.oxinst.com


\(^{347}\)http://www.pco.de/
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: **PCORAW**
We currently have:
- a few example datasets
We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▲

**Additional Information**
Source Code: PCORAWReader.java

Notes:

### 18.97 PCX (PC Paintbrush)

Extensions: .pcx
Developer: ZSoft Corporation

**Support**

BSD-licensed: ✔
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: **PCX (PC Paintbrush)**
We currently have:
- several .pcx files
- the ability to generate additional .pcx file
We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**
Source Code: PCXReader.java

Notes:

---

349 [https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/PCXReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/PCXReader.java)
Commercial applications that support PCX include Zeiss LSM Image Browser\textsuperscript{350}.

### 18.98 Perkin Elmer Densitometer

Extensions: .pds  
Developer: Perkin Elmer\textsuperscript{351}

**Support**

BSD-licensed: ✗  
Export: ✗  

Officially Supported Versions:

Supported Metadata Fields: Perkin Elmer Densitometer

We currently have:

- a few PDS datasets

We would like to have:

- an official specification document  
- more PDS datasets

**Ratings**

Pixels: 
Metadata: 
Openness: 
Presence: 
Utility:

**Additional Information**

Source Code: PDSReader.java\textsuperscript{352}

Notes:

### 18.99 PerkinElmer Operetta

Extensions: .tiff, .xml  
Developer: PerkinElmer\textsuperscript{353}

**Support**

BSD-licensed: ✗  
Export: ✗  

Officially Supported Versions:

Supported Metadata Fields: PerkinElmer Operetta

We currently have:

- a few sample datasets

\textsuperscript{351}\url{http://www.perkinelmer.com}  
\textsuperscript{352}\url{https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/PDSReader.java}  
\textsuperscript{353}\url{http://www.perkinelmer.com/}
We would like to have:

- an official specification document
- more sample datasets

**Ratings**

- **Pixels:** 🟢
- **Metadata:** 🟢
- **Openness:** 🟢
- **Presence:** 🟢
- **Utility:** 🟢

**Additional Information**

Source Code: OperettaReader.java

Notes:

**18.100 PerkinElmer UltraView**

Extensions: .tif, .2, .3, .4, etc.

Owner: PerkinElmer

**Support**

- BSD-licensed: ✗
- Export: ✗

Officially Supported Versions:

Supported Metadata Fields: PerkinElmer UltraView

We currently have:

- several UltraView datasets

We would like to have:

**Ratings**

- **Pixels:** 🟢
- **Metadata:** 🟢
- **Openness:** 🟢
- **Presence:** 🟢
- **Utility:** 🟢

**Additional Information**

Source Code: PerkinElmerReader.java

Notes:

Other associated extensions include: .tim, .zpo, .csv, .htm, .cfg, .ano, .rec

Commercial applications that support this format include:

- Bitplane Imaris

---

18.100. PerkinElmer UltraView
• Image-Pro Plus

See also:
PerkinElmer UltraView system overview

18.101 PGM (Portable Gray Map)

Extensions: .pgm
Developer: Netpbm developers

Support
BSD-licensed: ✓
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: PGM (Portable Gray Map)

Freely Available Software:
• Netpbm graphics filter

We currently have:
• a PGM specification document (from 2003 October 3, in HTML)
• a few PGM files

We would like to have:

Ratings
Pixels: ▲
Metadata: ▼
Openness: ▲
Presence: ▼
Utility: ▼

Additional Information
Source Code: PGMReader.java

Notes:

18.102 Adobe Photoshop PSD

Extensions: .psd
Developer: Adobe

Support
BSD-licensed: ❌
Export: ❌

Notes:

¡http://www.mediacom.com/
359http://www.perkinelmer.com/pages/020/cellularimaging/products/ultraviewvoxsystemsoverview.xhtml
360http://netpbm.sourceforge.net/
361http://netpbm.sourceforge.net/doc/pgm.html
362https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/PGMReader.java
363http://www.adobe.com/
Officially Supported Versions: 1.0

Supported Metadata Fields: *Adobe Photoshop PSD*

We currently have:

- a PSD specification document (v3.0.4, 16 July 1995)
- a few PSD files

We would like to have:

- more PSD files

### Ratings

- Pixels: 
- Metadata: 
- Openness: 
- Presence: 
- Utility: 

### Additional Information

Source Code: [PSDReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/PSDReader.java)

Notes:

### 18.103 Photoshop TIFF

Extensions: .tif, .tiff

Developer: *Adobe*[^365]

#### Support

- BSD-licensed: ✗
- Export: ✗

Officially Supported Versions:

Supported Metadata Fields: *Photoshop TIFF*

We currently have:

- a Photoshop TIFF specification document
- a few Photoshop TIFF files

We would like to have:

### Ratings

- Pixels: 
- Metadata: 
- Openness: 
- Presence: 
- Utility: 

### Additional Information


[^365]: [http://www.adobe.com](http://www.adobe.com)
Source Code: PhotoshopTiffReader.java\textsuperscript{366}

Notes:

18.104 PicoQuant Bin

Extensions: .bin

Developer: PicoQuant\textsuperscript{367}

Support

BSD-licensed:

Export:

Officially Supported Versions:

Supported Metadata Fields: PicoQuant Bin

Freely Available Software:

- SymphoTime64\textsuperscript{368}

We currently have:

- a few example datasets

We would like to have:

Ratings

Pixels:

Metadata:

Openness:

Presence:

Utility:

Additional Information

Source Code: PQBinReader.java\textsuperscript{369}

Notes:

18.105 PICT (Macintosh Picture)

Extensions: .pict

Developer: Apple Computer\textsuperscript{370}

Support

BSD-licensed:

Export:

Officially Supported Versions:

Supported Metadata Fields: PICT (Macintosh Picture)

We currently have:

\textsuperscript{366}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/PhotoshopTiffReader.java
\textsuperscript{367}http://www.picoquant.com/
\textsuperscript{368}http://www.picoquant.com/products/category/software/symphotime-64-fluorescence-lifetime-imaging-and-correlation-software
\textsuperscript{369}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/PQBinReader.java
\textsuperscript{370}http://www.apple.com
many PICT datasets

We would like to have:

**Ratings**

- **Pixels:**
- **Metadata:**
- **Openness:**
- **Presence:**
- **Utility:**

**Additional Information**

Source Code: PictReader.java

Notes:

QuickTime for Java is required for reading vector files and some compressed files.

See also:

PICT technical overview Another PICT technical overview

### 18.106 PNG (Portable Network Graphics)

Extensions: .png

Developer: PNG Development Group

**Support**

- BSD-licensed: ✓
- Export: ✓

**Officially Supported Versions:**

**Supported Metadata Fields:** PNG (Portable Network Graphics)

**Freely Available Software:**

- PNG Writer plugin for ImageJ

We currently have:

- a PNG specification document (W3C/ISO/IEC version, from 2003 November 10, in HTML)
- several PNG datasets

We would like to have:

**Ratings**

- **Pixels:**
- **Metadata:**
- **Openness:**
- **Presence:**

---

371 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/PictReader.java


374 http://www.prepressure.com/formats/pict/fileformat.htm

375 http://www.libpng.org/pub/png/pngnews.html


377 http://www.libpng.org/pub/png/spec/iso/
Utility:

Additional Information

Source Code: APNGReader.java

Notes:

Bio-Formats uses the Java Image I/O API to read and write PNG files.

See also:

PNG technical overview

18.107 Prairie Technologies TIFF

Extensions: .tif, .xml, .cfg

Developer: Prairie Technologies

Support

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: Prairie Technologies TIFF

We currently have:

• many Prairie datasets

We would like to have:

Ratings

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

Additional Information

Source Code: PrairieReader.java

Notes:

18.108 Quesant

Extensions: .afm

Developer: Quesant Instrument Corporation

Owner: KLA-Tencor Corporation

Support

378 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/APNGReader.java
379 http://docs.oracle.com/javase/6/docs/technotes/guides/imageio/
380 http://www.libpng.org/pub/png/
381 http://www.prairie-technologies.com/
382 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/PrairieReader.java
BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: *Quesant*

We currently have:
- Pascal code that can read Quesant files (from ImageSXM)
- several Quesant files

We would like to have:
- an official specification document
- more Quesant files

**Ratings**

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: *QuesantReader.java*[^384]

Notes:

**18.109 QuickTime Movie**

Extensions: .mov
Owner: *Apple Computer*[^385]

**Support**

BSD-licensed: ✔
Export: ✔

Officially Supported Versions:
Supported Metadata Fields: *QuickTime Movie*

Freely Available Software:
- *QuickTime Player*[^386]

We currently have:
- a *QuickTime specification document*[^387] (from 2001 March 1, in HTML)
- several QuickTime datasets
- the ability to produce more datasets

We would like to have:
- more QuickTime datasets, including:

[^384]: https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/QuesantReader.java
[^385]: http://www.apple.com/
files compressed with a common, unsupported codec
files with audio tracks and/or multiple video tracks

Ratings

Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ▲
Utility: ▼

Additional Information

Source Code: NativeQTReader.java388 Source Code: QTWriter.java389

Notes:

Bio-Formats has two modes of operation for QuickTime:

• QTJava mode requires QuickTime390 to be installed.
• Native mode works on systems with no QuickTime (e.g. Linux).

Bio-Formats can save image stacks as QuickTime movies. The following table shows supported codecs:

<table>
<thead>
<tr>
<th>Codec</th>
<th>Description</th>
<th>Native</th>
<th>QTJava</th>
</tr>
</thead>
<tbody>
<tr>
<td>raw</td>
<td>Full Frames (Uncompressed)</td>
<td>read &amp; write</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>iraw</td>
<td>Intel YUV Uncompressed</td>
<td>read only</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>rle</td>
<td>Animation (run length encoded RGB)</td>
<td>read only</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>jpeg</td>
<td>Still Image JPEG DIB</td>
<td>read only</td>
<td>read only</td>
</tr>
<tr>
<td>rpza</td>
<td>Apple Video 16 bit “road pizza”</td>
<td>read only (partial)</td>
<td>read only</td>
</tr>
<tr>
<td>mjpb</td>
<td>Motion JPEG codec</td>
<td>read only</td>
<td>read only</td>
</tr>
<tr>
<td>cvid</td>
<td>Cinepak</td>
<td>•</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>svq1</td>
<td>Sorenson Video</td>
<td>•</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>svq3</td>
<td>Sorenson Video 3</td>
<td>•</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>mp4v</td>
<td>MPEG-4</td>
<td>•</td>
<td>read &amp; write</td>
</tr>
<tr>
<td>h263</td>
<td>H.263</td>
<td>•</td>
<td>read &amp; write</td>
</tr>
</tbody>
</table>

See also:

QuickTime software overview391

18.110 RHK

Extensions: .sm2, .sm3

Owner: RHK Technologies392

Support

388https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/NativeQTReader.java
389https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/QTWriter.java
391http://www.apple.com/quicktime/
392http://www.rhk-tech.com
Bio-Formats Documentation, Release 5.0.4

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: **RHK**

We currently have:
- Pascal code that can read RHK files (from ImageSXM)
- a few RHK files

We would like to have:
- an official specification document
- more RHK files

**Ratings**

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: [RHKReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/RHKReader.java)

Notes:

**18.111 SBIG**

Owner: Santa Barbara Instrument Group (SBIG)

**Support**

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: **SBIG**

We currently have:
- an official SBIG specification document
- a few SBIG files

We would like to have:
- more SBIG files

**Ratings**

Pixels: ▲
Metadata: ▼
Openness: ▲

---

393 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/RHKReader.java
394 http://www.sbig.com
395 http://sbig.impulse.net/pdffiles/file.format.pdf
Presence: ▼
Utility: ▼

Additional Information
Source Code: SBIGReader.java

Notes:

18.112 Seiko

Extensions: .xqd, .xqf
Owner: Seiko

Support
BSD-licensed: 
Export: 

Officially Supported Versions:
Supported Metadata Fields: Seiko
We currently have:
• Pascal code that can read Seiko files (from ImageSXM)
• a few Seiko files
We would like to have:
• an official specification document
• more Seiko files

Ratings
Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: SeikoReader.java

Notes:

18.113 SimplePCI & HCImage

Extensions: .cxd
Developer: Compix

Support
BSD-licensed: 

---

396 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/SBIGReader.java
398 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/SeikoReader.java
399 http://hcimage.com
Export: 

Officially Supported Versions:

Supported Metadata Fields: SimplePCI & HCImage

We currently have:
  • several SimplePCI files

We would like to have:

**Ratings**

Pixels: ☑️

Metadata: ☐

Openness: ☑️

Presence: ☐

Utility: ☑️

**Additional Information**

Source Code: PCIReader.java

Notes:

Bio-Formats uses a modified version of the Apache Jakarta POI library to read CXD files.

See also:

SimplePCI software overview

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### 18.114 SimplePCI & HCImage TIFF

Extensions: .tiff

Developer: Hamamatsu

**Support**

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: SimplePCI & HCImage TIFF

We currently have:
  • a few SimplePCI TIFF datasets

We would like to have:
  • more SimplePCI TIFF datasets

**Ratings**

Pixels: ☑️

Metadata: ☐

Openness: ☑️

Presence: ☑️
Utility:

Additional Information
Source Code: SimplePCITiffReader.java\(^\text{404}\)
Notes:

### 18.115 SM Camera

**Support**

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: *SM Camera*

We currently have:

- Pascal code that can read SM-Camera files (from ImageSXM)
- a few SM-Camera files

We would like to have:

- an official specification document
- more SM-Camera files

**Ratings**

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: SMCameraReader.java\(^\text{405}\)

Notes:

### 18.116 SPIDER

Extensions: .spi, .stk

Developer: Wadsworth Center\(^\text{406}\)

**Support**

BSD-licensed: ❌
Export: ❌

Officially Supported Versions:

Supported Metadata Fields: *SPIDER*

Freely Available Software:

\(^\text{404}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/SimplePCITiffReader.java
\(^\text{405}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/SMCameraReader.java
\(^\text{406}\)http://www.wadsworth.org/spider_doc/spider/docs/spider.html
- **SPIDER**

We currently have:

- a few example datasets
- official file format documentation

We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: [SpiderReader.java](http://www.wadsworth.org/spider_doc/spider/docs/spider.html)

Notes:

18.117 **Targa**

Extensions: .tga

Developer: [Truevision](http://www.truevision.com)

**Support**

BSD-licensed: ✗
Export: ✗

Of officially Supported Versions:

Supported Metadata Fields: *Targa*

We currently have:

- a Targa specification document
- a few Targa files

We would like to have:

**Ratings**

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: [TargaReader.java](https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/TargaReader.java)

---


410 [http://www.truevision.com](http://www.truevision.com)

Notes:

18.118 Text

Extensions: .txt

Support

BSD-licensed: ✓

Export: ❌

Officially Supported Versions:

Supported Metadata Fields: Text

We currently have:

We would like to have:

Ratings

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

Additional Information

Source Code: TextReader.java

Notes:

Reads tabular pixel data produced by a variety of software.

18.119 TIFF (Tagged Image File Format)

Extensions: .tif

Developer: Aldus and Microsoft

Owner: Adobe

Support

BSD-licensed: ✓

Export: ✓

Officially Supported Versions:

Supported Metadata Fields: TIFF (Tagged Image File Format)

Sample Datasets:

• LZW TIFF data gallery

• Big TIFF

We currently have:

---

412 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/TextReader.java

413 http://www.adobe.com

414 http://marlin.life.utsa.edu/Data_Gallery.html

415 http://tiffcentral.com/
• a TIFF specification document\textsuperscript{416} (v6.0, from 1992 June 3, in PDF)
• many TIFF datasets
• a few BigTIFF datasets

We would like to have:

\textbf{Ratings}

Pixels: \uparrow

Metadata: \uparrow

Openness: \uparrow

Presence: \uparrow

Utility: \downarrow

\textbf{Additional Information}

Source Code: TiffReader.java\textsuperscript{417} Source Code: TiffWriter.java\textsuperscript{418}

Notes:

Bio-Formats can also read BigTIFF files (TIFF files larger than 4 GB). Bio-Formats can save image stacks as TIFF or BigTIFF.

\textbf{See also:}

TIFF technical overview\textsuperscript{419} BigTIFF technical overview\textsuperscript{420}

\section*{18.120 TillPhotonics TillVision}

Extensions: .vws

Developer: Till Photonics\textsuperscript{421}

\textbf{Support}

BSD-licensed: \xmark

Export: \xmark

Officially Supported Versions:

Supported Metadata Fields: \textit{TillPhotonics TillVision}

We currently have:

• several TillVision datasets

We would like to have:

• an official specification document

\textbf{Ratings}

Pixels: \xmark

Metadata: \xmark

Openness: \xmark

Presence: \xmark

Utility: \xmark

\textsuperscript{416} http://partners.adobe.com/asn/developer/PDFS/TN/TIFF6.pdf
\textsuperscript{417} https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/in/TiffReader.java
\textsuperscript{418} https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-bsd/src/loci/formats/out/TiffWriter.java
\textsuperscript{419} http://www.awaresystems.be/imaging/tiff/faq.html#q3
\textsuperscript{420} http://www.awaresystems.be/imaging/tiff/bigtiff.html
\textsuperscript{421} http://www.till-photonics.com/
Additional Information
Source Code: TillVisionReader.java\textsuperscript{422}

Notes:

18.121 Topometrix

Extensions: .tfr, .ffr, .zfr, .zfp, .2fl
Owner: TopoMetrix (now Veeco)\textsuperscript{423}

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: Topometrix

We currently have:
• Pascal code that reads Topometrix files (from ImageSXM)
• a few Topometrix files

We would like to have:
• an official specification document
• more Topometrix files

Ratings
Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

Additional Information
Source Code: TopometrixReader.java\textsuperscript{424}

Notes:

18.122 Trestle

Extensions: .tif, .sld, .jpg

Support
BSD-licensed: ❌
Export: ❌

Officially Supported Versions:
Supported Metadata Fields: Trestle

Sample Datasets:
\textsuperscript{422}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/TillVisionReader.java
\textsuperscript{423}http://www.veeco.com/
\textsuperscript{424}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/TopometrixReader.java
We currently have:

- a few example datasets
- developer documentation from the OpenSlide project\(^{426}\)

We would like to have:

**Ratings**

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

**Additional Information**

Source Code: [TrestleReader.java]\(^{427}\)

Notes:

**18.123 UBM**

Extensions: .pr3

**Support**

BSD-licensed: 

Export: 

Officially Supported Versions:

Supported Metadata Fields: **UBM**

We currently have:

- Pascal code that can read UBM files (from ImageSXM)
- one UBM file

We would like to have:

- an official specification document
- more UBM files

**Ratings**

Pixels: 

Metadata: 

Openness: 

Presence: 

Utility: 

**Additional Information**

Source Code: [UBMReader.java]\(^{428}\)

---

\(^{425}\)http://openslide.cs.cmu.edu/download/openslide-testdata/Trestle/
\(^{426}\)http://openslide.org/Trestle%20format/
\(^{427}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/TrestleReader.java
\(^{428}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/UBMReader.java
Notes:

### 18.124 Unisoku

Extensions: .dat, .hdr  
Owner: Unisoku

**Support**

BSD-licensed: ✗  
Export: ✗

Officially Supported Versions:  
Supported Metadata Fields: Unisoku

We currently have:  
- Pascal code that can read Unisoku files (from ImageSXM)  
- a few Unisoku files

We would like to have:  
- an official specification document  
- more Unisoku files

**Ratings**

Pixels: ▼  
Metadata: ▼  
Openness: ▼  
Presence: ▼  
Utility: ▼

**Additional Information**

Source Code: UnisokuReader.java

Notes:

### 18.125 Varian FDF

Extensions: .fdf  
Developer: Varian, Inc.

