Bio-Formats and NGFF

David Gault
University of Dundee

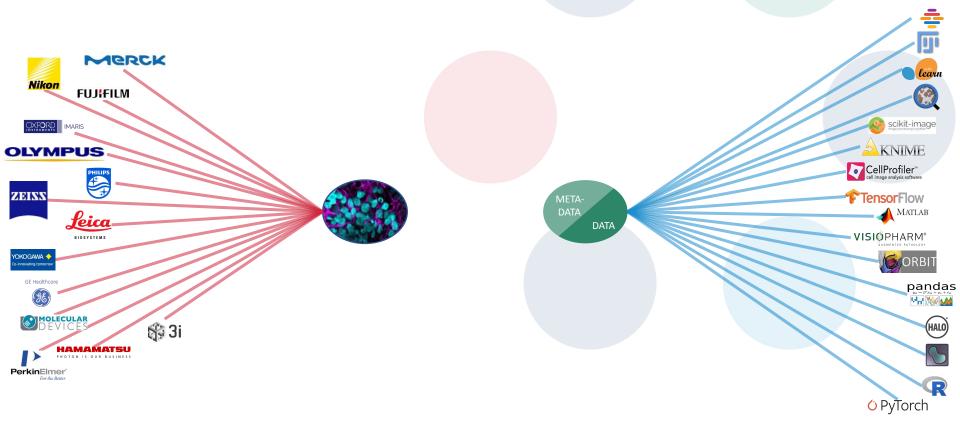
The OME Consortium openmicroscopy.org
@openmicroscopy