**Support**

BSD-licensed: ✗  
Export: ✗

Officially Supported Versions:  
Supported Metadata Fields: Varian FDF

We currently have:  
- a few Varian FDF datasets

---

429 http://www.unisoku.com  
430 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/UnisokuReader.java  
431 http://www.varianinc.com
We would like to have:

- an official specification document
- more Varian FDF datasets

**Ratings**

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: VarianFDFReader.java\(^{432}\)

Notes:

---

18.126 VG SAM

Extensions: .dti

**Support**

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:

Supported Metadata Fields: VG SAM

We currently have:

- a few VG-SAM files

We would like to have:

- an official specification document
- more VG-SAM files

**Ratings**

Pixels: ▼
Metadata: ▼
Openness: ▼
Presence: ▼
Utility: ▼

**Additional Information**

Source Code: VGSAMReader.java\(^{433}\)

Notes:

---

\(^{432}\)[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/VarianFDFReader.java]

\(^{433}\)[https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/VGSAMReader.java]
18.127 VisiTech XYS

Extensions: .xys, .html
Developer: VisiTech International

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: VisiTech XYS

We currently have:
• several VisiTech datasets

We would like to have:
• an official specification document

Ratings
Pixels: ▲
Metadata: ◄
Openness: ▼
Presence: ◄
Utility: ◄

Additional Information
Source Code: VisitechReader.java

Notes:

18.128 Volocity

Extensions: .mvd2
Developer: PerkinElmer

Support
BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: Volocity

Sample Datasets:
• PerkinElmer Downloads

We currently have:
• many example Volocity datasets

We would like to have:

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434 http://www.visitech.co.uk/
435 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/VisitechReader.java
436 http://www.perkinelmer.com/pages/020/cellularimaging/products/volocity.xhtml
437 http://cellularimaging.perkinelmer.com/downloads/
• an official specification document
• any Volocity datasets that do not open correctly

Ratings

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▲
Utility: ▲

Additional Information
Source Code: VolocityReader.java\textsuperscript{438}

Notes:

.mvd2 files are Metakit database files\textsuperscript{439}.

18.129 Volocity Library Clipping

Extensions: .acff
Developer: PerkinElmer\textsuperscript{440}

Support

BSD-licensed: x
Export: x

Officially Supported Versions:
Supported Metadata Fields: Volocity Library Clipping

We currently have:
• several Volocity library clipping datasets

We would like to have:
• any datasets that do not open correctly
• an official specification document

Ratings

Pixels: ▲
Metadata: ▲
Openness: ▲
Presence: ▲
Utility: ▲

Additional Information
Source Code: VolocityClippingReader.java\textsuperscript{441}

Notes:

\textsuperscript{438}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/VolocityReader.java
\textsuperscript{439}http://equi4.com/metakit/
\textsuperscript{440}http://www.perkinelmer.com/pages/020/cellularimaging/products/volocity.xhtml
\textsuperscript{441}https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/VolocityClippingReader.java
RGB .acff files are not yet supported. See #6413.

18.130 WA-TOP

Extensions: .wat
Developer: WA Technology
Owner: Oxford Instruments

Support

BSD-licensed: ✗
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: WA-TOP
We currently have:
- Pascal code that can read WA-TOP files (from ImageX)
- a few WA-TOP files
We would like to have:
- an official specification document
- more WA-TOP files

Ratings

Pixels: 🟣
Metadata: 🟣
Openness: 🟣
Presence: 🟣
Utility: 🟣

Additional Information

Source Code: WATOPReader.java

Notes:

18.131 Windows Bitmap

Extensions: .bmp
Developer: Microsoft and IBM

Support

BSD-licensed: ✓
Export: ✗

Officially Supported Versions:
Supported Metadata Fields: Windows Bitmap

Freely Available Software:

https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/WATOPReader.java
• BMP Writer plugin for ImageJ

We currently have:
• many BMP datasets

We would like to have:

Ratings

Pixels: ▲
Metadata: ▲
Openness: ▼
Presence: ▲
Utility: ▼

Additional Information

Source Code: BMPReader.java

Notes:

Compressed BMP files are currently not supported.

See also:

Technical Overview

18.132 Woolz

Extensions: .wlz

Developer: MRC Human Genetics Unit

Support

BSD-licensed: ❌

Export: ✔

Officially Supported Versions:

Supported Metadata Fields: Woolz

Freely Available Software:
• Woolz

We currently have:
• a few Woolz datasets

We would like to have:

Ratings

Pixels: ▲
Metadata: ▼
Openness: ▼
Presence: ▲
Utility: ▼

Additional Information

Source Code: WlzReader.java\(^{450}\) Source Code: WlzWriter.java\(^{451}\)

Notes:

### 18.133 Zeiss AxioVision TIFF

**Extensions:** .xml, .tiff

**Developer:** Carl Zeiss Microscopy GmbH\(^{452}\)

**Owner:** Carl Zeiss Microscopy GmbH\(^{453}\)

**Support**

BSD-licensed: ✗

Export: ✗

**Officially Supported Versions:**

**Supported Metadata Fields:** *Zeiss AxioVision TIFF*

**Freely Available Software:**

- *Zeiss ZEN Lite*\(^{454}\)

We currently have:

- many example datasets

We would like to have:

- an official specification document

**Ratings**

**Pixels:** ▲

**Metadata:** ▲

**Openness:** ▼

**Presence:** ▼

**Utility:** ▼

**Additional Information**

Source Code: ZeissTIFFReader.java\(^{455}\)

Notes:

### 18.134 Zeiss AxioVision ZVI (Zeiss Vision Image)

**Extensions:** .zvi

**Developer:** Carl Zeiss Microscopy GmbH (AxioVision)\(^{456}\)

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\(^{450}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/WlzReader.java

\(^{451}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/out/WlzWriter.java

\(^{452}\)http://www.zeiss.com/microscopy/

\(^{453}\)http://www.zeiss.com/microscopy/


\(^{455}\)https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ZeissTIFFReader.java

Owner: Carl Zeiss Microscopy GmbH

Support

BSD-licensed: ❌

Export: ❌

Officially Supported Versions: 1.0, 2.0

Supported Metadata Fields: Zeiss AxioVision ZVI (Zeiss Vision Image)

Freely Available Software:
- Zeiss Axiovision LE

We currently have:
- a ZVI specification document (v2.0.5, from 2010 August, in PDF)
- an older ZVI specification document (v2.0.2, from 2006 August 23, in PDF)
- an older ZVI specification document (v2.0.1, from 2005 April 21, in PDF)
- an older ZVI specification document (v1.0.26.01.01, from 2001 January 29, in DOC)
- Zeiss’ ZvImageReader code (v1.0, from 2001 January 25, in C++)
- many ZVI datasets

We would like to have:

Ratings

Pixels: ▲

Metadata: ▲

Openness: ▲

Presence: ❌

Utility: ❌

Additional Information

Source Code: ZeissZVIReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

Bio-Formats uses a modified version of the Apache Jakarta POI library to read ZVI files.

Commercial applications that support ZVI include Bitplane Imaris.

See also:

Axiovision software overview

18.135 Zeiss CZI

Extensions: .czi

Developer: Carl Zeiss Microscopy GmbH

---

18.135. Zeiss CZI
Support

BSD-licensed: ✗

Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Zeiss CZI

Freely Available Software:

- Zeiss ZEN 2012

We currently have:

- many example datasets
- official specification documents

We would like to have:

Ratings

Pixels: ★

Metadata: ★

Openness: ★

Presence: ▼

Utility: ★

Additional Information

Source Code: ZeissCZIReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

18.136 Zeiss LSM (Laser Scanning Microscope) 510/710

Extensions: .lsm, .mdb

Owner: Carl Zeiss Microscopy GmbH

Support

BSD-licensed: ✗

Export: ✗

Officially Supported Versions:

Supported Metadata Fields: Zeiss LSM (Laser Scanning Microscope) 510/710

Freely Available Software:

- Zeiss LSM Image Browser
- LSM Toolbox plugin for ImageJ
- LSM Reader plugin for ImageJ
- DIMIN
We currently have:

- LSM specification v3.2, from 2003 March 12, in PDF
- LSM specification v5.5, from 2009 November 23, in PDF
- LSM specification v6.0, from 2010 September 28, in PDF
- many LSM datasets

We would like to have:

**Ratings**

Pixels: 🔺

Metadata: 🔺

Openness: 

Presence: 🔺

Utility: 

**Additional Information**

Source Code: ZeissLSMReader.java

Notes:

Please note that while we have specification documents for this format, we are not able to distribute them to third parties.

Bio-Formats uses the [MDB Tools Java port](http://mdbtools.sourceforge.net/)

Commercial applications that support this format include:

- SVI Huygens
- Bitplane Imaris
- Amira
- Image-Pro Plus

---

472 https://github.com/openmicroscopy/bioformats/blob/v5.0.4/components/formats-gpl/src/loci/formats/in/ZeissLSMReader.java
473 http://mdbtools.sourceforge.net/
474 http://www2.svi.nl/
475 http://www.bitplane.com/
476 http://www.amira.com/
477 http://www mediacy.com/
## SUMMARY OF SUPPORTED METADATA FIELDS

### 19.1 Format readers

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69 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#DoubleAnnotation_Value  
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138 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ExperimenterRef_ID
139 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ExperimenterGroupRef_ID
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143 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#MicrobeamManipulationRef_ID
144 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
145 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROF_xsd.html#ROIRef_ID
146 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROF_xsd.html#ImagingEnvironment_AirPressure
149 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROF_xsd.html#ImagingEnvironment_Temperature
151 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROF_xsd.html#Shape_FillColor
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158 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROF_xsd.html#Shape_Locked
159 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROF_xsd.html#Shape_StrokeColor
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162 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ROI_xsd.html#Shape_Text
163 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ROI_xsd.html#Shape_TheC
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171 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ome_xsd.html#LightSource_ID
173 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ome_xsd.html#ManufacturerSpec_LotNumber
174 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
179 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ome_xsd.html#Laser_RepetitionRate
180 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
182 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OMF-2013-06/ome_xsd.html#Laser_Type

19.2. Metadata fields
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236 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth

237 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Text

238 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheC

239 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheT

240 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ

241 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Transform

242 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Visible

243 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Mask_Width

244 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Mask_X

245 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Mask_Y

246 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ExperimenterRef_ID


248 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ROIRef_ID

249 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#MicrobeamManipulation_Type

250 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_Attenuation

251 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_Wavelength

252 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_LotNumber

253 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer

254 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model

255 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber

256 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Microscope_Type

257 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification

258 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction

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19.2. Metadata fields

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284 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
286 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
287 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
289 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
290 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationRef_ID
291 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
293 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_HashSHA1
298 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
299 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
300 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationRef_ID
301 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ColumnNamingConvention
302 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Columns
303 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Description
304 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ExternalIdentifier
305 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ID
306 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Name
307 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_RowNamingConvention

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311 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_WellOriginY
312 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_Description
313 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_EndTime
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316 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_MaximumFieldCount
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319 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FillColor
320 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FillRule
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19.2. Metadata fields

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406 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Text
407 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheC
408 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheT
409 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ
410 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Transform
411 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Visible
412 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Width
413 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_X
414 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Y
415 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationRef_ID
416 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#Annotation_Description
417 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_Description
418 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_ID
419 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_Name
420 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_PlateRef_ID
421 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_ProtocolDescription
422 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_ProtocolIdentifier
423 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_ReagentSetDescription
424 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_ReagentSetIdentifier
425 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Screen_Type
426 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Name
428 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Y
429 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationRef_ID
430 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#Annotation_Description

19.2. Metadata fields
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452 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TransmittanceRange_CutOutTolerance
453 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TransmittanceRange_Transmittance
454 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TiffData_TiffData_UUID_FileName
455 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#UniversallyUniqueIdentifier
456 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationRef_ID
457 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Color
458 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Column
459 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellExternalDescription
460 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellExternalIdentifier
461 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellID
462 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_ID
464 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellRow
465 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellType
466 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#ReagentRef_ID
467 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSampleIndex
469 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSamplePositionY
470 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSampleTimepoint
471 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationRef_ID
472 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationID
473 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#AnnotationNamespace

19.2. Metadata fields
### 19.2.1 SlidebookReader

This page lists supported metadata fields for the Bio-Formats Olympus Slidebook format reader.

These fields are from the OME data model[^76]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 34 of them (7%).
- Of those, Bio-Formats fully or partially converts 34 (100%).

#### Supported fields

These fields are fully supported by the Bio-Formats Olympus Slidebook format reader:

- **Channel**: ID[^77]
- **Channel**: NDFilter[^78]
- **Channel**: Name[^79]
- **Channel**: SamplesPerPixel[^80]
- **Image**: AcquisitionDate[^81]
- **Image**: Description[^82]
- **Image**: ID[^83]
- **Image**: InstrumentRef[^84]
- **Image**: Name[^85]
- **Instrument**: ID[^86]
- **Objective**: Correction[^87]
- **Objective**: ID[^88]
- **Objective**: Immersion[^89]
- **Objective**: Model[^90]
- **Objective**: NominalMagnification[^91]
- **ObjectiveSettings**: ID[^92]

[^19]: Table 19.2 – continued from previous page

<table>
<thead>
<tr>
<th>Field</th>
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[^475]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SA_xsd.html#XMLAnnotation_Value
[^476]: http://www.openmicroscopy.org/site/support/ome-model/
[^477]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
[^478]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_NDFilter
[^479]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
[^480]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
[^481]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
[^482]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
[^487]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
[^488]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
[^489]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
[^490]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
[^491]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
[^492]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
19.2.2 AIMReader

This page lists supported metadata fields for the Bio-Formats AIM format reader. These fields are from the OME data model\(^5\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

19.2. Metadata fields
Supported fields

These fields are fully supported by the Bio-Formats AIM format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 22
Total unknown or missing: 453
19.2.3 AliconaReader

This page lists supported metadata fields for the Bio-Formats Alicona AL3D format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 33 of them (6%).
- Of those, Bio-Formats fully or partially converts 33 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Alicona AL3D format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Detector: ID
- Detector: Type
- DetectorSettings: ID
- DetectorSettings: Voltage
- Image: AcquisitionDate
- Image: ID
- Image: InstrumentRef
- Image: Name
- Instrument: ID
- Objective: CalibratedMagnification
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: WorkingDistance
- ObjectiveSettings: ID
- Pixels: BigEndian

534 http://www.openmicroscopy.org/site/support/ome-model/
535 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
536 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
537 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
538 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
539 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
540 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Voltage
541 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
544 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
546 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
547 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
548 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
549 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
550 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_WorkingDistance
551 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
552 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
Bio-Formats Documentation, Release 5.0.4

• Pixels : DimensionOrder\textsuperscript{553}
• Pixels : ID\textsuperscript{554}
• Pixels : Interleaved\textsuperscript{555}
• Pixels : PhysicalSizeX\textsuperscript{556}
• Pixels : PhysicalSizeY\textsuperscript{557}
• Pixels : SignificantBits\textsuperscript{558}
• Pixels : SizeC\textsuperscript{559}
• Pixels : SizeT\textsuperscript{560}
• Pixels : SizeX\textsuperscript{561}
• Pixels : SizeY\textsuperscript{562}
• Pixels : SizeZ\textsuperscript{563}
• Pixels : Type\textsuperscript{564}
• Plane : THE\textsuperscript{565}
• Plane : THE\textsuperscript{566}
• Plane : THE\textsuperscript{567}

Total supported: 33
Total unknown or missing: 442

19.2.4 GelReader

This page lists supported metadata fields for the Bio-Formats Amersham Biosciences GEL format reader.

These fields are from the OME data model\textsuperscript{568}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 21 of them (4%).
• Of those, Bio-Formats fully or partially converts 21 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Amersham Biosciences GEL format reader:

• Channel : ID\textsuperscript{569}
• Channel : SamplesPerPixel\textsuperscript{570}

\textsuperscript{553}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\textsuperscript{554}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
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\textsuperscript{568}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{569}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\textsuperscript{570}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 21
Total unknown or missing: 454

19.2.5 AmiraReader

This page lists supported metadata fields for the Bio-Formats Amira format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

571 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
573 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
574 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
575 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
576 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
577 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
579 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
580 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
581 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
582 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
583 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
584 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
585 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
586 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
588 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
589 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
590 http://www.openmicroscopy.org/site/support/ome-model/

19.2. Metadata fields
Supported fields

These fields are fully supported by the Bio-Formats Amira format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 22

Total unknown or missing: 453

591 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
592 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
593 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
595 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
596 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
597 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
598 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
599 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
600 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
603 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
604 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
605 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
606 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
607 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
608 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
609 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
610 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
611 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
612 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
19.2.6 AnalyzeReader

This page lists supported metadata fields for the Bio-Formats Analyze 7.5 format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 24 of them (5%).
- Of those, Bio-Formats fully or partially converts 24 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Analyze 7.5 format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: Description
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY

http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
• Pixels : SizeZ
• Pixels : TimeIncrement
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 24
Total unknown or missing: 451

19.2.7 AFIReader

This page lists supported metadata fields for the Bio-Formats Aperio AFI format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 30 of them (6%).
• Of those, Bio-Formats fully or partially converts 30 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Aperio AFI format reader:

• Channel : EmissionWavelength
• Channel : ExcitationWavelength
• Channel : ID
• Channel : Name
• Channel : SamplesPerPixel
• Image : AcquisitionDate
• Image : ID
• Image : InstrumentRef
• Image : Name
• Instrument : ID
• Objective : ID
This page lists supported metadata fields for the Bio-Formats Aperio SVS format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**19.2.8 SVSReader**

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 29 of them (6%).
- Of those, Bio-Formats fully or partially converts 29 (100%).

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650 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
651 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
652 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
653 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
654 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
655 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
657 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
658 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
659 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
660 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
661 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
662 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
663 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
664 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
665 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
666 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
667 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
668 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
669 http://www.openmicroscopy.org/site/support/ome-model/
Supported fields

These fields are fully supported by the Bio-Formats Aperio SVS format reader:

- Channel: EmissionWavelength
- Channel: ExcitationWavelength
- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: Description
- Image: ID
- Instrument: ID
- Objective: ID
- Objective: NominalMagnification
- ObjectiveSettings: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY

671 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
672 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
673 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
674 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
675 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
676 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
678 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
679 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveNominalMagnification
680 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
681 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
682 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
683 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
684 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
687 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
688 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
689 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
690 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
691 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
• Pixels : SizeZ\textsuperscript{694}
• Pixels : Type\textsuperscript{695}
• Plane : TheC\textsuperscript{696}
• Plane : TheT\textsuperscript{697}
• Plane : TheZ\textsuperscript{698}

Total supported: 29
Total unknown or missing: 446

19.2.9 CellWorxReader

This page lists supported metadata fields for the Bio-Formats CellWorx format reader.

These fields are from the OME data model\textsuperscript{699}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 45 of them (9%).
• Of those, Bio-Formats fully or partially converts 45 (100%).

Supported fields

These fields are fully supported by the Bio-Formats CellWorx format reader:

• Channel : EmissionWavelength\textsuperscript{700}
• Channel : ExcitationWavelength\textsuperscript{701}
• Channel : ID\textsuperscript{702}
• Channel : Name\textsuperscript{703}
• Channel : SamplesPerPixel\textsuperscript{704}
• Detector : ID\textsuperscript{705}
• DetectorSettings : Gain\textsuperscript{706}
• DetectorSettings : ID\textsuperscript{707}
• Image : AcquisitionDate\textsuperscript{708}
• Image : ID\textsuperscript{709}
• Image : InstrumentRef\textsuperscript{710}
• Image : Name\textsuperscript{711}

\textsuperscript{694}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{695}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{696}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{697}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{698}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{699}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{700}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_EmissionWavelength
\textsuperscript{701}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
\textsuperscript{702}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\textsuperscript{703}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
\textsuperscript{704}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\textsuperscript{705}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
\textsuperscript{706}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Gain
\textsuperscript{707}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
\textsuperscript{708}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\textsuperscript{709}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\textsuperscript{710}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#InstrumentRef_ID
\textsuperscript{711}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
• Instrument: ID
• Microscope: SerialNumber
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Plate: ID
• Plate: Name
• PlateAcquisition: EndTime
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount
• PlateAcquisition: StartTime
• PlateAcquisition: WellSampleRef
• Well: Column
19.2.10 AVIReader

This page lists supported metadata fields for the Bio-Formats Audio Video Interleave format reader.

These fields are from the OME data model\(^{745}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Audio Video Interleave format reader:

- Channel : ID\(^{746}\)
- Channel : SamplesPerPixel\(^{747}\)
- Image : AcquisitionDate\(^{748}\)
- Image : ID\(^{749}\)
- Image : Name\(^{750}\)
- Pixels : BigEndian\(^{751}\)
- Pixels : DimensionOrder\(^{752}\)
- Pixels : ID\(^{753}\)
- Pixels : Interleaved\(^{754}\)
- Pixels : SignificantBits\(^{755}\)

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\(^{738}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_ID
\(^{739}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Row
\(^{740}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_ID
\(^{741}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ImageRef_ID
\(^{742}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_Index
\(^{743}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_PositionX
\(^{744}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_PositionY
\(^{745}\)http://www.openmicroscopy.org/Site/support/ome-model/
\(^{746}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^{747}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^{748}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^{749}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^{750}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^{751}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^{752}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^{753}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^{754}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^{755}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
19.2. Metadata fields

Total supported: 19
Total unknown or missing: 456

19.2.11 ARFReader

This page lists supported metadata fields for the Bio-Formats ARF format reader.

These fields are from the OME data model\(^\text{765}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats ARF format reader:

- Channel: ID\(^\text{766}\)
- Channel: SamplesPerPixel\(^\text{767}\)
- Image: AcquisitionDate\(^\text{768}\)
- Image: ID\(^\text{769}\)
- Image: Name\(^\text{770}\)
- Pixels: BigEndian\(^\text{771}\)
- Pixels: DimensionOrder\(^\text{772}\)
- Pixels: ID\(^\text{773}\)

\(^{756}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\(^{757}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\(^{758}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\(^{759}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\(^{760}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^{761}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^{762}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\(^{763}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\(^{764}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\(^{765}\)http://www.openmicroscopy.org/site/support/ome-model/
\(^{766}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^{767}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^{768}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^{769}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^{770}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^{771}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^{772}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^{773}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.12 BDReader

This page lists supported metadata fields for the Bio-Formats BD Pathway format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 57 of them (12%).
- Of those, Bio-Formats fully or partially converts 57 (100%).

Supported fields

These fields are fully supported by the Bio-Formats BD Pathway format reader:

- Channel: EmissionWavelength
- Channel: ExcitationWavelength
- Channel: ID
- Channel: Name
- Channel: SamplesPerPixel
- Detector: ID

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774 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
775 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
776 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
777 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
778 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
779 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
780 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
782 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
783 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
784 http://www.openmicroscopy.org/site/support/ome-model/
786 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_EmissionWavelength
787 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
788 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
789 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
790 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
791 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
• DetectorSettings : Binning
• DetectorSettings : Gain
• DetectorSettings : ID
• DetectorSettings : Offset
• Image : AcquisitionDate
• Image : ID
• Image : InstrumentRef
• Image : Name
• Image : ROIRef
• Instrument : ID
• Objective : ID
• Objective : LensNA
• Objective : Manufacturer
• Objective : NominalMagnification
• ObjectiveSettings : ID
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Offset
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ROIRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Plate: ColumnNamingConvention
• Plate: Description
• Plate: ID
• Plate: Name
• Plate: RowNamingConvention
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount
• PlateAcquisition: WellSampleRef
• ROI: ID
• Rectangle: Height
• Rectangle: ID
• Rectangle: Width
• Rectangle: X
• Rectangle: Y
• Well: Column
• Well: ID
• Well: Row
• WellSample: ID
• WellSample: ImageRef
• WellSample: Index

Total supported: 57

Total unknown or missing: 418
19.2.13 SDTReader

This page lists supported metadata fields for the Bio-Formats SPCImage Data format reader.

These fields are from the OME data model[843]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats SPCImage Data format reader:

- Channel : ID[844]
- Channel : SamplesPerPixel[845]
- Image : AcquisitionDate[846]
- Image : ID[847]
- Image : Name[848]
- Pixels : BigEndian[849]
- Pixels : DimensionOrder[850]
- Pixels : ID[851]
- Pixels : Interleaved[852]
- Pixels : SignificantBits[853]
- Pixels : SizeC[854]
- Pixels : SizeT[855]
- Pixels : SizeX[856]
- Pixels : SizeY[857]
- Pixels : SizeZ[858]
- Pixels : Type[859]
- Plane : TheC[860]
- Plane : TheT[861]
19.2.14 BioRadGelReader

This page lists supported metadata fields for the Bio-Formats Bio-Rad GEL format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 21 of them (4%).
- Of those, Bio-Formats fully or partially converts 21 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Bio-Rad GEL format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY

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862 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
863 http://www.openmicroscopy.org/site/support/ome-model/
864 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
865 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
866 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
868 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
869 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
870 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
871 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
872 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
874 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
875 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
876 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
877 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
878 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
879 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 21
Total unknown or missing: 454

19.2.15 BioRadReader

This page lists supported metadata fields for the Bio-Formats Bio-Rad PIC format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 40 of them (8%).
• Of those, Bio-Formats fully or partially converts 40 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Bio-Rad PIC format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Detector: Gain
• Detector: ID
• Detector: Offset
• Detector: Type
• DetectorSettings: Gain
• DetectorSettings: ID
• DetectorSettings: Offset
• Experiment: ID
• Experiment: Type
• Image: AcquisitionDate

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/

19.2. Metadata fields
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Model
• Objective: NominalMagnification
• ObjectiveSettings: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC

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100| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
102| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
103| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
104| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
105| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
106| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
107| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
108| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
109| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
110| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
111| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
112| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
114| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
116| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
117| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
118| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
119| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
120| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
121| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
122| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
123| http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC

19.2. Metadata fields
19.2.16 BioRadSCNReader

This page lists supported metadata fields for the Bio-Formats Bio-Rad SCN format reader. These fields are from the OME data model\(^{24}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 29 of them (6%).
- Of those, Bio-Formats fully or partially converts 29 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Bio-Rad SCN format reader:

- Channel: ID\(^{27}\)
- Channel: SamplesPerPixel\(^{28}\)
- Detector: ID\(^{29}\)
- DetectorSettings: Binning\(^{30}\)
- DetectorSettings: Gain\(^{31}\)
- DetectorSettings: ID\(^{32}\)
- Image: AcquisitionDate\(^{33}\)
- Image: ID\(^{34}\)
- Image: Name\(^{35}\)
- Instrument: ID\(^{36}\)
- Microscope: Model\(^{37}\)
- Microscope: SerialNumber\(^{38}\)
- Pixels: BigEndian\(^{39}\)
- Pixels: DimensionOrder\(^{40}\)
- Pixels: ID\(^{41}\)

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\(^{24}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT]
\(^{25}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ]
\(^{26}\)[http://www.openmicroscopy.org/site/support/ome-model/]
\(^{27}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID]
\(^{28}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel]
\(^{29}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID]
\(^{30}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning]
\(^{31}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Gain]
\(^{32}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID]
\(^{33}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate]
\(^{34}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID]
\(^{35}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name]
\(^{36}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_ID]
\(^{37}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model]
\(^{38}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber]
\(^{39}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian]
\(^{40}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder]
\(^{41}\)[http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID]
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: Size
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: SizeC
• Pixels: SizeT
• Plane: ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 29
Total unknown or missing: 446

19.2.17 ImarisHDFReader

This page lists supported metadata fields for the Bio-Formats Bitplane Imaris 5.5 (HDF) format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 23 of them (4%).
• Of those, Bio-Formats fully or partially converts 23 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Bitplane Imaris 5.5 (HDF) format reader:

• Channel: Color
• Channel: ID
• Channel: SamplesPerPixel

942 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
944 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
945 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
946 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
947 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
948 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
949 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
950 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
951 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
952 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
954 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
955 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
956 http://www.openmicroscopy.org/site/support/ome-model/
957 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Channel_Color
958 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Channel_ID
959 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Channel_SamplesPerPixel
19.2.18 BrukerReader

This page lists supported metadata fields for the Bio-Formats Bruker format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 23
Total unknown or missing: 452
• The file format itself supports 23 of them (4%).
• Of those, Bio-Formats fully or partially converts 23 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Bruker format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Experimenter: ID
- Experimenter: Institution
- Experimenter: LastName
- Image: AcquisitionDate
- Image: ExperimenterRef
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

981 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
982 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
983 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_ID
984 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_Institution
985 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_LastName
986 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
987 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Image_ExperimenterRef_ID
988 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
989 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
990 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
991 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
992 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
993 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
994 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
995 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
996 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
997 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
998 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
999 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1000 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1001 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1002 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1003 http://www.openmicroscopy.org/Schemas/DOCUmentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
Total supported: 23
Total unknown or missing: 452

19.2.19 BurleighReader

This page lists supported metadata fields for the Bio-Formats Burleigh format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Burleigh format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
Total supported: 22
Total unknown or missing: 453

19.2.20 DNGReader

This page lists supported metadata fields for the Bio-Formats DNG format reader.

These fields are from the OME data model\(^{1027}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats DNG format reader:

- Channel: ID\(^{1028}\)
- Channel: SamplesPerPixel\(^{1029}\)
- Image: AcquisitionDate\(^{1030}\)
- Image: ID\(^{1031}\)
- Image: Name\(^{1032}\)
- Pixels: BigEndian\(^{1033}\)
- Pixels: DimensionOrder\(^{1034}\)
- Pixels: ID\(^{1035}\)
- Pixels: Interleaved\(^{1036}\)
- Pixels: SignificantBits\(^{1037}\)
- Pixels: SizeC\(^{1038}\)
- Pixels: SizeT\(^{1039}\)

\(^{1022}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^{1023}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^{1024}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\(^{1025}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\(^{1026}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\(^{1027}\)http://www.openmicroscopy.org/site/support/ome-model/
\(^{1028}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^{1029}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^{1030}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^{1031}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^{1032}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^{1033}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^{1034}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^{1035}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^{1036}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^{1037}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\(^{1038}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\(^{1039}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
Total supported: 19

19.2.21 CellomicsReader

This page lists supported metadata fields for the Bio-Formats Cellomics C01 format reader.

These fields are from the OME data model[^1047]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 31 of them (6%).
- Of those, Bio-Formats fully or partially converts 31 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Cellomics C01 format reader:

- Channel: ID[^1048]
- Channel: SamplesPerPixel[^1049]
- Image: AcquisitionDate[^1050]
- Image: ID[^1051]
- Image: Name[^1052]
- Pixels: BigEndian[^1053]
- Pixels: DimensionOrder[^1054]
- Pixels: ID[^1055]
- Pixels: Interleaved[^1056]
- Pixels: PhysicalSizeX[^1057]

[^1040]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
[^1041]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
[^1042]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
[^1043]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
[^1045]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
[^1046]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
[^1047]: http://www.openmicroscopy.org/site/support/ome-model/
[^1048]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
[^1049]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
[^1050]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
[^1051]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
[^1052]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
[^1053]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
[^1054]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
[^1056]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
• Pixels : PhysicalSizeY
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ
• Plate : ColumnNamingConvention
• Plate : ID
• Plate : Name
• Plate : RowNamingConvention
• Well : Column
• Well : ID
• Well : Row
• WellSample : ID
• WellSample : ImageRef
• WellSample : Index

Total supported: 31
Total unknown or missing: 444

19.2.22 CellSensReader

This page lists supported metadata fields for the Bio-Formats CellSens VSI format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ColumnNamingConvention
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_RowNamingConvention
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Column
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Row
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_Index

19.2. Metadata fields
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats CellSens VSI format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19

Total unknown or missing: 456

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
19.2.23 CellVoyagerReader

This page lists supported metadata fields for the Bio-Formats CellVoyager format reader. These fields are from the OME data model\(^\text{1099}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 34 of them (7%).
- Of those, Bio-Formats fully or partially converts 34 (100%).

Supported fields

These fields are fully supported by the Bio-Formats CellVoyager format reader:

- Channel: ID\(^\text{1100}\)
- Channel: Name\(^\text{1101}\)
- Channel: PinholeSize\(^\text{1102}\)
- Channel: SamplesPerPixel\(^\text{1103}\)
- Image: AcquisitionDate\(^\text{1104}\)
- Image: ID\(^\text{1105}\)
- Image: Name\(^\text{1106}\)
- Pixels: BigEndian\(^\text{1107}\)
- Pixels: DimensionOrder\(^\text{1108}\)
- Pixels: ID\(^\text{1109}\)
- Pixels: Interleaved\(^\text{1110}\)
- Pixels: SignificantBits\(^\text{1111}\)
- Pixels: SizeC\(^\text{1112}\)
- Pixels: SizeT\(^\text{1113}\)
- Pixels: SizeX\(^\text{1114}\)
- Pixels: SizeY\(^\text{1115}\)
- Pixels: SizeZ\(^\text{1116}\)
- Pixels: Type\(^\text{1117}\)

\(^\text{1099}\)http://www.openmicroscopy.org/site/support/ome-model/
\(^\text{1100}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^\text{1101}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
\(^\text{1102}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize
\(^\text{1103}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^\text{1104}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^\text{1105}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^\text{1106}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^\text{1107}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^\text{1108}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^\text{1109}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^\text{1110}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^\text{1111}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\(^\text{1112}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\(^\text{1113}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\(^\text{1114}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\(^\text{1115}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\(^\text{1116}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^\text{1117}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type

19.2. Metadata fields
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Plate: Columns
• Plate: Rows
• PlateAcquisition: EndTime
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount
• PlateAcquisition: StartTime
• Well: Column
• Well: ID
• Well: Row
• WellSample: ID
• WellSample: Index
• WellSample: PositionX
• WellSample: PositionY

Total supported: 34
Total unknown or missing: 441

19.2.24 DeltavisionReader

This page lists supported metadata fields for the Bio-Formats Deltavision format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 52 of them (10%).
• Of those, Bio-Formats fully or partially converts 52 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Deltavision format reader:

• Channel: EmissionWavelength

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/omer_xml.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/omer_xml.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/omer_xml.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#Plate_Columns
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#Plate_Rows
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#PlateAcquisition_EndTime
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#PlateAcquisition_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#PlateAcquisition_MaximumFieldCount
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#PlateAcquisition_StartTime
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#Well_Column
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#Well_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#Well_Row
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#WellSample_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#WellSample_Index
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xml.html#WellSample_PositionY
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/omer_xml.html#Channel_EmissionWavelength
• Channel : ExcitationWavelength
• Channel : ID
• Channel : NDFilter
• Channel : Name
• Channel : SamplesPerPixel
• Detector : ID
• Detector : Model
• Detector : Type
• DetectorSettings : Binning
• DetectorSettings : Gain
• DetectorSettings : ID
• DetectorSettings : ReadOutRate
• Image : AcquisitionDate
• Image : Description
• Image : ID
• Image : InstrumentRef
• Image : Name
• ImagingEnvironment : Temperature
• Instrument : ID
• Objective : CalibratedMagnification
• Objective : Correction
• Objective : ID
• Objective : Immersion
• Objective : LensNA
• Objective : Manufacturer
• Objective : Model

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1136 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
1137 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1138 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_NDFilter
1139 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
1140 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1141 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
1142 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1143 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
1144 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
1146 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
1147 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ReadOutRate
1148 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
1149 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
1150 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
1152 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
1153 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ImagingEnvironment_Temperature
1155 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
1156 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
1157 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
1158 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
1159 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
1160 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
• Objective: NominalMagnification
  • Objective: WorkingDistance
• ObjectiveSettings: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 52
Total unknown or missing: 423
19.2.25 DicomReader

This page lists supported metadata fields for the Bio-Formats DICOM format reader. These fields are from the OME data model\[1187\]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 23 of them (4%).
- Of those, Bio-Formats fully or partially converts 23 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats DICOM format reader:

- Channel : ID\[1188\]
- Channel : SamplesPerPixel\[1189\]
- Image : AcquisitionDate\[1190\]
- Image : Description\[1191\]
- Image : ID\[1192\]
- Image : Name\[1193\]
- Pixels : BigEndian\[1194\]
- Pixels : DimensionOrder\[1195\]
- Pixels : ID\[1196\]
- Pixels : Interleaved\[1197\]
- Pixels : PhysicalSizeX\[1198\]
- Pixels : PhysicalSizeY\[1199\]
- Pixels : PhysicalSizeZ\[1200\]
- Pixels : SignificantBits\[1201\]
- Pixels : SizeC\[1202\]
- Pixels : SizeT\[1203\]
- Pixels : SizeX\[1204\]
- Pixels : SizeY\[1205\]

\[1187\]http://www.openmicroscopy.org/site/support/ome-model/
\[1188\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\[1189\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\[1190\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\[1191\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
\[1192\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\[1193\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\[1194\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\[1195\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\[1196\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\[1197\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\[1198\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\[1199\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\[1200\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeZ
\[1201\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\[1202\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\[1203\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\[1204\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\[1205\]http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
19.2.26 Ecat7Reader

This page lists supported metadata fields for the Bio-Formats Ecat7 format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 23 of them (4%).
- Of those, Bio-Formats fully or partially converts 23 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Ecat7 format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: Description
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
**19.2.27 EPSReader**

This page lists supported metadata fields for the Bio-Formats Encapsulated PostScript format reader.

These fields are from the OME data model\(^\text{1235}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Encapsulated PostScript format reader:

- Channel : ID\(^\text{1236}\)
- Channel : SamplesPerPixel\(^\text{1237}\)
- Image : AcquisitionDate\(^\text{1238}\)
- Image : ID\(^\text{1239}\)
- Image : Name\(^\text{1240}\)
- Pixels : BigEndian\(^\text{1241}\)

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\(^{1224}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeZ>

\(^{1225}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits>

\(^{1226}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC>

\(^{1227}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT>

\(^{1228}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX>

\(^{1229}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY>

\(^{1230}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ>

\(^{1231}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type>

\(^{1232}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC>

\(^{1233}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT>

\(^{1234}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ>

\(^{1235}\)<http://www.openmicroscopy.org/site/support/ome-model/>

\(^{1236}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID>

\(^{1237}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel>

\(^{1238}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate>

\(^{1239}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID>

\(^{1240}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name>

\(^{1241}\)<http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian>
19.2.28 FlexReader

This page lists supported metadata fields for the Bio-Formats Evotec Flex format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 69 of them (14%).
- Of those, Bio-Formats fully or partially converts 69 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Evotec Flex format reader:

- Channel : ID
- Channel : LightSourceSettingsID
- Channel : Name
- Channel : SamplesPerPixel
- Plane : TheC
- Plane : TheT
- Plane : TheZ
• Detector: ID\textsuperscript{1260}
• Detector: Type\textsuperscript{1261}
• DetectorSettings: Binning\textsuperscript{1262}
• DetectorSettings: ID\textsuperscript{1263}
• Dichroic: ID\textsuperscript{1264}
• Dichroic: Model\textsuperscript{1265}
• Filter: FilterWheel\textsuperscript{1266}
• Filter: ID\textsuperscript{1267}
• Filter: Model\textsuperscript{1268}
• Image: AcquisitionDate\textsuperscript{1269}
• Image: ID\textsuperscript{1270}
• Image: InstrumentRef\textsuperscript{1271}
• Image: Name\textsuperscript{1272}
• Instrument: ID\textsuperscript{1273}
• Laser: ID\textsuperscript{1274}
• Laser: LaserMedium\textsuperscript{1275}
• Laser: Type\textsuperscript{1276}
• Laser: Wavelength\textsuperscript{1277}
• LightPath: DichroicRef\textsuperscript{1278}
• LightPath: EmissionFilterRef\textsuperscript{1279}
• LightPath: ExcitationFilterRef\textsuperscript{1280}
• Objective: CalibratedMagnification\textsuperscript{1281}
• Objective: Correction\textsuperscript{1282}
• Objective: ID\textsuperscript{1283}
• Objective: Immersion\textsuperscript{1284}
• Objective: LensNA\textsuperscript{1285}

\textsuperscript{1260}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
\textsuperscript{1261}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
\textsuperscript{1262}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
\textsuperscript{1263}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
\textsuperscript{1264}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Dichroic_ID
\textsuperscript{1265}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
\textsuperscript{1266}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_FilterWheel
\textsuperscript{1267}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
\textsuperscript{1268}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
\textsuperscript{1269}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\textsuperscript{1270}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\textsuperscript{1271}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#InstrumentRef_ID
\textsuperscript{1272}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\textsuperscript{1273}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_ID
\textsuperscript{1274}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSource_ID
\textsuperscript{1275}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_LaserMedium
\textsuperscript{1276}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Type
\textsuperscript{1277}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Wavelength
\textsuperscript{1278}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DichroicRef_ID
\textsuperscript{1279}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
\textsuperscript{1280}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
\textsuperscript{1281}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
\textsuperscript{1282}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
\textsuperscript{1283}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
\textsuperscript{1284}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
\textsuperscript{1285}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA

19.2. Metadata fields
• ObjectiveSettings : ID
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : DeltaT
• Plane : ExposureTime
• Plane : PositionX
• Plane : PositionY
• Plane : PositionZ
• Plane : TheC
• Plane : TheT
• Plane : TheZ
• Plate : ColumnNamingConvention
• Plate : ExternalIdentifier
• Plate : ID
• Plate : Name

1286 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
1287 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
1288 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
1289 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
1290 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1291 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
1292 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
1293 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1294 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1295 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1296 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1297 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1298 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1299 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1300 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
1301 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
1306 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1307 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1308 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ColumnNamingConvention
1309 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ExternalIdentifier
1310 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ID
1311 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Name
• Plate: RowNamingConvention
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount
• PlateAcquisition: StartTime
• PlateAcquisition: WellSampleRef
• Well: Column
• Well: ID
• Well: Row
• WellSample: ID
• WellSample: ImageRef
• WellSample: Index
• WellSample: PositionX
• WellSample: PositionY

Total supported: 69
Total unknown or missing: 406

19.2.29 FEIReader

This page lists supported metadata fields for the Bio-Formats FEI/Philips format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats FEI/Philips format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID

1312 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_RowNamingConvention
1313 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_ID
1314 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_MaximumFieldCount
1315 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_StartTime
1316 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSampleRef_ID
1317 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Column
1318 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_ID
1319 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Well_Row
1320 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_ID
1321 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/one_xsd.html#ImageRef_ID
1322 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_Index
1324 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_PositionY
1325 http://www.openmicroscopy.org/site/support/ome-model/
1326 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1327 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1328 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
• Image: Name\textsuperscript{1330}
• Pixels: BigEndian\textsuperscript{1331}
• Pixels: DimensionOrder\textsuperscript{1332}
• Pixels: ID\textsuperscript{1333}
• Pixels: Interleaved\textsuperscript{1334}
• Pixels: SignificantBits\textsuperscript{1335}
• Pixels: SizeC\textsuperscript{1336}
• Pixels: SizeT\textsuperscript{1337}
• Pixels: SizeX\textsuperscript{1338}
• Pixels: SizeY\textsuperscript{1339}
• Pixels: SizeZ\textsuperscript{1340}
• Pixels: Type\textsuperscript{1341}
• Plane: TheC\textsuperscript{1342}
• Plane: TheT\textsuperscript{1343}
• Plane: TheZ\textsuperscript{1344}

Total supported: 19
Total unknown or missing: 456

19.2.30 FEITiffReader

This page lists supported metadata fields for the Bio-Formats FEI TIFF format reader.