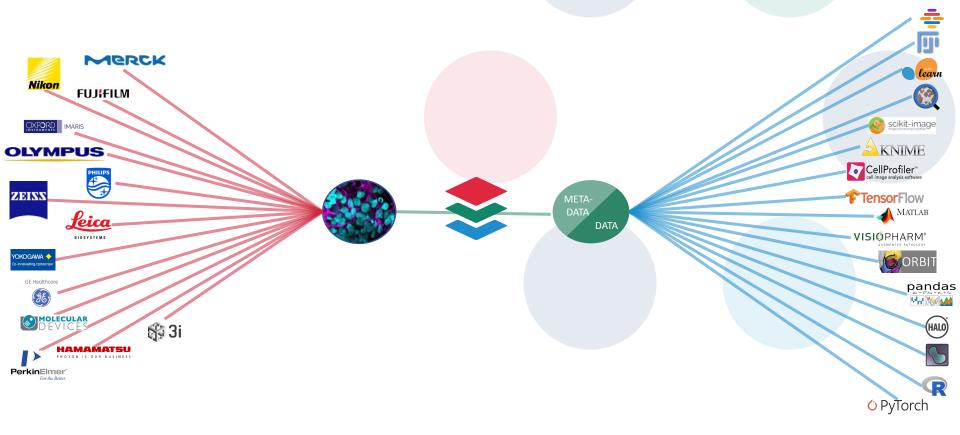


Proprietary Formats



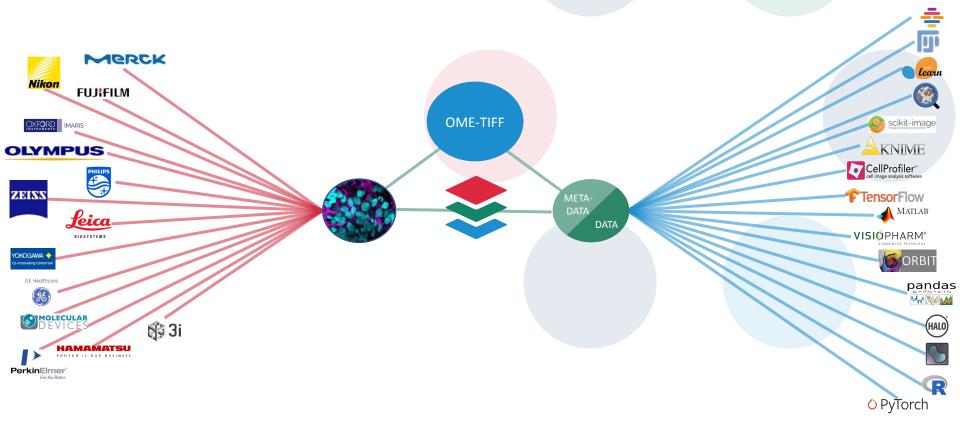


Proprietary Formats



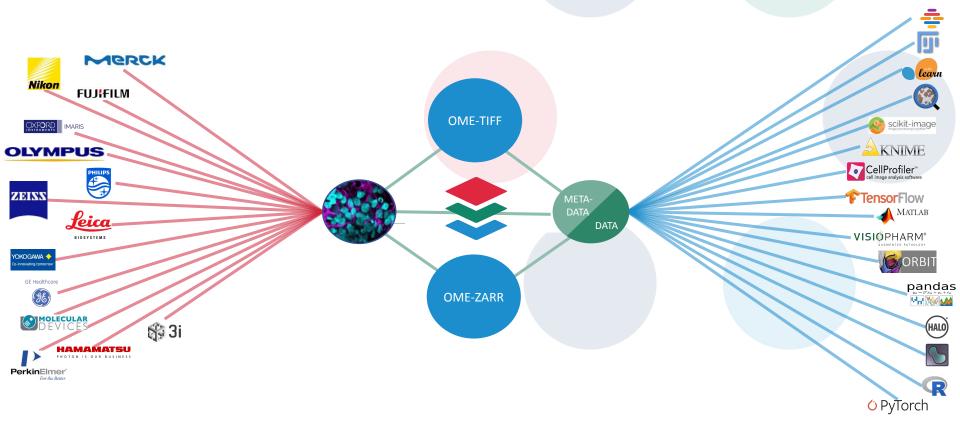


Open Standard Formats



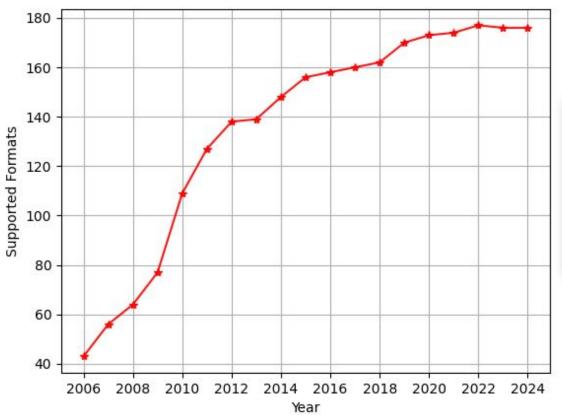


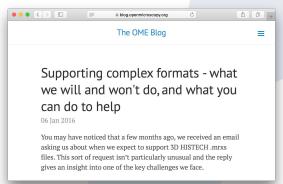
Open Standard Formats





Supporting complex formats





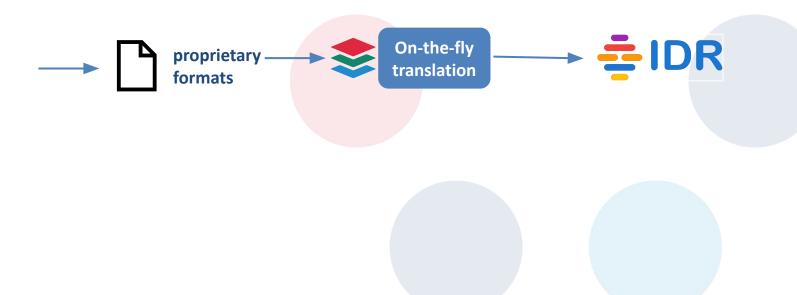




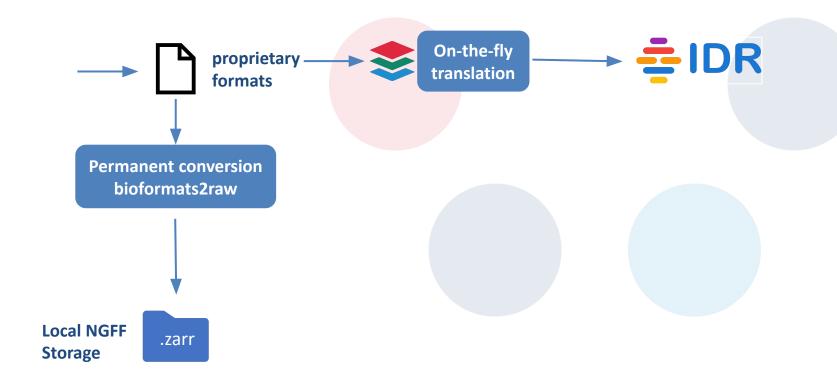
- o 7.3.0 (April 2024)
 - Java 21 compatibility
- 7.2.0 (February 2024)
 - o format improvements for Aperio SVS, Olympus OIR and Zeiss CZI and more
- o 7.1.0 (December 2023)
 - API additions for compressed tiles
- o 7.0.1 (October 2023)
 - o Format improvements for Leica LIF, NDPI, TillVision, Gatan DM3, DICOM and more
- o 7.0.0 (August 2023)
 - Removed a number of deprecated components
 - Added support for dual personality DICOM
- o 6.14.0 (July 2023)
 - Format improvements for CV7000/CV8000, KLB, MicroManager
- o 6.13.0 (May 2023)
 - Dependency updates and API additions