These fields are from the OME data model\textsuperscript{1345}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 39 of them (8%).
• Of those, Bio-Formats fully or partially converts 39 (100%).

Supported fields

These fields are fully supported by the Bio-Formats FEI TIFF format reader:

• Channel: ID\textsuperscript{1346}
• Channel: SamplesPerPixel\textsuperscript{1347}

\textsuperscript{1330}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\textsuperscript{1331}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\textsuperscript{1332}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\textsuperscript{1333}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\textsuperscript{1334}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\textsuperscript{1335}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\textsuperscript{1336}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\textsuperscript{1337}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\textsuperscript{1338}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\textsuperscript{1339}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\textsuperscript{1340}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{1341}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{1342}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{1343}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{1344}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{1345}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{1346}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\textsuperscript{1347}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
• Detector: ID
• Detector: Model
• Detector: Type
• Experimenter: ID
• Experimenter: LastName
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Microscope: Model
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: NominalMagnification
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_LastName
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• StageLabel: Name
• StageLabel: X
• StageLabel: Y
• StageLabel: Z

Total supported: 39
Total unknown or missing: 436

19.2.31 FitsReader

This page lists supported metadata fields for the Bio-Formats Flexible Image Transport System format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Flexible Image Transport System format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian

1374 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1375 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1376 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_TimeIncrement
1377 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1378 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1379 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1380 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1381 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Name
1383 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Y
1384 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Z
1385 http://www.openmicroscopy.org/site/support/ome-model/
1386 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1387 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1388 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
1390 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
1391 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian

19.2. Metadata fields
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 19
Total unknown or missing: 456

19.2.32 GatanDM2Reader

This page lists supported metadata fields for the Bio-Formats Gatan DM2 format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 30 of them (6%).
• Of those, Bio-Formats fully or partially converts 30 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Gatan DM2 format reader:

• Channel : ID
• Channel : SamplesPerPixel
• Detector : ID
• DetectorSettings : Binning

1392 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
1393 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
1394 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1395 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1396 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1397 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1398 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1399 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1400 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1401 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1402 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1403 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1404 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1405 http://www.openmicroscopy.org/site/support/ome-model/
1406 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1407 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1408 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
1409 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning

19.2. Metadata fields
Bio-Formats Documentation, Release 5.0.4

• DetectorSettings: ID
• Experimenter: FirstName
• Experimenter: ID
• Experimenter: LastName
• Image: AcquisitionDate
• Image: ExperimenterRef
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

1410 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
1411 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_FirstName
1412 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_ID
1413 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_LastName
1414 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
1415 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_RefExperimenter
1416 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
1417 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_RefID
1418 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
1420 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
1421 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
1422 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
1423 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1425 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
1426 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1427 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1428 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1429 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1430 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1431 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1432 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1433 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1434 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1435 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ

19.2. Metadata fields
Total supported: 30
Total unknown or missing: 445

19.2.33 GatanReader

This page lists supported metadata fields for the Bio-Formats Gatan Digital Micrograph format reader. These fields are from the OME data model. Bio-Formats standardizes each format's original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 36 of them (7%).
- Of those, Bio-Formats fully or partially converts 36 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Gatan Digital Micrograph format reader:
- Channel: AcquisitionMode
- Channel: ID
- Channel: SamplesPerPixel
- Detector: ID
- DetectorSettings: ID
- DetectorSettings: Voltage
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Instrument: ID
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: NominalMagnification
- ObjectiveSettings: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID\textsuperscript{1454}
- Pixels: Interleaved\textsuperscript{1455}
- Pixels: PhysicalSizeX\textsuperscript{1456}
- Pixels: PhysicalSizeY\textsuperscript{1457}
- Pixels: PhysicalSizeZ\textsuperscript{1458}
- Pixels: SignificantBits\textsuperscript{1459}
- Pixels: SizeC\textsuperscript{1460}
- Pixels: SizeT\textsuperscript{1461}
- Pixels: SizeX\textsuperscript{1462}
- Pixels: SizeY\textsuperscript{1463}
- Pixels: SizeZ\textsuperscript{1464}
- Pixels: Type\textsuperscript{1465}
- Plane: ExposureTime\textsuperscript{1466}
- Plane: PositionX\textsuperscript{1467}
- Plane: PositionY\textsuperscript{1468}
- Plane: PositionZ\textsuperscript{1469}
- Plane: TheC\textsuperscript{1470}
- Plane: TheT\textsuperscript{1471}
- Plane: TheZ\textsuperscript{1472}

Total supported: 36

Total unknown or missing: 439

19.2.34 GIFReader

This page lists supported metadata fields for the Bio-Formats Graphics Interchange Format format reader.

These fields are from the OME data model\textsuperscript{1473}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

\textsuperscript{1454} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\textsuperscript{1455} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\textsuperscript{1456} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\textsuperscript{1457} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\textsuperscript{1458} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeZ
\textsuperscript{1459} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\textsuperscript{1460} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\textsuperscript{1461} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\textsuperscript{1462} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\textsuperscript{1463} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\textsuperscript{1464} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{1465} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{1466} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
\textsuperscript{1467} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PositionX
\textsuperscript{1468} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PositionY
\textsuperscript{1469} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PositionZ
\textsuperscript{1470} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{1471} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{1472} http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{1473} http://www.openmicroscopy.org/site/support/ome-model/
Supported fields

These fields are fully supported by the Bio-Formats Graphics Interchange Format format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.35 NAFReader

This page lists supported metadata fields for the Bio-Formats Hamamatsu Aquacosmos format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

1474 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1475 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1476 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
1478 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
1479 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
1480 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
1481 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
1482 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1483 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1484 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1485 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1486 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1487 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1488 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1489 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1490 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1491 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1492 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1493 http://www.openmicroscopy.org/site/support/ome-model/
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Hamamatsu Aquacosmos format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19

Total unknown or missing: 456
19.2.36 **HISReader**

This page lists supported metadata fields for the Bio-Formats Hamamatsu HIS format reader. These fields are from the OME data model\(^1\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 27 of them (5%).
- Of those, Bio-Formats fully or partially converts 27 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Hamamatsu HIS format reader:

- Channel: ID\(^2\)
- Channel: SamplesPerPixel\(^3\)
- Detector: ID\(^4\)
- Detector: Offset\(^5\)
- Detector: Type\(^6\)
- DetectorSettings: Binning\(^7\)
- DetectorSettings: ID\(^8\)
- Image: AcquisitionDate\(^9\)
- Image: ID\(^10\)
- Image: InstrumentRef\(^11\)
- Image: Name\(^12\)
- Instrument: ID\(^13\)
- Pixels: BigEndian\(^14\)
- Pixels: DimensionOrder\(^15\)
- Pixels: ID\(^16\)
- Pixels: Interleaved\(^17\)
- Pixels: SignificantBits\(^18\)
- Pixels: SizeC\(^19\)

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\(^1\) [http://www.openmicroscopy.org/site/support/ome-model/](http://www.openmicroscopy.org/site/support/ome-model/)


\(^3\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel)


\(^6\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type)


\(^9\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate)


\(^12\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name)


\(^14\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian)


\(^17\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved)


\(^19\) [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC)
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 27
Total unknown or missing: 448

19.2.37 NDPIReader

This page lists supported metadata fields for the Bio-Formats Hamamatsu NDPI format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 21 of them (4%).
• Of those, Bio-Formats fully or partially converts 21 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Hamamatsu NDPI format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID

19.2. Metadata fields
19.2. Metadata fields

• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 21
Total unknown or missing: 454

19.2.38 HamamatsuVMSReader

This page lists supported metadata fields for the Bio-Formats Hamamatsu VMS format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 26 of them (5%).
• Of those, Bio-Formats fully or partially converts 26 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Hamamatsu VMS format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID

1550 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1552 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
1553 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1554 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1555 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1556 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1557 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1558 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1559 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1560 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1561 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1562 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1563 http://www.openmicroscopy.org/site/support/ome-model/
1564 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1565 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1566 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Objective: ID
• Objective: NominalMagnification
• ObjectiveSettings: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 26
Total unknown or missing: 449

19.2.39 HitachiReader

This page lists supported metadata fields for the Bio-Formats Hitachi format reader.

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
These fields are from the OME data model\(^\text{1590}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 31 of them (6%).
- Of those, Bio-Formats fully or partially converts 31 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Hitachi format reader:

- Channel : ID\(^\text{1591}\)
- Channel : SamplesPerPixel\(^\text{1592}\)
- Image : AcquisitionDate\(^\text{1593}\)
- Image : ID\(^\text{1594}\)
- Image : InstrumentRef\(^\text{1595}\)
- Image : Name\(^\text{1596}\)
- Instrument : ID\(^\text{1597}\)
- Microscope : Model\(^\text{1598}\)
- Microscope : SerialNumber\(^\text{1599}\)
- Objective : ID\(^\text{1600}\)
- Objective : WorkingDistance\(^\text{1601}\)
- ObjectiveSettings : ID\(^\text{1602}\)
- Pixels : BigEndian\(^\text{1603}\)
- Pixels : DimensionOrder\(^\text{1604}\)
- Pixels : ID\(^\text{1605}\)
- Pixels : Interleaved\(^\text{1606}\)
- Pixels : PhysicalSizeX\(^\text{1607}\)
- Pixels : PhysicalSizeY\(^\text{1608}\)
- Pixels : SignificantBits\(^\text{1609}\)
- Pixels : SizeC\(^\text{1610}\)

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\(^{1590}\)http://www.openmicroscopy.org/site/support/ome-model/

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19.2. Metadata fields
• Pixels: SizeT\textsuperscript{1611}
• Pixels: SizeX\textsuperscript{1612}
• Pixels: SizeY\textsuperscript{1613}
• Pixels: SizeZ\textsuperscript{1614}
• Pixels: Type\textsuperscript{1615}
• Plane: PositionX\textsuperscript{1616}
• Plane: PositionY\textsuperscript{1617}
• Plane: PositionZ\textsuperscript{1618}
• Plane: TheC\textsuperscript{1619}
• Plane: TheT\textsuperscript{1620}
• Plane: TheZ\textsuperscript{1621}

Total supported: 31

Total unknown or missing: 444

19.2.40 ICSReader

This page lists supported metadata fields for the Bio-Formats Image Cytometry Standard format reader. These fields are from the OME data model\textsuperscript{1622}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 72 of them (15%).
• Of those, Bio-Formats fully or partially converts 72 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Image Cytometry Standard format reader:

• Channel: EmissionWavelength\textsuperscript{1623}
• Channel: ExcitationWavelength\textsuperscript{1624}
• Channel: ID\textsuperscript{1625}
• Channel: Name\textsuperscript{1626}
• Channel: PinholeSize\textsuperscript{1627}
• Channel: SamplesPerPixel\textsuperscript{1628}
• Detector: ID
• Detector: Manufacturer
• Detector: Model
• Detector: Type
• DetectorSettings: Gain
• DetectorSettings: ID
• Dichroic: ID
• Dichroic: Model
• Experiment: ID
• Experiment: Type
• Experimenter: ID
• Experimenter: LastName
• Filter: ID
• Filter: Model
• FilterSet: DichroicRef
• FilterSet: EmissionFilterRef
• FilterSet: ExcitationFilterRef
• FilterSet: ID
• FilterSet: Model
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Laser: ID

1629 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
1630 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
1631 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1632 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
1634 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
1635 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Dichroic_ID
1636 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1637 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experiment_ID
1638 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experiment_Type
1639 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_ID
1640 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_LastName
1641 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
1642 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1643 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DichroicRef_ID
1644 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
1645 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
1646 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterSet_ID
1647 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1648 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
1649 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
1650 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
1653 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSource_ID
- Laser: LaserMedium
- Laser: Manufacturer
- Laser: Model
- Laser: Power
- Laser: RepetitionRate
- Laser: Type
- Laser: Wavelength
- Microscope: Manufacturer
- Microscope: Model
- Objective: CalibratedMagnification
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: LensNA
- Objective: Model
- Objective: WorkingDistance
- ObjectiveSettings: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC

1656 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
1657 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1659 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_RepetitionRate
1660 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Type
1661 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Wavelength
1662 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1663 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1664 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
1665 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
1666 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
1667 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
1668 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
1669 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
1670 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_WorkingDistance
1671 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
1672 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
1673 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
1674 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
1675 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1677 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
1679 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1680 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC

19.2. Metadata fields
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 72
Total unknown or missing: 403

19.2.41 ImaconReader

This page lists supported metadata fields for the Bio-Formats Imacon format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 23 of them (4%).
• Of those, Bio-Formats fully or partially converts 23 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Imacon format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Experimenter: FirstName

1681 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1682 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1683 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1684 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1685 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_TimeIncrement
1686 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1687 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
1688 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
1692 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1693 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1694 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1695 http://www.openmicroscopy.org/site/support/ome-model/
1696 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1697 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1698 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_FirstName
• Experimenter : ID\textsuperscript{1699}
• Experimenter : LastName\textsuperscript{1700}
• Image : AcquisitionDate\textsuperscript{1701}
• Image : ExperimenterRef\textsuperscript{1702}
• Image : ID\textsuperscript{1703}
• Image : Name\textsuperscript{1704}
• Pixels : BigEndian\textsuperscript{1705}
• Pixels : DimensionOrder\textsuperscript{1706}
• Pixels : ID\textsuperscript{1707}
• Pixels : Interleaved\textsuperscript{1708}
• Pixels : SignificantBits\textsuperscript{1709}
• Pixels : SizeC\textsuperscript{1710}
• Pixels : SizeT\textsuperscript{1711}
• Pixels : SizeX\textsuperscript{1712}
• Pixels : SizeY\textsuperscript{1713}
• Pixels : SizeZ\textsuperscript{1714}
• Pixels : Type\textsuperscript{1715}
• Plane : TheC\textsuperscript{1716}
• Plane : TheT\textsuperscript{1717}
• Plane : TheZ\textsuperscript{1718}

Total supported: 23
Total unknown or missing: 452

19.2.42 SEQReader

This page lists supported metadata fields for the Bio-Formats Image-Pro Sequence format reader.

These fields are from the OME data model\textsuperscript{1719}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

\textsuperscript{1699}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_ID
\textsuperscript{1700}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_LastName
\textsuperscript{1701}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\textsuperscript{1702}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ExperimenterRef_ID
\textsuperscript{1703}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\textsuperscript{1704}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\textsuperscript{1705}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\textsuperscript{1706}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\textsuperscript{1707}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\textsuperscript{1708}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\textsuperscript{1709}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\textsuperscript{1710}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\textsuperscript{1711}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\textsuperscript{1712}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\textsuperscript{1713}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\textsuperscript{1714}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{1715}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{1716}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{1717}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{1718}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{1719}http://www.openmicroscopy.org/site/support/ome-model/
- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Image-Pro Sequence format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : SignificantBits
- Pixels : SizeC
- Pixels : SizeT
- Pixels : SizeX
- Pixels : SizeY
- Pixels : SizeZ
- Pixels : Type
- Plane : TheC
- Plane : TheT
- Plane : TheZ

**Total supported: 19**

**Total unknown or missing: 456**

1720 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1721 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1722 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
1723 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
1724 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
1725 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
1726 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
1727 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
1728 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1729 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1730 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1731 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1732 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1733 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1734 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1735 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1736 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1737 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1738 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
19.2.43 IPWReader

This page lists supported metadata fields for the Bio-Formats Image-Pro Workspace format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 20 of them (4%).
- Of those, Bio-Formats fully or partially converts 20 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Image-Pro Workspace format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : Description
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : SignificantBits
- Pixels : SizeC
- Pixels : SizeT
- Pixels : SizeX
- Pixels : SizeY
- Pixels : SizeZ
- Pixels : Type
- Plane : TheC


19.2. Metadata fields
• Plane: TheT\textsuperscript{1758}
• Plane: TheZ\textsuperscript{1759}

Total supported: 20
Total unknown or missing: 455

19.2.44 ImagicReader

This page lists supported metadata fields for the Bio-Formats IMAGIC format reader.

These fields are from the OME data model\textsuperscript{1760}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats IMAGIC format reader:

• Channel: ID\textsuperscript{1761}
• Channel: SamplesPerPixel\textsuperscript{1762}
• Image: AcquisitionDate\textsuperscript{1763}
• Image: ID\textsuperscript{1764}
• Image: Name\textsuperscript{1765}
• Pixels: BigEndian\textsuperscript{1766}
• Pixels: DimensionOrder\textsuperscript{1767}
• Pixels: ID\textsuperscript{1768}
• Pixels: Interleaved\textsuperscript{1769}
• Pixels: PhysicalSizeX\textsuperscript{1770}
• Pixels: PhysicalSizeY\textsuperscript{1771}
• Pixels: PhysicalSizeZ\textsuperscript{1772}
• Pixels: SignificantBits\textsuperscript{1773}
• Pixels: SizeC\textsuperscript{1774}
• Pixels: SizeT\textsuperscript{1775}

\textsuperscript{1758}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{1759}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{1760}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{1761}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\textsuperscript{1762}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\textsuperscript{1763}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\textsuperscript{1764}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\textsuperscript{1765}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\textsuperscript{1766}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\textsuperscript{1767}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\textsuperscript{1768}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\textsuperscript{1769}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\textsuperscript{1770}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\textsuperscript{1771}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\textsuperscript{1772}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeZ
\textsuperscript{1773}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\textsuperscript{1774}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\textsuperscript{1775}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
• Pixels: SizeX\textsuperscript{1776}
• Pixels: SizeY\textsuperscript{1777}
• Pixels: SizeZ\textsuperscript{1778}
• Pixels: Type\textsuperscript{1779}
• Plane: TheC\textsuperscript{1780}
• Plane: TheT\textsuperscript{1781}
• Plane: TheZ\textsuperscript{1782}

Total supported: 22
Total unknown or missing: 453

19.2.45 IMODReader

This page lists supported metadata fields for the Bio-Formats IMOD format reader.

These fields are from the OME data model\textsuperscript{1783}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 44 of them (9%).
• Of those, Bio-Formats fully or partially converts 44 (100%).

Supported fields

These fields are fully supported by the Bio-Formats IMOD format reader:

• Channel: ID\textsuperscript{1784}
• Channel: SamplesPerPixel\textsuperscript{1785}
• Image: AcquisitionDate\textsuperscript{1786}
• Image: ID\textsuperscript{1787}
• Image: Name\textsuperscript{1788}
• Image: ROIRef\textsuperscript{1789}
• Pixels: BigEndian\textsuperscript{1790}
• Pixels: DimensionOrder\textsuperscript{1791}
• Pixels: ID\textsuperscript{1792}
• Pixels: Interleaved\textsuperscript{1793}

\textsuperscript{1776}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX}
\textsuperscript{1777}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY}
\textsuperscript{1778}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ}
\textsuperscript{1779}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type}
\textsuperscript{1780}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC}
\textsuperscript{1781}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT}
\textsuperscript{1782}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ}
\textsuperscript{1783}\url{http://www.openmicroscopy.org/site/support/ome-model/}
\textsuperscript{1784}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID}
\textsuperscript{1785}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel}
\textsuperscript{1786}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate}
\textsuperscript{1787}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID}
\textsuperscript{1788}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name}
\textsuperscript{1789}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROIRef_ID}
\textsuperscript{1790}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian}
\textsuperscript{1791}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder}
\textsuperscript{1792}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID}
\textsuperscript{1793}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved}
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Point: ID
• Point: StrokeColor
• Point: StrokeDashArray
• Point: StrokeWidth
• Point: TheZ
• Point: X
• Point: Y
• Polygon: ID
• Polygon: Points
• Polygon: StrokeColor
• Polygon: StrokeDashArray
• Polygon: StrokeWidth
• Polygon: TheZ

1795 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
1797 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1798 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1799 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1800 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
1801 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1802 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1803 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1804 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
1805 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1806 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1807 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
1808 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeColor
1809 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeDashArray
1810 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
1811 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ
1812 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Point_X
1813 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Point_Y
1814 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Point_z
1815 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Polygon_Points
1816 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeColor
1817 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeDashArray
1818 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
1819 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ
• Polyline: ID
• Polyline: Points
• Polyline: StrokeColor
• Polyline: StrokeDashArray
• Polyline: StrokeWidth
• Polyline: TheZ
• ROI: ID
• ROI: Name

Total supported: 44
Total unknown or missing: 431

19.2.46 OpenlabReader

This page lists supported metadata fields for the Bio-Formats Openlab LIFF format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 32 of them (6%).
• Of those, Bio-Formats fully or partially converts 32 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Openlab LIFF format reader:

• Channel: ID
• Channel: Name
• Channel: SamplesPerPixel
• Detector: ID
• Detector: Type
• DetectorSettings: Gain
• DetectorSettings: ID
• DetectorSettings: Offset
• Image: AcquisitionDate

\[1820\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
\[1821\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Polyline_Points
\[1822\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeColor
\[1823\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeDashArray
\[1824\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
\[1825\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ
\[1826\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROI_ID
\[1827\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROI_Name
\[1828\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\[1829\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
\[1830\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\[1831\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
\[1832\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
\[1833\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Gain
\[1834\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Offset
\[1835\] http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate

19.2. Metadata fields
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 32

Total unknown or missing: 443
19.2.47 OpenlabRawReader

This page lists supported metadata fields for the Bio-Formats Openlab RAW format reader.

These fields are from the OME data model\(^1\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Openlab RAW format reader:

- Channel: ID\(^2\)
- Channel: SamplesPerPixel\(^3\)
- Image: AcquisitionDate\(^4\)
- Image: ID\(^5\)
- Image: Name\(^6\)
- Pixels: BigEndian\(^7\)
- Pixels: DimensionOrder\(^8\)
- Pixels: ID\(^9\)
- Pixels: Interleaved\(^10\)
- Pixels: SignificantBits\(^11\)
- Pixels: SizeC\(^12\)
- Pixels: SizeT\(^13\)
- Pixels: SizeX\(^14\)
- Pixels: SizeY\(^15\)
- Pixels: SizeZ\(^16\)
- Pixels: Type\(^17\)
- Plane: TheC\(^18\)
- Plane: TheT\(^19\)

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\(^1\) http://www.openmicroscopy.org/site/support/ome-model/
\(^2\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^3\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^4\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^5\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^6\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^7\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^8\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^9\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^10\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^11\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\(^12\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\(^13\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\(^14\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\(^15\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\(^16\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^17\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^18\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\(^19\) http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT

19.2. Metadata fields
19.2.48 ImprovisionTiffReader

This page lists supported metadata fields for the Bio-Formats Improvision TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 25 of them (5%).
- Of those, Bio-Formats fully or partially converts 25 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Improvision TIFF format reader:

- Channel: ID
- Channel: Name
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: Description
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
This page lists supported metadata fields for the Bio-Formats OBF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format's original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats OBF format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 25

Total unknown or missing: 450
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.50 InCellReader

This page lists supported metadata fields for the Bio-Formats InCell 1000/2000 format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 67 of them (14%).
• Of those, Bio-Formats fully or partially converts 67 (100%).

Supported fields

These fields are fully supported by the Bio-Formats InCell 1000/2000 format reader:
• Channel: EmissionWavelength
• Channel: ExcitationWavelength
• Channel: ID
• Channel: Name
• Channel: SamplesPerPixel
• Detector: ID

1916 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
1917 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
1918 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
1919 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
1921 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
1922 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
1923 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
1925 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
1926 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
1927 http://www.openmicroscopy.org/site/support/ome-model/
1929 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
1930 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
1931 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
1932 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
1933 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
• Detector: Model
• Detector: Type
• DetectorSettings: Binning
• DetectorSettings: Gain
• DetectorSettings: ID
• Experiment: ID
• Experiment: Type
• Image: AcquisitionDate
• Image: Description
• Image: ExperimentRef
• Image: ID
• Image: InstrumentRef
• Image: Name
• ImagingEnvironment: Temperature
• Instrument: ID
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Manufacturer
• Objective: NominalMagnification
• ObjectiveSettings: ID
• ObjectiveSettings: RefractiveIndex
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID

19.2. Metadata fields
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Plate: ColumnNamingConvention
• Plate: ID
• Plate: Name
• Plate: RowNamingConvention
• Plate: WellOriginX
• Plate: WellOriginY
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ColumnNamingConvention
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_RowNamingConvention
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_WellOriginY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#PlateAcquisition_maximumFieldCount

19.2. Metadata fields
19.2.51 InCell3000Reader

This page lists supported metadata fields for the Bio-Formats InCell 3000 format reader.

These fields are from the OME data model. Bio-Formats standardizes each format's original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats InCell 3000 format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.52 INRReader

This page lists supported metadata fields for the Bio-Formats INR format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats INR format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian

2006 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
2007 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
2011 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
2015 http://www.openmicroscopy.org/site/support/ome-model/
2017 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2018 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2020 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
2021 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
- Pixels: DimensionOrder\textsuperscript{2022}
- Pixels: ID\textsuperscript{2023}
- Pixels: Interleaved\textsuperscript{2024}
- Pixels: PhysicalSizeX\textsuperscript{2025}
- Pixels: PhysicalSizeY\textsuperscript{2026}
- Pixels: PhysicalSizeZ\textsuperscript{2027}
- Pixels: SignificantBits\textsuperscript{2028}
- Pixels: SizeC\textsuperscript{2029}
- Pixels: SizeT\textsuperscript{2030}
- Pixels: SizeX\textsuperscript{2031}
- Pixels: SizeY\textsuperscript{2032}
- Pixels: SizeZ\textsuperscript{2033}
- Pixels: Type\textsuperscript{2034}
- Plane: TheC\textsuperscript{2035}
- Plane: TheT\textsuperscript{2036}
- Plane: TheZ\textsuperscript{2037}

Total supported: 22

Total unknown or missing: 453

19.2.53 InveonReader

This page lists supported metadata fields for the Bio-Formats Inveon format reader.

These fields are from the OME data model\textsuperscript{2038}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 30 of them (6%).
- Of those, Bio-Formats fully or partially converts 30 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Inveon format reader:
- Channel: ID\textsuperscript{2039}

\textsuperscript{2022}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\textsuperscript{2023}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\textsuperscript{2024}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\textsuperscript{2025}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\textsuperscript{2026}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\textsuperscript{2027}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeZ
\textsuperscript{2028}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\textsuperscript{2029}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\textsuperscript{2030}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\textsuperscript{2031}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\textsuperscript{2032}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\textsuperscript{2033}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{2034}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{2035}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{2036}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{2037}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{2038}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{2039}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
• Channel: SamplesPerPixel
• Experimenter: ID
• Experimenter: Institution
• Experimenter: UserName
• Image: AcquisitionDate
• Image: Description
• Image: ExperimenterRef
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Microscope: Model
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 30
Total unknown or missing: 445

19.2.54 IvisionReader

This page lists supported metadata fields for the Bio-Formats I Vision format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 34 of them (7%).
• Of those, Bio-Formats fully or partially converts 34 (100%).

Supported fields

These fields are fully supported by the Bio-Formats I Vision format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Detector: ID
• Detector: Type
• DetectorSettings: Binning
• DetectorSettings: Gain
• DetectorSettings: ID
• Image: AcquisitionDate
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Objective: Correction
• Objective: ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID

19.2. Metadata fields
• Objective: Immersion
• Objective: LensNA
• Objective: NominalMagnification
• ObjectiveSettings: ID
• ObjectiveSettings: RefractiveIndex
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 34
Total unknown or missing: 441

19.2.55 IPLabReader

This page lists supported metadata fields for the Bio-Formats IPLab format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 31 of them (6%).
• Of those, Bio-Formats fully or partially converts 31 (100%).

Supported fields

These fields are fully supported by the Bio-Formats IPLab format reader:

• Channel : ID
• Channel : SamplesPerPixel
• Image : AcquisitionDate
• Image : Description
• Image : ID
• Image : Name
• Image : ROIRef
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : TimeIncrement
• Pixels : Type
• Plane : DeltaT
• Plane : TheC

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ROIRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_TimeIncrement
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
• Plane: TheT
• Plane: TheZ
• ROI: ID
• Rectangle: Height
• Rectangle: ID
• Rectangle: Width
• Rectangle: X
• Rectangle: Y

Total supported: 31
Total unknown or missing: 444

19.2.56 JEOLReader

This page lists supported metadata fields for the Bio-Formats JEOL format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats JEOL format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved

2128 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
2129 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
2130 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROI_ID
2131 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Height
2132 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
2133 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Width
2135 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Y
2136 http://www.openmicroscopy.org/site/support/ome-model/
2137 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
2138 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2139 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2140 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
2141 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
2142 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
2143 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
2144 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
2145 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
• Pixels: SignificantBits\textsuperscript{2146}
• Pixels: SizeC\textsuperscript{2147}
• Pixels: SizeT\textsuperscript{2148}
• Pixels: SizeX\textsuperscript{2149}
• Pixels: SizeY\textsuperscript{2150}
• Pixels: SizeZ\textsuperscript{2151}
• Pixels: Type\textsuperscript{2152}
• Plane: TheC\textsuperscript{2153}
• Plane: TheT\textsuperscript{2154}
• Plane: TheZ\textsuperscript{2155}

Total supported: 19
Total unknown or missing: 456

19.2.57 JPEG2000Reader

This page lists supported metadata fields for the Bio-Formats JPEG-2000 format reader.

These fields are from the OME data model\textsuperscript{2156}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats JPEG-2000 format reader:

• Channel: ID\textsuperscript{2157}
• Channel: SamplesPerPixel\textsuperscript{2158}
• Image: AcquisitionDate\textsuperscript{2159}
• Image: ID\textsuperscript{2160}
• Image: Name\textsuperscript{2161}
• Pixels: BigEndian\textsuperscript{2162}
• Pixels: DimensionOrder\textsuperscript{2163}

\textsuperscript{2146}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\textsuperscript{2147}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\textsuperscript{2148}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\textsuperscript{2149}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\textsuperscript{2150}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\textsuperscript{2151}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{2152}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{2153}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{2154}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{2155}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{2156}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{2157}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\textsuperscript{2158}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\textsuperscript{2159}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\textsuperscript{2160}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\textsuperscript{2161}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\textsuperscript{2162}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\textsuperscript{2163}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.58 JPEGReader

This page lists supported metadata fields for the Bio-Formats JPEG format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats JPEG format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
Bio-Formats Documentation, Release 5.0.4

- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.59 JPKReader

This page lists supported metadata fields for the Bio-Formats JPK Instruments format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats JPK Instruments format reader:
- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
19.2.60 JPXReader

This page lists supported metadata fields for the Bio-Formats JPX format reader.

These fields are from the OME data model\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats JPX format reader:

- Channel : ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID}
- Image : ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name}
- Pixels : BigEndian\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian}
- Pixels : DimensionOrder\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder}
- Pixels : ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID}
- Pixels : Interleaved\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved}
- Pixels : SignificantBits\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits}
- Pixels : SizeC\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC}
- Pixels : SizeT\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT}
- Pixels : SizeX\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX}
- Pixels : SizeY\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY}
- Pixels : SizeZ\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ}
- Pixels : Type\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type}
- Plane : TheC\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC}
- Plane : TheT\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT}
- Plane : TheZ\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ}

Total supported: 19

Total unknown or missing: 456
• Channel: SamplesPerPixel
• Image: AcquisitiionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.61 KhorosReader

This page lists supported metadata fields for the Bio-Formats Khoros XV format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
Supported fields

These fields are fully supported by the Bio-Formats Khoros XV format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.62 KodakReader

This page lists supported metadata fields for the Bio-Formats Kodak Molecular Imaging format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 26 of them (5%).
- Of those, Bio-Formats fully or partially converts 26 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Kodak Molecular Imaging format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: InstrumentRef
- Image: Name
- ImagingEnvironment: Temperature
- Instrument: ID
- Microscope: Model
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
19.2.63 LiFlimReader

This page lists supported metadata fields for the Bio-Formats LI-FLIM format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 25 of them (5%).
- Of those, Bio-Formats fully or partially converts 25 (100%).

Supported fields

These fields are fully supported by the Bio-Formats LI-FLIM format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Image: ROIRef
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : DeltaT
• Plane : ExposureTime
• Plane : TheC
• Plane : TheT
• Plane : TheZ
• Polygon : ID
• Polygon : Points
• ROI : ID

Total supported: 25
Total unknown or missing: 450

19.2.64 InspectorReader

This page lists supported metadata fields for the Bio-Formats Lavision Inspector format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Lavision Inspector format reader:

• Channel : ID
• Channel : SamplesPerPixel
• Image : AcquisitionDate
• Image : ID
• Image : Name

2297 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
2298 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
2299 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
2300 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
2301 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
2302 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
2303 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
2304 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
2305 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
2306 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_Flame
2307 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ROI_Shape_ID
2308 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ROI_Polygon_Points
2309 http://www.openmicroscopy.org/site/support/ome-model/
2310 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
2311 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2312 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2314 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name

19.2. Metadata fields
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19

Total unknown or missing: 456

19.2.65 LeicaReader

This page lists supported metadata fields for the Bio-Formats Leica format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 56 of them (11%).
• Of those, Bio-Formats fully or partially converts 56 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Leica format reader:

• Channel: Color
• Channel: EmissionWavelength
• Channel: ExcitationWavelength
• Channel: ID
• Channel: Name
• Channel: PinholeSize
• Channel: SamplesPerPixel
• Detector: ID
• Detector: Offset
• Detector: Type
• Detector: Voltage
• DetectorSettings: ID
• Filter: ID
• Filter: Model
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• LightPath: EmissionFilterRef
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Model
• Objective: NominalMagnification
• Objective: SerialNumber
• ObjectiveSettings: ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Offset
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Voltage
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
- ObjectiveSettings: RefractiveIndex
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: TimeIncrement
- Pixels: Type
- Plane: DeltaT
- Plane: ExposureTime
- Plane: PositionX
- Plane: PositionY
- Plane: TheC
- Plane: TheT
- Plane: TheZ
- StageLabel: Name
- StageLabel: Z
- TransmittanceRange: CutIn
• TransmittanceRange : CutOut

Total supported: 56
Total unknown or missing: 419

19.2.66 LIFReader

This page lists supported metadata fields for the Bio-Formats Leica Image File Format format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 85 of them (17%).
• Of those, Bio-Formats fully or partially converts 85 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Leica Image File Format format reader:

• Channel : Color
• Channel : ExcitationWavelength
• Channel : ID
• Channel : LightSourceSettings: Attenuation
• Channel : LightSourceSettings: ID
• Channel : Name
• Channel : PinholeSize
• Channel : SamplesPerPixel
• Detector : ID
• Detector : Model
• Detector : Offset
• Detector : Type
• Detector : Zoom
• DetectorSettings : Gain
• DetectorSettings : ID
• DetectorSettings : Offset

2388 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Color
2389 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
2390 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_Attenuation
2391 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_ID
2392 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
2393 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize
2394 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2395 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
2396 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
2397 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Offset
2398 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
2399 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Zoom
2401 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
2402 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Offset
• Filter: ID
• Filter: Model
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: InstrumentRef
• Image: Name
• Image: ROIRef
• Instrument: ID
• Label: FontSize
• Label: ID
• Label: StrokeWidth
• Label: Text
• Label: X
• Label: Y
• Laser: ID
• Laser: LaserMedium
• Laser: Type
• Laser: Wavelength
• LightPath: EmissionFilterRef
• Line: ID
• Line: X1
• Line: X2
• Line: Y1
• Line: Y2
• Microscope: Model

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROIRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FontSize
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Text
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Label_Y
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#LightSource_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Shape_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
• Microscope: Type
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Model
• Objective: NominalMagnification
• Objective: SerialNumber
• ObjectiveSettings: ID
• ObjectiveSettings: RefractiveIndex
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: DeltaT

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Microscope_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_RefractiveIndex
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_TimeIncrement
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Polygon: ID
• Polygon: Points
• ROI: ID
• Rectangle: Height
• Rectangle: ID
• Rectangle: Width
• Rectangle: X
• Rectangle: Y
• TransmittanceRange: CutIn
• TransmittanceRange: CutOut

Total supported: 85
Total unknown or missing: 390

19.2.67 LeicaSCNReader

This page lists supported metadata fields for the Bio-Formats Leica SCN format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 33 of them (6%).
• Of those, Bio-Formats fully or partially converts 33 (100%).
Bio-Formats Documentation, Release 5.0.4

Supported fields

These fields are fully supported by the Bio-Formats Leica SCN format reader:

- Channel : ID
- Channel : IlluminationType
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : Description
- Image : ID
- Image : InstrumentRef
- Image : Name
- Instrument : ID
- Objective : CalibratedMagnification
- Objective : ID
- Objective : LensNA
- Objective : NominalMagnification
- ObjectiveSettings : ID
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : PhysicalSizeX
- Pixels : PhysicalSizeY
- Pixels : PhysicalSizeZ
- Pixels : SignificantBits
- Pixels : SizeC
- Pixels : SizeT

2473 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
2474 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_IlluminationType
2475 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2476 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2477 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
2478 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
2480 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
2482 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
2483 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
2484 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
2485 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
2486 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
2487 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
2488 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
2489 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
2490 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
2492 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
2494 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
2495 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
2496 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT

19.2. Metadata fields
19.2.68 LEOReader

This page lists supported metadata fields for the Bio-Formats LEO format reader. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 27 of them (5%).
- Of those, Bio-Formats fully or partially converts 27 (100%).

Supported fields

These fields are fully supported by the Bio-Formats LEO format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: InstrumentRef
- Image: Name
- Instrument: ID
- Objective: Correction

Total supported: 33
Total unknown or missing: 442
• Objective : ID
• Objective : Immersion
• Objective : WorkingDistance
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 27
Total unknown or missing: 448

19.2.69 L2DReader

This page lists supported metadata fields for the Bio-Formats Li-Cor L2D format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 29 of them (6%).
• Of those, Bio-Formats fully or partially converts 29 (100%).