NGFF workflow for IDR

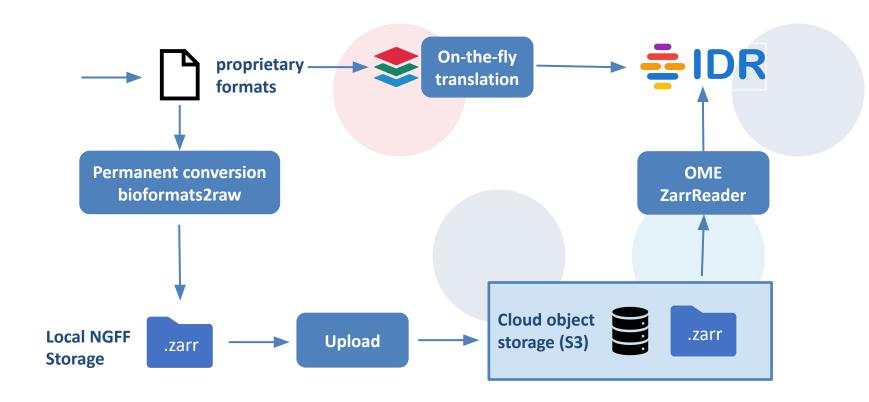


NGFF workflow for IDR





NGFF workflow for IDR





Reading NGFF

OME ZarrReader

- A traditional Bio-Formats style reader for OME-Zarr
- o Released as an external reader which can be used with Bio-Formats
- Development is driven by IDR
- o Future NGFF spec updates will be supported via new releases
- https://github.com/dgault/ZarrReader

OMERO Plus

- An OMERO specific implementation
- o Makes use of OMERO Zarr Pixel Buffer
- https://github.com/glencoesoftware/omero-zarr-pixel-buffer



Converting to NGFF

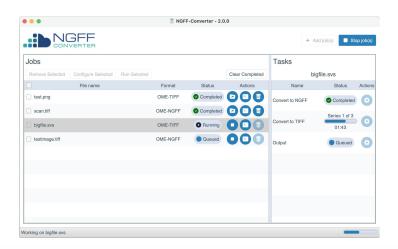
- Glencoe Software has developed new tools specifically for NGFF
 - o Bioformats2raw
 - Raw2ometiff

Formats

- NGFF Converter
- Benefit from parallel reading and writing for improved performance
- Especially useful for whole slide images
- Automatically generates down-sampled resolutions for image pyramids









EASY TO USE INTERFACE

The intuitive interface was designed with users in mind, providing the essential tools and functionality.



CLOUD OPTIMIZED

OME-NGFF is the cloud-friendly multi-dimensional bioimaging data format, allowing storage and analysis of bioimaging data in the cloud.



FULL CONTROL

Customizable conversion options for the expert user.



UNIFIED DATA

Support for novel OME-NGFF and classic OME-TIFF, both options for unifying data into a common, open format for streamlined data access and analysis.

Future Plans

- Continued development for Bio-Formats
 - Better support for reading and writing 3D data
 - Improved performance for data such as HCS
- Ongoing support for OME ZarrReader
- Support for Zarr in Java ecosystem
 - All the Java implementations rely on the JZarr library
 - The Java tooling must evolve alongside the NGFF spec

Thanks to all contributors



Jason Swedlow

Frances Wong

Dominik Lindner

Josh Moore

Jean-Marie Burel

Petr Walczysko

Will Moore

David Gault

Khaled Mohamed



Bio-Formats contributors

Shaquille Louisa - Lambert Instruments

Chris Allen

Andreas Knah

Alexandr Virodov

Melissa Linkert

Sebastien Besson

Kevin Kozlowski

Maadhay Kothari

Nicolas Chiaruttini - EPFL

Emil Rozbicki

Chris MacLeod

Can Gokhan Dogan

Erin Diel

David Stirling

Peter Haub - DIP Systems

Kyle Balis-West

Stephen S

Stephan Wagner-Conrad - Carl Zeiss Microscopy

Marc Bruce

Liza Unson

Tim Blackmore - WEHI

Lindsay Durward

Muhanad Zahra

Edward Scanlon - Proscia

Nichola Haxton



Presentations available @

https://downloads.openmicroscopy.org/ presentations/2024/Dundee