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_WorkingDistance
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
Supported fields

These fields are fully supported by the Bio-Formats Li-Cor L2D format reader:

- Channel: ID\textsuperscript{2535}
- Channel: LightSourceSettingsID\textsuperscript{2536}
- Channel: SamplesPerPixel\textsuperscript{2537}
- Image: AcquisitionDate\textsuperscript{2538}
- Image: Description\textsuperscript{2539}
- Image: ID\textsuperscript{2540}
- Image: InstrumentRef\textsuperscript{2541}
- Image: Name\textsuperscript{2542}
- Instrument: ID\textsuperscript{2543}
- Laser: ID\textsuperscript{2544}
- Laser: LaserMedium\textsuperscript{2545}
- Laser: Type\textsuperscript{2546}
- Laser: Wavelength\textsuperscript{2547}
- Microscope: Model\textsuperscript{2548}
- Microscope: Type\textsuperscript{2549}
- Pixels: BigEndian\textsuperscript{2550}
- Pixels: DimensionOrder\textsuperscript{2551}
- Pixels: ID\textsuperscript{2552}
- Pixels: Interleaved\textsuperscript{2553}
- Pixels: SignificantBits\textsuperscript{2554}
- Pixels: SizeC\textsuperscript{2555}
- Pixels: SizeT\textsuperscript{2556}
- Pixels: SizeX\textsuperscript{2557}
- Pixels: SizeY\textsuperscript{2558}

\textsuperscript{2535}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID}
\textsuperscript{2536}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_ID}
\textsuperscript{2537}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel}
\textsuperscript{2538}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate}
\textsuperscript{2539}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description}
\textsuperscript{2540}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID}
\textsuperscript{2541}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#InstrumentRef_ID}
\textsuperscript{2542}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name}
\textsuperscript{2543}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_ID}
\textsuperscript{2544}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSource_ID}
\textsuperscript{2545}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_LaserMedium}
\textsuperscript{2546}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Type}
\textsuperscript{2547}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Wavelength}
\textsuperscript{2548}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model}
\textsuperscript{2549}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Microscope_Type}
\textsuperscript{2550}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian}
\textsuperscript{2551}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder}
\textsuperscript{2552}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID}
\textsuperscript{2553}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved}
\textsuperscript{2554}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits}
\textsuperscript{2555}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC}
\textsuperscript{2556}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT}
\textsuperscript{2557}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX}
\textsuperscript{2558}\url{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY}
This page lists supported metadata fields for the Bio-Formats Laboratory Imaging format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Laboratory Imaging format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : SignificantBits
- Pixels : SizeC
- Pixels : SizeT
- Plane : TheC
- Plane : TheT
- Plane : TheZ

Total supported: 29
Total unknown or missing: 446
• Pixels: SizeX\textsuperscript{2577}
• Pixels: SizeY\textsuperscript{2578}
• Pixels: SizeZ\textsuperscript{2579}
• Pixels: Type\textsuperscript{2580}
• Plane: TheC\textsuperscript{2581}
• Plane: TheT\textsuperscript{2582}
• Plane: TheZ\textsuperscript{2583}

Total supported: 19
Total unknown or missing: 456

19.2.71 MetamorphTiffReader

This page lists supported metadata fields for the Bio-Formats Metamorph TIFF format reader. These fields are from the OME data model\textsuperscript{2584}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 38 of them (8%).
• Of those, Bio-Formats fully or partially converts 38 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Metamorph TIFF format reader:

• Channel: ID\textsuperscript{2585}
• Channel: Name\textsuperscript{2586}
• Channel: SamplesPerPixel\textsuperscript{2587}
• Image: AcquisitionDate\textsuperscript{2588}
• Image: Description\textsuperscript{2589}
• Image: ID\textsuperscript{2590}
• Image: Name\textsuperscript{2591}
• ImagingEnvironment: Temperature\textsuperscript{2592}
• Pixels: BigEndian\textsuperscript{2593}
• Pixels: DimensionOrder\textsuperscript{2594}

\textsuperscript{2577}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\textsuperscript{2578}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\textsuperscript{2579}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\textsuperscript{2580}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\textsuperscript{2581}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\textsuperscript{2582}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\textsuperscript{2583}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\textsuperscript{2584}http://www.openmicroscopy.org/site/support/ome-model/
\textsuperscript{2585}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\textsuperscript{2586}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
\textsuperscript{2587}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\textsuperscript{2588}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\textsuperscript{2589}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
\textsuperscript{2590}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\textsuperscript{2591}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\textsuperscript{2592}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ImagingEnvironment_Temperature
\textsuperscript{2593}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\textsuperscript{2594}http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : PhysicalSizeZ
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : DeltaT
• Plane : ExposureTime
• Plane : PositionX
• Plane : PositionY
• Plane : TheC
• Plane : TheT
• Plane : TheZ
• Plate : ColumnNamingConvention
• Plate : ID
• Plate : RowNamingConvention
• Well : Column
• Well : ID
• Well : Row
• WellSample : ID
• WellSample : ImageRef
• WellSample : Index

Total supported: 38
Total unknown or missing: 437

19.2.72 MetamorphReader

This page lists supported metadata fields for the Bio-Formats Metamorph STK format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 43 of them (9%).
• Of those, Bio-Formats fully or partially converts 43 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Metamorph STK format reader:

• Channel : ID
• Channel : LightSourceSettingsID
• Channel : LightSourceSettingsWavelength
• Channel : Name
• Channel : SamplesPerPixel
• Detector : ID
• Detector : Type
• DetectorSettings : Binning
• DetectorSettings : Gain
• DetectorSettings : ID
• DetectorSettings : ReadOutRate
• Image : AcquisitionDate
• Image : Description
• Image : ID
• Image : InstrumentRef

2622 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_Index
2623 http://www.openmicroscopy.org/site/support/ome-model/
2624 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
2625 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_ID
2626 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSourceSettings_Wavelength
2627 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
2628 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2629 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
2630 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
2631 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
2632 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Gain
2633 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
2634 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ReadOutRate
2635 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2636 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
• Image: Name
• ImagingEnvironment: Temperature
• Instrument: ID
• Laser: ID
• Laser: LaserMedium
• Laser: Type
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
19.2.73 MIASReader

This page lists supported metadata fields for the Bio-Formats MIAS format reader. These fields are from the OME data model\[2667]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 64 of them (13%).
- Of those, Bio-Formats fully or partially converts 64 (100%).

Supported fields

These fields are fully supported by the Bio-Formats MIAS format reader:

- Channel: Color\[2668]
- Channel: ID\[2669]
- Channel: Name\[2670]
- Channel: SamplesPerPixel\[2671]
- Ellipse: ID\[2672]
- Ellipse: RadiusX\[2673]
- Ellipse: RadiusY\[2674]
- Ellipse: Text\[2675]
- Ellipse: TheT\[2676]
- Ellipse: TheZ\[2677]
- Ellipse: X\[2678]
- Ellipse: Y\[2679]
- Experiment: Description\[2680]
- Experiment: ID\[2681]
- Experiment: Type\[2682]
• Image: AcquisitionDate
• Image: ExperimentRef
• Image: ID
• Image: InstrumentRef
• Image: Name
• Image: ROIRef
• Instrument: ID
• Mask: FillColor
• Mask: Height
• Mask: ID
• Mask: StrokeColor
• Mask: Width
• Mask: X
• Mask: Y
• Objective: ID
• Objective: Model
• Objective: NominalMagnification
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: ExposureTime
- Plane: TheC
- Plane: TheT
- Plane: TheZ
- Plate: ColumnNamingConvention
- Plate: ExternalIdentifier
- Plate: ID
- Plate: Name
- Plate: RowNamingConvention
- PlateAcquisition: ID
- PlateAcquisition: MaximumFieldCount
- PlateAcquisition: WellSampleRef
- ROI: ID
- Well: Column
- Well: ID
- Well: Row
- WellSample: ID
- WellSample: ImageRef
- WellSample: Index

Total supported: 64

Total unknown or missing: 411
19.2.74 MicromanagerReader

This page lists supported metadata fields for the Bio-Formats Micro-Manager format reader.

These fields are from the OME data model\(^{2732}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 38 of them (8%).
- Of those, Bio-Formats fully or partially converts 38 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Micro-Manager format reader:

- Channel: ID\(^{2733}\)
- Channel: Name\(^{2734}\)
- Channel: SamplesPerPixel\(^{2735}\)
- Detector: ID\(^{2736}\)
- Detector: Manufacturer\(^{2737}\)
- Detector: Model\(^{2738}\)
- Detector: SerialNumber\(^{2739}\)
- Detector: Type\(^{2740}\)
- DetectorSettings: Binning\(^{2741}\)
- DetectorSettings: Gain\(^{2742}\)
- DetectorSettings: ID\(^{2743}\)
- DetectorSettings: Voltage\(^{2744}\)
- Image: AcquisitionDate\(^{2745}\)
- Image: Description\(^{2746}\)
- Image: ID\(^{2747}\)
- Image: InstrumentRef\(^{2748}\)
- Image: Name\(^{2749}\)
- ImagingEnvironment: Temperature\(^{2750}\)

\(^{2732}\)http://www.openmicroscopy.org/site/support/ome-model/

\(^{2733}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID

\(^{2734}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name

\(^{2735}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel

\(^{2736}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID

\(^{2737}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer

\(^{2738}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model

\(^{2739}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber

\(^{2740}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type

\(^{2741}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning

\(^{2742}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Gain

\(^{2743}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID

\(^{2744}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Voltage

\(^{2745}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate

\(^{2746}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description

\(^{2747}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID

\(^{2748}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#InstrumentRef_ID

\(^{2749}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name

\(^{2750}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ImagingEnvironment_Temperature
• Instrument: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 38
Total unknown or missing: 437

19.2.75 MINCReader

This page lists supported metadata fields for the Bio-Formats MINC MRI format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
• The file format itself supports 23 of them (4%).
• Of those, Bio-Formats fully or partially converts 23 (100%).

Supported fields

These fields are fully supported by the Bio-Formats MINC MRI format reader:

• Channel : ID
• Channel : SamplesPerPixel
• Image : AcquisitionDate
• Image : Description
• Image : ID
• Image : Name
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : PhysicalSizeZ
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ

19.2. Metadata fields
Total supported: 23
Total unknown or missing: 452

19.2.76 MRWReader

This page lists supported metadata fields for the Bio-Formats Minolta MRW format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Minolta MRW format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC

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2795 http://www.openmicroscopy.org/site/support/ome-model/
2796 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
2797 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2798 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2799 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
2800 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
2801 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
2802 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
2803 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
2804 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
2805 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
2806 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
2807 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
2808 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
2809 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
2810 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
2811 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
2812 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC

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19.2. Metadata fields
19.2.77 MNGReader

This page lists supported metadata fields for the Bio-Formats Multiple Network Graphics format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Multiple Network Graphics format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ

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2813 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
2814 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
2815 http://www.openmicroscopy.org/site/support/ome-model/
2816 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
2817 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
2818 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
2819 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
2820 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
2821 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
2822 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
2823 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
2824 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
2825 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
2826 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
2827 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
2828 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
2829 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
2830 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
19.2.78 MolecularImagingReader

This page lists supported metadata fields for the Bio-Formats Molecular Imaging format reader.

These fields are from the OME data model\(^{2835}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 21 of them (4%).
- Of those, Bio-Formats fully or partially converts 21 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Molecular Imaging format reader:

- Channel: ID\(^{2836}\)
- Channel: SamplesPerPixel\(^{2837}\)
- Image: AcquisitionDate\(^{2838}\)
- Image: ID\(^{2839}\)
- Image: Name\(^{2840}\)
- Pixels: BigEndian\(^{2841}\)
- Pixels: DimensionOrder\(^{2842}\)
- Pixels: ID\(^{2843}\)
- Pixels: Interleaved\(^{2844}\)
- Pixels: PhysicalSizeX\(^{2845}\)
- Pixels: PhysicalSizeY\(^{2846}\)
- Pixels: SignificantBits\(^{2847}\)
- Pixels: SizeC\(^{2848}\)

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\(^{2831}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^{2832}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\(^{2833}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\(^{2834}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\(^{2835}\)http://www.openmicroscopy.org/site/support/ome-model/
\(^{2836}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^{2837}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^{2838}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^{2839}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^{2840}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^{2841}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^{2842}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^{2843}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^{2844}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^{2845}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\(^{2846}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\(^{2847}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\(^{2848}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 21
Total unknown or missing: 454

19.2.79 MRCReader

This page lists supported metadata fields for the Bio-Formats Medical Research Council format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Medical Research Council format reader:
• Channel : ID
• Channel : SamplesPerPixel
• Image : AcquisitionDate
• Image : ID
• Image : Name
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved

2849 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
2850 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
2851 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
2852 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
2853 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
2854 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
2855 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
2856 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
Bio-Formats Documentation, Release 5.0.4

- Pixels : PhysicalSizeX
- Pixels : PhysicalSizeY
- Pixels : PhysicalSizeZ
- Pixels : SignificantBits
- Pixels : Size
- Pixels : SizeT
- Pixels : SizeX
- Pixels : SizeY
- Pixels : SizeZ
- Pixels : Type
- Plane : TheC
- Plane : TheT
- Plane : TheZ

Total supported: 22
Total unknown or missing: 453

19.2.80 NikonReader

This page lists supported metadata fields for the Bio-Formats Nikon NEF format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Nikon NEF format reader:
- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : ID
This page lists supported metadata fields for the Bio-Formats NIfTI format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 24 of them (5%).
- Of those, Bio-Formats fully or partially converts 24 (100%).

Supported fields

These fields are fully supported by the Bio-Formats NIfTI format reader:

- Channel : ID
- Channel : SamplesPerPixel

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2885: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
2886: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
2887: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
2888: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
2889: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
2890: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
2891: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
2892: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
2893: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
2894: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
2895: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
2897: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
2898: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
2899: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_C
2900: http://www.openmicroscopy.org/site/support/ome-model/
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 24
Total unknown or missing: 451

19.2.82 NikonElementsTiffReader

This page lists supported metadata fields for the Bio-Formats Nikon Elements TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_TimeIncrement
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 50 of them (10%).
- Of those, Bio-Formats fully or partially converts 50 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Nikon Elements TIFF format reader:

- Channel: AcquisitionMode\(^\text{2926}\)
- Channel: EmissionWavelength\(^\text{2927}\)
- Channel: ExcitationWavelength\(^\text{2928}\)
- Channel: ID\(^\text{2929}\)
- Channel: Name\(^\text{2930}\)
- Channel: PinholeSize\(^\text{2931}\)
- Channel: SamplesPerPixel\(^\text{2932}\)
- Detector: ID\(^\text{2933}\)
- Detector: Model\(^\text{2934}\)
- Detector: Type\(^\text{2935}\)
- DetectorSettings: Binning\(^\text{2936}\)
- DetectorSettings: Gain\(^\text{2937}\)
- DetectorSettings: ID\(^\text{2938}\)
- DetectorSettings: ReadOutRate\(^\text{2939}\)
- DetectorSettings: Voltage\(^\text{2940}\)
- Image: AcquisitionDate\(^\text{2941}\)
- Image: ID\(^\text{2942}\)
- Image: InstrumentRef\(^\text{2943}\)
- Image: Name\(^\text{2944}\)
- ImagingEnvironment: Temperature\(^\text{2945}\)
- Instrument: ID\(^\text{2946}\)

\(^{2926}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_AcquisitionMode

\(^{2927}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_EmissionWavelength

\(^{2928}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength

\(^{2929}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID

\(^{2930}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name

\(^{2931}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize

\(^{2932}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel

\(^{2933}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID

\(^{2934}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model

\(^{2935}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type

\(^{2936}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning

\(^{2937}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Gain

\(^{2938}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ReadOutRate

\(^{2939}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Voltage

\(^{2940}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate

\(^{2941}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID

\(^{2942}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name

\(^{2943}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#InstrumentRef_ID

\(^{2944}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_ID

\(^{2945}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ImagingEnvironment_Temperature

\(^{2946}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_ID
• Objective: CalibratedMagnification
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Model
• ObjectiveSettings: ID
• ObjectiveSettings: RefractiveIndex
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ

2947 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
2948 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
2949 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
2950 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
2951 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
2952 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
2953 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
2954 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_RefractiveIndex
2955 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
2956 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
2957 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
2958 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
2960 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
2962 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
2963 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
2964 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
2965 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
2966 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
2967 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
2968 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
2969 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 50
Total unknown or missing: 425

19.2.83 NikonTiffReader

This page lists supported metadata fields for the Bio-Formats Nikon TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 47 of them (9%).
• Of those, Bio-Formats fully or partially converts 47 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Nikon TIFF format reader:

• Channel: EmissionWavelength
• Channel: ExcitationWavelength
• Channel: ID
• Channel: PinholeSize
• Channel: SamplesPerPixel
• Detector: Gain
• Detector: ID
• Detector: Type
• Dichroic: ID
• Dichroic: Model
• Filter: ID
• Filter: Model
• Image: AcquisitionDate
• Image: Description

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_EmissionWavelength
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Dichroic_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description

19.2. Metadata fields
- Image: ID
- Image: InstrumentRef
- Image: Name
- Instrument: ID
- Laser: ID
- Laser: LaserMedium
- Laser: Model
- Laser: Type
- Laser: Wavelength
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: LensNA
- Objective: NominalMagnification
- Objective: WorkingDistance
- ObjectiveSettings: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 47
Total unknown or missing: 428

19.2.84 NativeND2Reader

This page lists supported metadata fields for the Bio-Formats Nikon ND2 format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 52 of them (10%).
• Of those, Bio-Formats fully or partially converts 52 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Nikon ND2 format reader:

• Channel : AcquisitionMode
• Channel : Color
• Channel : EmissionWavelength
• Channel : ExcitationWavelength
• Channel : ID
• Channel : Name
• Channel : PinholeSize
• Channel : SamplesPerPixel
• Detector : ID
• Detector : Model

3017 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
3018 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
3019 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
3020 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
3021 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
3022 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
3023 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
3024 http://www.openmicroscopy.org/site/support/ome-model/
3025 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_AcquisitionMode
3026 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Color
3027 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_EmissionWavelength
3028 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ExcitationWavelength
3029 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
3030 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
3031 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize
3032 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
3033 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
3034 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
• Detector: Type
• DetectorSettings: Binning
• DetectorSettings: Gain
• DetectorSettings: ID
• DetectorSettings: ReadOutRate
• DetectorSettings: Voltage
• Image: AcquisitionDate
• Image: ID
• Image: InstrumentRef
• Image: Name
• ImagingEnvironment: Temperature
• Instrument: ID
• Objective: CalibratedMagnification
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Model
• ObjectiveSettings: ID
• ObjectiveSettings: RefractiveIndex
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY

19.2. Metadata fields
supported metadata fields for the Bio-Formats NRRD format reader.

These fields are from the OME data model[^3]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

### Supported fields

These fields are fully supported by the Bio-Formats NRRD format reader:

- **Channel**: ID[^7]

[^3]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
[^4]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
[^6]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
[^7]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
[^8]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
[^14]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
[^15]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
[^16]: http://www.openmicroscopy.org/site/support/ome-model/
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.86 APLReader

This page lists supported metadata fields for the Bio-Formats Olympus APL format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 21 of them (4%).
- Of those, Bio-Formats fully or partially converts 21 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Olympus APL format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 21

Total unknown or missing: 454

103. http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
19.2.87 FV1000Reader

This page lists supported metadata fields for the Bio-Formats Olympus FV1000 format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 113 of them (23%).
- Of those, Bio-Formats fully or partially converts 113 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Olympus FV1000 format reader:

- Channel: EmissionWavelength
- Channel: ExcitationWavelength
- Channel: ID
- Channel: IlluminationType
- Channel: LightSourceSettingsID
- Channel: LightSourceSettingsWavelength
- Channel: Name
- Channel: SamplesPerPixel
- Detector: Gain
- Detector: ID
- Detector: Type
- Detector: Voltage
- DetectorSettings: ID
- Dichroic: ID
- Dichroic: Model
- Ellipse: FontSize
- Ellipse: ID
- Ellipse: RadiusX
- Ellipse: RadiusY

• Ellipse: StrokeWidth
• Ellipse: TheT
• Ellipse: TheZ
• Ellipse: Transform
• Ellipse: X
• Ellipse: Y
• Filter: ID
• Filter: Model
• Image: AcquisitionDate
• Image: ID
• Image: InstrumentRef
• Image: Name
• Image: ROIRef
• Instrument: ID
• Laser: ID
• Laser: LaserMedium
• Laser: Type
• Laser: Wavelength
• LightPath: DichroicRef
• LightPath: EmissionFilterRef
• Line: FontSize
• Line: ID
• Line: StrokeWidth
• Line: TheT
• Line: TheZ
• Line: Transform
- Line: X1
- Line: X2
- Line: Y1
- Line: Y2
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: LensNA
- Objective: Model
- Objective: NominalMagnification
- Objective: WorkingDistance
- ObjectiveSettings: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: PhysicalSizeZ
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: TimeIncrement

19.2. Metadata fields
• Pixels: Type
• Plane: DeltaT
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Point: FontSize
• Point: ID
• Point: StrokeWidth
• Point: TheT
• Point: TheZ
• Point: X
• Point: Y
• Polygon: FontSize
• Polygon: ID
• Polygon: Points
• Polygon: StrokeWidth
• Polygon: TheT
• Polygon: TheZ
• Polygon: Transform
• Polyline: FontSize
• Polyline: ID
• Polyline: Points
• Polyline: StrokeWidth

\[19.2. \text{Metadata fields}\]
19.2.88 FluoviewReader

This page lists supported metadata fields for the Bio-Formats Olympus Fluoview/ABD TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 49 of them (10%).
- Of those, Bio-Formats fully or partially converts 49 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Olympus Fluoview/ABD TIFF format reader:

- Channel: ID

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http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Transform
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROI_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FontSize
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Height
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Transform
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Width
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Y
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TransmittanceRange_CutIn
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
• Channel: Name
• Channel: SamplesPerPixel
• Detector: ID
• Detector: Manufacturer
• Detector: Model
• Detector: Type
• DetectorSettings: Gain
• DetectorSettings: ID
• DetectorSettings: Offset
• DetectorSettings: ReadOutRate
• DetectorSettings: Voltage
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: InstrumentRef
• Image: Name
• ImagingEnvironment: Temperature
• Instrument: ID
• Objective: CalibratedMagnification
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Model
• ObjectiveSettings: ID
• Pixels: BigEndian
• **Pixels**: `DimensionOrder`3264
• **Pixels**: `ID`3265
• **Pixels**: `Interleaved`3266
• **Pixels**: `PhysicalSizeX`3267
• **Pixels**: `PhysicalSizeY`3268
• **Pixels**: `PhysicalSizeZ`3269
• **Pixels**: `SignificantBits`3270
• **Pixels**: `SizeC`3271
• **Pixels**: `SizeT`3272
• **Pixels**: `SizeX`3273
• **Pixels**: `SizeY`3274
• **Pixels**: `SizeZ`3275
• **Pixels**: `TimeIncrement`3276
• **Pixels**: `Type`3277
• **Plane**: `DeltaT`3278
• **Plane**: `ExposureTime`3279
• **Plane**: `PositionX`3280
• **Plane**: `PositionY`3281
• **Plane**: `PositionZ`3282
• **Plane**: `TheC`3283
• **Plane**: `TheT`3284
• **Plane**: `TheZ`3285

**Total supported**: 49

**Total unknown or missing**: 426

### 19.2.89 ScanRReader

This page lists supported metadata fields for the Bio-Formats Olympus ScanR format reader.

These fields are from the [OME data model](http://www.openmicroscopy.org/site/support/ome-model/). Bio-Formats standardizes each format’s original metadata to and from the OME.
data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 43 of them (9%).
- Of those, Bio-Formats fully or partially converts 43 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Olympus ScanR format reader:

- Channel: ID
- Channel: Name
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: DeltaT
- Plane: ExposureTime

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3287: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
3288: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
3289: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
3290: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
3292: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
3293: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
3294: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
3295: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
3296: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
3298: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
3299: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
3300: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
3301: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
3302: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
3303: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
3304: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
3305: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
3306: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
3307: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
Bio-Formats Documentation, Release 5.0.4

• Plane: PositionX
• Plane: PositionY
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Plate: ColumnNamingConvention
• Plate: Columns
• Plate: ID
• Plate: Name
• Plate: RowNamingConvention
• Plate: Rows
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount
• PlateAcquisition: WellSampleRef
• Well: Column
• Well: ID
• Well: Row
• WellSample: ID
• WellSample: ImageRef
• WellSample: Index
• WellSample: PositionX
• WellSample: PositionY

Total supported: 43
Total unknown or missing: 432

19.2.90 SISReader

This page lists supported metadata fields for the Bio-Formats Olympus SIS TIFF format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME.

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ColumnNamingConvention
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Rows
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_Reference
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_Index
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#WellSample_PositionY
http://www.openmicroscopy.org/site/support/ome-model/
data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 33 of them (6%).
- Of those, Bio-Formats fully or partially converts 33 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Olympus SIS TIFF format reader:

- Channel: ID
- Channel: Name
- Channel: SamplesPerPixel
- Detector: ID
- Detector: Model
- Detector: Type
- DetectorSettings: ID
- Image: AcquisitionDate
- Image: ID
- Image: InstrumentRef
- Image: Name
- Instrument: ID
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: NominalMagnification
- ObjectiveSettings: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 33
Total unknown or missing: 442

19.2.91 OMETiffReader

This page lists supported metadata fields for the Bio-Formats OME-TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats OME-TIFF format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name

3353 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PixelsPhysicalSizeY
3354 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PixelsSignificantBits
3355 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PixelsSizeC
3356 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PixelsSizeT
3358 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PixelsSizeY
3360 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PixelsType
3362 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PlaneTheT
3363 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PlaneTheZ
3364 http://www.openmicroscopy.org/site/support/ome-model/
3365 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Channel_ID
3366 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Channel_SamplesPerPixel
3367 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ImageAcquisitionDate
3369 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Image_Name

19.2. Metadata fields
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.92 OMEXMLReader

This page lists supported metadata fields for the Bio-Formats OME-XML format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats OME-XML format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19

Total unknown or missing: 456

19.2.93 OxfordInstrumentsReader

This page lists supported metadata fields for the Bio-Formats Oxford Instruments format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Oxford Instruments format reader:

• Channel: ID

3404. http://www.openmicroscopy.org/site/support/ome-model/

19.2. Metadata fields
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.94 PCORAWReader

This page lists supported metadata fields for the Bio-Formats PCO-RAW format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ

19.2. Metadata fields
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 26 of them (5%).
- Of those, Bio-Formats fully or partially converts 26 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats PCO-RAW format reader:

- Channel: ID \(^{3428}\)
- Channel: SamplesPerPixel \(^{3429}\)
- Detector: ID \(^{3430}\)
- Detector: SerialNumber \(^{3431}\)
- DetectorSettings: Binning \(^{3432}\)
- DetectorSettings: ID \(^{3433}\)
- Image: AcquisitionDate \(^{3434}\)
- Image: Description \(^{3435}\)
- Image: ID \(^{3436}\)
- Image: Name \(^{3437}\)
- Instrument: ID \(^{3438}\)
- Pixels: BigEndian \(^{3439}\)
- Pixels: DimensionOrder \(^{3440}\)
- Pixels: ID \(^{3441}\)
- Pixels: Interleaved \(^{3442}\)
- Pixels: SignificantBits \(^{3443}\)
- Pixels: SizeC \(^{3444}\)
- Pixels: SizeT \(^{3445}\)
- Pixels: SizeX \(^{3446}\)
- Pixels: SizeY \(^{3447}\)
- Pixels: SizeZ \(^{3448}\)
- Pixels: Type \(^{3449}\)
• Plane: ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 26
Total unknown or missing: 449

19.2.95 PCXReader

This page lists supported metadata fields for the Bio-Formats PCX format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats PCX format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX

3450http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
3451http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
3452http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
3453http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
3454http://www.openmicroscopy.org/site/support/ome-model/
3455http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
3456http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
3457http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
3459http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
3460http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
3461http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
3462http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
3463http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
3464http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
3465http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
3466http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
3467http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.96 PDSReader

This page lists supported metadata fields for the Bio-Formats Perkin Elmer Densitometer format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 23 of them (4%).
• Of those, Bio-Formats fully or partially converts 23 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Perkin Elmer Densitometer format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeX
• Pixels : SizeY
• Pixels : SizeZ
• Pixels : Type
• Plane : PositionX
• Plane : PositionY
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 23
Total unknown or missing: 452

19.2.97 OperettaReader

This page lists supported metadata fields for the Bio-Formats PerkinElmer Operetta format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 43 of them (9%).
• Of those, Bio-Formats fully or partially converts 43 (100%).

Supported fields

These fields are fully supported by the Bio-Formats PerkinElmer Operetta format reader:
• Channel : ID
• Channel : Name
• Channel : SamplesPerPixel
• Experimenter : ID
• Experimenter : LastName

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
19.2. Metadata fields

- Image: AcquisitionDate
- Image: ExperimenterRef
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: PositionX
- Plane: PositionY
- Plane: PositionZ
- Plane: TheC
- Plane: TheT
- Plane: TheZ
- Plate: Columns
- Plate: Description
- Plate: ExternalIdentifier

3504 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
3505 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ExperimenterRef_ID
3507 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
3508 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
3509 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
3510 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
3511 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
3513 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
3514 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
3515 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
3516 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
3517 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
3518 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
3519 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
3520 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
3524 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
3525 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
3526 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
3527 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Columns
3528 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_Description
3529 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/SPW_xsd.html#Plate_ExternalIdentifier
• Plate: ID
• Plate: Name
• Plate: Rows
• PlateAcquisition: ID
• PlateAcquisition: MaximumFieldCount
• PlateAcquisition: WellSampleRef
• Well: Column
• Well: ID
• Well: Row
• WellSample: ID
• WellSample: ImageRef
• WellSample: Index

Total supported: 43

Total unknown or missing: 432

19.2.98 PerkinElmerReader

This page lists supported metadata fields for the Bio-Formats PerkinElmer format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 30 of them (6%).
• Of those, Bio-Formats fully or partially converts 30 (100%).

Supported fields

These fields are fully supported by the Bio-Formats PerkinElmer format reader:
• Channel: EmissionWavelength
• Channel: ExcitationWavelength
• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate

19.2. Metadata fields
19.2. Metadata fields

- Image: ID
- Image: InstrumentRef
- Image: Name
- Instrument: ID
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: DeltaT
- Plane: ExposureTime
- Plane: PositionX
- Plane: PositionY
- Plane: PositionZ
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 30

Total unknown or missing: 445
19.2.99 PGMReader

This page lists supported metadata fields for the Bio-Formats Portable Gray Map format reader. These fields are from the OME data model. Bio-Formats standardizes each format's original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Portable Gray Map format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT

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3576. http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.100 PSDReader

This page lists supported metadata fields for the Bio-Formats Adobe Photoshop format reader. These fields are from the OME data model. Bio-Formats standardizes each format's original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Adobe Photoshop format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type

19.2. Metadata fields
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.101 PhotoshopTiffReader

This page lists supported metadata fields for the Bio-Formats Adobe Photoshop TIFF format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Adobe Photoshop TIFF format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY

19.2. Metadata fields
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.102 PQBinReader

This page lists supported metadata fields for the Bio-Formats PicoQuant Bin format reader.

These fields are from the OME data model. Bio-Formats standardizes each format's original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 21 of them (4%).
• Of those, Bio-Formats fully or partially converts 21 (100%).

Supported fields

These fields are fully supported by the Bio-Formats PicoQuant Bin format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits

References:

3628 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
3629 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
3630 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
3631 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
3632 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
3633 http://www.openmicroscopy.org/site/support/ome-model/
3634 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
3635 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
3636 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
3638 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
3639 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
3640 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
3641 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
3642 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
3644 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
3645 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 21
Total unknown or missing: 454

19.2.103 PictReader

This page lists supported metadata fields for the Bio-Formats PICT format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats PICT format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
19.2.104 APNGReader

This page lists supported metadata fields for the Bio-Formats Animated PNG format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Animated PNG format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : ID
- Image : Name
- Pixels : BigEndian

Total supported: 19
Total unknown or missing: 456
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.105 PrairieReader

This page lists supported metadata fields for the Bio-Formats Prairie TIFF format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 45 of them (9%).
- Of those, Bio-Formats fully or partially converts 45 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Prairie TIFF format reader:
- Channel: ID
- Channel: Name
- Channel: SamplesPerPixel
- Detector: ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
• Detector: Type
• Detector: Zoom
• DetectorSettings: Gain
• DetectorSettings: ID
• DetectorSettings: Offset
• Image: AcquisitionDate
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Laser: ID
• Laser: Power
• Microscope: Model
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: LensNA
• Objective: Manufacturer
• Objective: NominalMagnification
• ObjectiveSettings: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: DeltaT
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 45
Total unknown or missing: 430

### 19.2.106 QuesantReader

This page lists supported metadata fields for the Bio-Formats Quesant AFM format reader.

These fields are from the [OME data model](http://www.openmicroscopy.org/site/support/ome-model/). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

#### Supported fields

These fields are fully supported by the Bio-Formats Quesant AFM format reader:

- Channel: ID
- Channel: SamplesPerPixel

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**3727** [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC)


**3729** [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX)


**3733** [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type)


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**3742** [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel)
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.107 NativeQTReader

This page lists supported metadata fields for the Bio-Formats QuickTime format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ

19.2. Metadata fields
• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats QuickTime format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19

Total unknown or missing: 456
19.2.108 RHKReader

This page lists supported metadata fields for the Bio-Formats RHK Technologies format reader. These fields are from the OME data model\(^{3784}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats RHK Technologies format reader:

- Channel: ID\(^{3785}\)
- Channel: SamplesPerPixel\(^{3786}\)
- Image: AcquisitionDate\(^{3787}\)
- Image: Description\(^{3788}\)
- Image: ID\(^{3789}\)
- Image: Name\(^{3790}\)
- Pixels: BigEndian\(^{3791}\)
- Pixels: DimensionOrder\(^{3792}\)
- Pixels: ID\(^{3793}\)
- Pixels: Interleaved\(^{3794}\)
- Pixels: PhysicalSizeX\(^{3795}\)
- Pixels: PhysicalSizeY\(^{3796}\)
- Pixels: SignificantBits\(^{3797}\)
- Pixels: SizeC\(^{3798}\)
- Pixels: SizeT\(^{3799}\)
- Pixels: SizeX\(^{3800}\)
- Pixels: SizeY\(^{3801}\)
- Pixels: SizeZ\(^{3802}\)
- Pixels: Type\(^{3803}\)

\(^{3784}\)http://www.openmicroscopy.org/site/support/ome-model/

\(^{3785}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID

\(^{3786}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel

\(^{3787}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate

\(^{3788}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description

\(^{3789}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID

\(^{3790}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name

\(^{3791}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian

\(^{3792}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder

\(^{3793}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID

\(^{3794}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved

\(^{3795}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX

\(^{3796}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY

\(^{3797}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits

\(^{3798}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC

\(^{3799}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT

\(^{3800}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX

\(^{3801}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY

\(^{3802}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ

\(^{3803}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type

19.2. Metadata fields
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.109 SBJGReader

This page lists supported metadata fields for the Bio-Formats SBJG format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats SBJG format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC

19.2. Metadata fields
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.110 SeikoReader

This page lists supported metadata fields for the Bio-Formats Seiko format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Seiko format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: Size
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.111 PCIReader

This page lists supported metadata fields for the Bio-Formats Compix Simple-PCI format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 29 of them (6%).
• Of those, Bio-Formats fully or partially converts 29 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Compix Simple-PCI format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Detector: ID
• Detector: Type

3840 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
3842 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
3843 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
3844 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
3845 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
3846 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
3847 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
3848 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
3849 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
3850 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
3851 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
3852 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
3853 http://www.openmicroscopy.org/site/support/ome-model/
3854 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
3855 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
3856 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
3857 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
• DetectorSettings: Binning
• DetectorSettings: ID
• Image: AcquisitionDate
• Image: ID
• Image: InstrumentRef
• Image: Name
• Instrument: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: TimeIncrement
• Pixels: Type
• Plane: DeltaT
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 29

Total unknown or missing: 446

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_TimeIncrement
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ

19.2. Metadata fields
19.2.112 SimplePCITiffReader

This page lists supported metadata fields for the Bio-Formats SimplePCI TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 33 of them (6%).
- Of those, Bio-Formats fully or partially converts 33 (100%).

Supported fields

These fields are fully supported by the Bio-Formats SimplePCI TIFF format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Detector : ID
- Detector : Model
- Detector : Type
- DetectorSettings : Binning
- DetectorSettings : ID
- Image : AcquisitionDate
- Image : Description
- Image : ID
- Image : InstrumentRef
- Image : Name
- Instrument : ID
- Objective : ID
- Objective : Immersion
- Objective : NominalMagnification
- Pixels : BigEndian
- Pixels : DimensionOrder

http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT
• Pixels : SizeZ
• Pixels : Type
• Plane : ExposureTime
• Plane : TheC
• Plane : TheT
• Plane : TheZ

Total supported: 33
Total unknown or missing: 442

19.2.113 SMCameraReader

This page lists supported metadata fields for the Bio-Formats SM Camera format reader.

These fields are from the [OME data model](http://www.openmicroscopy.org/site/support/ome-model/). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats SM Camera format reader:

- Channel : ID
- Channel : SamplesPerPixel

19.2. Metadata fields
This page lists supported metadata fields for the Bio-Formats SPIDER format reader.

These fields are from the OME data model\textsuperscript{3937}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 21 of them (4%).
- Of those, Bio-Formats fully or partially converts 21 (100%).

\begin{itemize}
  \item Image: AcquisitionDate\textsuperscript{3920}
  \item Image: ID\textsuperscript{3921}
  \item Image: Name\textsuperscript{3922}
  \item Pixels: BigEndian\textsuperscript{3923}
  \item Pixels: DimensionOrder\textsuperscript{3924}
  \item Pixels: ID\textsuperscript{3925}
  \item Pixels: Interleaved\textsuperscript{3926}
  \item Pixels: SignificantBits\textsuperscript{3927}
  \item Pixels: SizeC\textsuperscript{3928}
  \item Pixels: SizeT\textsuperscript{3929}
  \item Pixels: SizeX\textsuperscript{3930}
  \item Pixels: SizeY\textsuperscript{3931}
  \item Pixels: SizeZ\textsuperscript{3932}
  \item Pixels: Type\textsuperscript{3933}
  \item Plane: TheC\textsuperscript{3934}
  \item Plane: TheT\textsuperscript{3935}
  \item Plane: TheZ\textsuperscript{3936}
\end{itemize}

Total supported: 19

Total unknown or missing: 456

\textbf{19.2.114 SpiderReader}

This page lists supported metadata fields for the Bio-Formats SPIDER format reader.

These fields are from the OME data model\textsuperscript{3937}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g., physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 21 of them (4%).
- Of those, Bio-Formats fully or partially converts 21 (100%).
Supported fields

These fields are fully supported by the Bio-Formats SPIDER format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: PhysicalSizeX
- Pixels: PhysicalSizeY
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 21

Total unknown or missing: 454
19.2.115 TargaReader

This page lists supported metadata fields for the Bio-Formats Truevision Targa format reader.

These fields are from the [OME data model](http://www.openmicroscopy.org/site/support/ome-model/). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 20 of them (4%).
- Of those, Bio-Formats fully or partially converts 20 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Truevision Targa format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : Description
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : SignificantBits
- Pixels : SizeC
- Pixels : SizeT
- Pixels : SizeX
- Pixels : SizeY
- Pixels : SizeZ
- Pixels : Type
- Plane : TheC

---

3961 [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel)
3962 [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate)
3976 [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type)
Total supported: 20
Total unknown or missing: 455

19.2.116 TextReader

This page lists supported metadata fields for the Bio-Formats Text format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Text format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate


http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
Total supported: 19  
Total unknown or missing: 456

19.2.117 TiffReader

This page lists supported metadata fields for the Bio-Formats Tagged Image File Format format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Tagged Image File Format format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : Description
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : PhysicalSizeZ
- Pixels : SignificantBits
- Pixels : SizeC
Total supported: 22
Total unknown or missing: 453

19.2.118 TillVisionReader

This page lists supported metadata fields for the Bio-Formats TillVision format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats TillVision format reader:
• Channel: ID
• Channel: SamplesPerPixel
• Experiment: ID
• Experiment: Type
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Plane: TheC
• Plane: TheT
• Plane: TheZ

19.2. Metadata fields
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: Size
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: ExposureTime
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.119 TopometrixReader

This page lists supported metadata fields for the Bio-Formats TopoMetrix format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats TopoMetrix format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate

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http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_ExposureTime
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 22
Total unknown or missing: 453

19.2.120 TrestleReader

This page lists supported metadata fields for the Bio-Formats Trestle format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
• The file format itself supports 26 of them (5%).
• Of those, Bio-Formats fully or partially converts 26 (100%).

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/site/support/ome-model/
Supported fields

These fields are fully supported by the Bio-Formats Trestle format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Image: ROIRef
- Mask: Height
- Mask: ID
- Mask: Width
- Mask: X
- Mask: Y
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT

/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROIRef_ID
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Mask_Height
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Mask_Width
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Mask_Y
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
/http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT

19.2. Metadata fields
• Plane: TheZ
• ROI: ID

Total supported: 26
Total unknown or missing: 449

19.2.121 UBMReader

This page lists supported metadata fields for the Bio-Formats UBM format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats UBM format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROI_ID
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.122 UnisokuReader

This page lists supported metadata fields for the Bio-Formats Unisoku STM format reader. These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 22 of them (4%).
• Of those, Bio-Formats fully or partially converts 22 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Unisoku STM format reader:

• Channel: ID
• Channel: SamplesPerPixel
• Image: AcquisitionDate
• Image: Description
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: SignificantBits

19.2. Metadata fields
19.2.123 VarianFDFReader

This page lists supported metadata fields for the Bio-Formats Varian FDF format reader.

These fields are from the OME data model\footnote{http://www.openmicroscopy.org/site/support/ome-model/}. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 25 of them (5%).
- Of those, Bio-Formats fully or partially converts 25 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Varian FDF format reader:
- Channel: ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID}
- Channel: SamplesPerPixel\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel}
- Image: AcquisitionDate\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate}
- Image: ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID}
- Image: Name\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name}
- Pixels: BigEndian\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian}
- Pixels: DimensionOrder\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder}
- Pixels: ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID}

Total supported: 22
Total unknown or missing: 453
19.2.124 VGSAMReader

This page lists supported metadata fields for the Bio-Formats VG SAM format reader. These fields are from the OME data model\(^{4165}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

\(^{4148}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^{4149}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\(^{4150}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\(^{4151}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeZ
\(^{4152}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\(^{4153}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\(^{4154}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\(^{4155}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\(^{4156}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\(^{4157}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^{4158}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^{4159}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PositionX
\(^{4160}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_PositionY
\(^{4162}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\(^{4163}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\(^{4164}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\(^{4165}\)http://www.openmicroscopy.org/site/support/ome-model/
Supported fields

These fields are fully supported by the Bio-Formats VG SAM format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.125 VisitechReader

This page lists supported metadata fields for the Bio-Formats Visitech XYS format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

4166 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
4167 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
4168 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
4169 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
4170 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
4171 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
4172 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
4173 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
4174 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
4175 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
4176 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
4177 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
4178 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
4179 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
4180 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
4181 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
4182 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
4183 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
4184 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
4185 http://www.openmicroscopy.org/site/support/ome-model/
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Visitech XYS format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19

Total unknown or missing: 456
19.2.126 VolocityClippingReader

This page lists supported metadata fields for the Bio-Formats Volocity Library Clipping format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 19 of them (4%).
- Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Volocity Library Clipping format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : SignificantBits
- Pixels : SizeC
- Pixels : SizeT
- Pixels : SizeX
- Pixels : SizeY
- Pixels : SizeZ
- Pixels : Type
- Plane : TheC
- Plane : TheT

4205 http://www.openmicroscopy.org/site/support/ome-model/
4206 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
4207 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
4208 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
4210 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
4211 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
4212 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
4213 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
4214 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
4215 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
4216 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
4217 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
4218 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
4219 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
4220 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
4221 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
4222 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
4223 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.127 VolocityReader

This page lists supported metadata fields for the Bio-Formats Volocity Library format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 38 of them (8%).
- Of those, Bio-Formats fully or partially converts 38 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Volocity Library format reader:

- Channel: ID
- Channel: Name
- Channel: SamplesPerPixel
- Detector: ID
- Detector: Model
- DetectorSettings: ID
- Image: AcquisitionDate
- Image: Description
- Image: ID
- Image: InstrumentRef
- Image: Name
- Instrument: ID
- Objective: Correction
- Objective: ID
- Objective: Immersion
- Objective: NominalMagnification

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification

19.2. Metadata fields
• ObjectiveSettings: ID
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: PhysicalSizeX
• Pixels: PhysicalSizeY
• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: DeltaT
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 38
Total unknown or missing: 437

19.2.128 WATOPReader

This page lists supported metadata fields for the Bio-Formats WA Technology TOP format reader.

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_DeltaT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
These fields are from the OME data model\(^{4264}\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

**Of the 475 fields documented in the metadata summary table:**

- The file format itself supports 22 of them (4%).
- Of those, Bio-Formats fully or partially converts 22 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats WA Technology TOP format reader:

- Channel: ID\(^{4265}\)
- Channel: SamplesPerPixel\(^{4266}\)
- Image: AcquisitionDate\(^{4267}\)
- Image: Description\(^{4268}\)
- Image: ID\(^{4269}\)
- Image: Name\(^{4270}\)
- Pixels: BigEndian\(^{4271}\)
- Pixels: DimensionOrder\(^{4272}\)
- Pixels: ID\(^{4273}\)
- Pixels: Interleaved\(^{4274}\)
- Pixels: PhysicalSizeX\(^{4275}\)
- Pixels: PhysicalSizeY\(^{4276}\)
- Pixels: SignificantBits\(^{4277}\)
- Pixels: SizeC\(^{4278}\)
- Pixels: SizeT\(^{4279}\)
- Pixels: SizeX\(^{4280}\)
- Pixels: SizeY\(^{4281}\)
- Pixels: SizeZ\(^{4282}\)
- Pixels: Type\(^{4283}\)
- Plane: TheC\(^{4284}\)

\(^{4264}\)http://www.openmicroscopy.org/site/support/ome-model/
\(^{4265}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^{4266}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^{4267}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^{4268}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
\(^{4269}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^{4270}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^{4271}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^{4272}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^{4273}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^{4274}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^{4275}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\(^{4276}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
\(^{4277}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
\(^{4278}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
\(^{4279}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
\(^{4280}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
\(^{4281}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\(^{4282}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^{4283}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^{4284}\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC

19.2. Metadata fields
Total supported: 22

Total unknown or missing: 453

19.2.129 BMPReader

This page lists supported metadata fields for the Bio-Formats Windows Bitmap format reader.

These fields are from the OME data model[4287]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 21 of them (4%).
- Of those, Bio-Formats fully or partially converts 21 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Windows Bitmap format reader:

- Channel: ID[4288]
- Channel: SamplesPerPixel[4289]
- Image: AcquisitionDate[4290]
- Image: ID[4291]
- Image: Name[4292]
- Pixels: BigEndian[4293]
- Pixels: DimensionOrder[4294]
- Pixels: ID[4295]
- Pixels: Interleaved[4296]
- Pixels: PhysicalSizeX[4297]
- Pixels: PhysicalSizeY[4298]
- Pixels: SignificantBits[4299]
- Pixels: SizeC[4300]
- Pixels: SizeT[4301]
- Pixels: SizeX[4302]

19.2.130 WlzReader

This page lists supported metadata fields for the Bio-Formats Woolz format reader.

These fields are from the OME data model\(^4\). Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

- The file format itself supports 26 of them (5%).
- Of those, Bio-Formats fully or partially converts 26 (100%).

**Supported fields**

These fields are fully supported by the Bio-Formats Woolz format reader:

- Channel : ID
- Channel : SamplesPerPixel
- Image : AcquisitionDate
- Image : ID
- Image : Name
- Pixels : BigEndian
- Pixels : DimensionOrder
- Pixels : ID
- Pixels : Interleaved
- Pixels : PhysicalSizeX
- Pixels : PhysicalSizeY

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\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeX
\(^4\)http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY

19.2. Metadata fields
Bio-Formats Documentation, Release 5.0.4

• Pixels: PhysicalSizeZ
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• StageLabel: Name
• StageLabel: X
• StageLabel: Y
• StageLabel: Z

Total supported: 26
Total unknown or missing: 449

19.2.131 ZeissTIFFReader

This page lists supported metadata fields for the Bio-Formats Zeiss AxioVision TIFF format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
  • Of those, Bio-Formats fully or partially converts 19 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Zeiss AxioVision TIFF format reader:

• Channel: ID
• Channel: SamplesPerPixel

19.2. Metadata fields

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Name
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_X
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Y
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#StageLabel_Z
http://www.openmicroscopy.org/site/support/ome-model/
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
• Image: AcquisitionDate
• Image: ID
• Image: Name
• Pixels: BigEndian
• Pixels: DimensionOrder
• Pixels: ID
• Pixels: Interleaved
• Pixels: SignificantBits
• Pixels: SizeC
• Pixels: SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: TheC
• Plane: TheT
• Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.132 ZeissZVIReader

This page lists supported metadata fields for the Bio-Formats Zeiss Vision Image (ZVI) format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:

• The file format itself supports 19 of them (4%).
• Of those, Bio-Formats fully or partially converts 19 (100%).
Supported fields

These fields are fully supported by the Bio-Formats Zeiss Vision Image (ZVI) format reader:

- Channel: ID
- Channel: SamplesPerPixel
- Image: AcquisitionDate
- Image: ID
- Image: Name
- Pixels: BigEndian
- Pixels: DimensionOrder
- Pixels: ID
- Pixels: Interleaved
- Pixels: SignificantBits
- Pixels: SizeC
- Pixels: SizeT
- Pixels: SizeX
- Pixels: SizeY
- Pixels: SizeZ
- Pixels: Type
- Plane: TheC
- Plane: TheT
- Plane: TheZ

Total supported: 19
Total unknown or missing: 456

19.2.133 ZeissCZIReader

This page lists supported metadata fields for the Bio-Formats Zeiss CZI format reader.

These fields are from the OME data model. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

[^4357]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
[^4358]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
[^4359]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
[^4361]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
[^4362]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
[^4363]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
[^4364]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
[^4365]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
[^4366]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
[^4367]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
[^4368]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
[^4369]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeX
[^4370]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeY
[^4371]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeZ
[^4372]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type
[^4374]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheT
[^4375]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Plane_TheZ
[^4376]: http://www.openmicroscopy.org/site/support/ome-model/
Of the 475 fields documented in the metadata summary table:

- The file format itself supports 158 of them (33%).
- Of those, Bio-Formats fully or partially converts 158 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Zeiss CZI format reader:

- Arc: LotNumber
- Arc: Manufacturer
- Arc: Model
- Arc: Power
- Arc: SerialNumber
- Channel: AcquisitionMode
- Channel: Color
- Channel: EmissionWavelength
- Channel: ExcitationWavelength
- Channel: FilterSetRef
- Channel: Fluor
- Channel: ID
- Channel: IlluminationType
- Channel: Name
- Channel: PinholeSize
- Channel: SamplesPerPixel
- Detector: AmplificationGain
- Detector: Gain
- Detector: ID
- Detector: LotNumber
- Detector: Manufacturer
- Detector: Model
• Detector: Offset
• Detector: SerialNumber
• Detector: Type
• Detector: Zoom
• DetectorSettings: Binning
• DetectorSettings: Gain
• DetectorSettings: ID
• Dichroic: ID
• Dichroic: LotNumber
• Dichroic: Manufacturer
• Dichroic: Model
• Dichroic: SerialNumber
• Ellipse: ID
• Ellipse: RadiusX
• Ellipse: RadiusY
• Ellipse: Text
• Ellipse: X
• Ellipse: Y
• Experimenter: Email
• Experimenter: FirstName
• Experimenter: ID
• Experimenter: Institution
• Experimenter: LastName
• Experimenter: MiddleName
• Experimenter: UserName
• Filament: LotNumber

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Offset
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Zoom
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Dichroic_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_LotNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Ellipse_RadiusY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Text
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Ellipse_Y
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_Email
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_FirstName
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_Institution
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_LastName
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_MiddleName
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Experimenter_UserName

19.2. Metadata fields
Bio-Formats Documentation, Release 5.0.4

- Filament: Manufacturer
- Filament: Model
- Filament: Power
- Filament: Serial Number
- Filter: Filter Wheel
- Filter: ID
- Filter: Lot Number
- Filter: Manufacturer
- Filter: Model
- Filter: Serial Number
- Filter: Type
- FilterSet: Dichroic Ref
- FilterSet: Emission Filter Ref
- FilterSet: Excitation Filter Ref
- FilterSet: ID
- FilterSet: Lot Number
- FilterSet: Manufacturer
- FilterSet: Model
- FilterSet: Serial Number
- Image: Acquisition Date
- Image: Description
- Image: Experimenter Ref
- Image: ID
- Image: Instrument Ref
- Image: Name
- Image: ROI Ref

http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_FilterWheel
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DichroicRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ExperimenterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROIRef_ID

19.2. Metadata fields
• ImagingEnvironment : AirPressure
• ImagingEnvironment : CO2Percent
• ImagingEnvironment : Humidity
• ImagingEnvironment : Temperature
• Instrument : ID
• Laser : LotNumber
• Laser : Manufacturer
• Laser : Model
• Laser : Power
• Laser : SerialNumber
• LightEmittingDiode : LotNumber
• LightEmittingDiode : Manufacturer
• LightEmittingDiode : Model
• LightEmittingDiode : Power
• LightEmittingDiode : SerialNumber
• Line : ID
• Line : Text
• Line : X1
• Line : X2
• Line : Y1
• Line : Y2
• Microscope : LotNumber
• Microscope : Manufacturer
• Microscope : Model
• Microscope : SerialNumber
• Microscope : Type

19.2. Metadata fields
• Objective : CalibratedMagnification
• Objective : Correction
• Objective : ID
• Objective : Immersion
• Objective : Iris
• Objective : LensNA
• Objective : LotNumber
• Objective : Manufacturer
• Objective : Model
• Objective : NominalMagnification
• Objective : SerialNumber
• Objective : WorkingDistance
• ObjectiveSettings : CorrectionCollar
• ObjectiveSettings : ID
• ObjectiveSettings : Medium
• ObjectiveSettings : RefractiveIndex
• Pixels : BigEndian
• Pixels : DimensionOrder
• Pixels : ID
• Pixels : Interleaved
• Pixels : PhysicalSizeX
• Pixels : PhysicalSizeY
• Pixels : PhysicalSizeZ
• Pixels : SignificantBits
• Pixels : SizeC
• Pixels : SizeT

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19.2. Metadata fields

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http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_CalibratedMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Iris
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_LotNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Manufacturer
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_SerialNumber
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_WorkingDistance
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_CorrectionCollar
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_Medium
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_RefractiveIndex
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Interleaved
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_PhysicalSizeY
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SignificantBits
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeC
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_SizeT
• Pixels: SizeX
• Pixels: SizeY
• Pixels: SizeZ
• Pixels: Type
• Plane: DeltaT
• Plane: ExposureTime
• Plane: PositionX
• Plane: PositionY
• Plane: PositionZ
• Plane: TheC
• Plane: TheT
• Plane: TheZ
• Polygon: ID
• Polygon: Points
• Polygon: Text
• Polyline: ID
• Polyline: Points
• Polyline: Text
• ROI: Description
• ROI: ID
• ROI: Name
• Rectangle: Height
• Rectangle: ID
• Rectangle: Text
• Rectangle: Width
• Rectangle: X
19.2.134 ZeissLSMReader

This page lists supported metadata fields for the Bio-Formats Zeiss Laser-Scanning Microscopy format reader. These fields are from the OME data model[^535]. Bio-Formats standardizes each format’s original metadata to and from the OME data model so that you can work with a particular piece of metadata (e.g. physical width of the image in microns) in a format-independent way.

Of the 475 fields documented in the metadata summary table:
- The file format itself supports 101 of them (21%).
- Of those, Bio-Formats fully or partially converts 101 (100%).

Supported fields

These fields are fully supported by the Bio-Formats Zeiss Laser-Scanning Microscopy format reader:

- Channel : Color[^536]
- Channel : ID[^537]
- Channel : Name[^538]
- Channel : PinholeSize[^539]
- Channel : SamplesPerPixel[^540]
- Detector : AmplificationGain[^541]
- Detector : Gain[^542]
- Detector : ID[^543]
- Detector : Type[^544]
- Detector : Zoom[^545]
- DetectorSettings : Binning[^546]

[^529]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Rectangle_Y
[^530]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TransmittanceRange_CutIn
[^531]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TransmittanceRange_CutInTolerance
[^534]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#TransmittanceRange_Transmittance
[^535]: http://www.openmicroscopy.org/site/support/ome-model/
[^536]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Color
[^537]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_ID
[^538]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_Name
[^539]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_PinholeSize
[^540]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Channel_SamplesPerPixel
[^541]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_AmplificationGain
[^543]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_ID
[^544]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Type
[^545]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Detector_Zoom
[^546]: http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_Binning
• DetectorSettings: ID
• Dichroic: ID
• Dichroic : Model
• Ellipse : FontSize
• Ellipse : ID
• Ellipse : RadiusX
• Ellipse : RadiusY
• Ellipse : StrokeWidth
• Ellipse : Transform
• Ellipse : X
• Ellipse : Y
• Experimenter : ID
• Experimenter : UserName
• Filter : ID
• Filter : Model
• Filter : Type
• Image : AcquisitionDate
• Image : Description
• Image : ID
• Image : InstrumentRef
• Image : Name
• Image : ROIRef
• Instrument : ID
• Label : FontSize
• Label : ID
• Label : StrokeWidth

4547 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DetectorSettings_ID
4548 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Dichroic_ID
4549 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
4550 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FontSize
4551 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
4553 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Ellipse_RadiusY
4554 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
4555 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Transform
4557 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Ellipse_Y
4558 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_ID
4559 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Experimenter_UserName
4560 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_ID
4561 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
4562 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Filter_Type
4563 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
4564 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
4565 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_ID
4567 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Instrument_Name
4568 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ROIRef_ID
4569 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
4570 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_AcquisitionDate
4571 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Description
4573 http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Image_Name
• Label: Text
• Label: X
• Label: Y
• Laser: ID
• Laser: LaserMedium
• Laser: Model
• Laser: Type
• Laser: Wavelength
• LightPath: DichroicRef
• LightPath: EmissionFilterRef
• Line: FontSize
• Line: ID
• Line: StrokeWidth
• Line: X1
• Line: X2
• Line: Y1
• Line: Y2
• Objective: Correction
• Objective: ID
• Objective: Immersion
• Objective: Iris
• Objective: LensNA
• Objective: NominalMagnification
• ObjectiveSettings: ID
• Pixels: BigEndian
• Pixels: DimensionOrder

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http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_Text
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Label_Y
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#LightSource_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ManufacturerSpec_Model
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Laser_Type
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#DichroicRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#FilterRef_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FontSize
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Line_X1
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Line_X2
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Line_Y1
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Correction
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Immersion
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_Iris
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_LensNA
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Objective_NominalMagnification
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#ObjectiveSettings_ID
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_BigEndian
http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_DimensionOrder
- **Pixels**: ID
- **Pixels**: Interleaved
- **Pixels**: PhysicalSizeX
- **Pixels**: PhysicalSizeY
- **Pixels**: PhysicalSizeZ
- **Pixels**: SignificantBits
- **Pixels**: SizeC
- **Pixels**: SizeT
- **Pixels**: SizeX
- **Pixels**: SizeY
- **Pixels**: SizeZ
- **Pixels**: TimeIncrement
- **Pixels**: Type
- **Plane**: DeltaT
- **Plane**: PositionX
- **Plane**: PositionY
- **Plane**: PositionZ
- **Plane**: TheC
- **Plane**: TheT
- **Plane**: TheZ
- **Polygon**: FontSize
- **Polygon**: ID
- **Polygon**: Points
- **Polygon**: StrokeWidth
- **Polyline**: FontSize
- **Polyline**: ID

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4611 [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ome_xsd.html#Pixels_Type)
4622 [http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth](http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth)
- Polyline: Points\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Polyline_Points}
- Polyline: StrokeWidth\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_StrokeWidth}
- ROI: ID\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#ROI_ID}
- Rectangle: FontSize\footnote{http://www.openmicroscopy.org/Schemas/Documentation/Generated/OME-2013-06/ROI_xsd.html#Shape_FontSize}
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