

# Talk Outline

- Thank you!
- This Meeting...
- The Problem
- Our Progress
- Funding...
- Future Priorities...

# Thank you!!!

- *Institut Pasteur*

- Christiane Pacaud
- Nathalie Aulner
- Anne Danckaert
- Sebastien Simard
- Spencer Shorte

- *University of Dundee*

- June Matthew
- Wilma Woudenberg

- *The OME Consortium*

Thank you!!!

**wellcome**trust



# OME Consortium

- Dundee – Jason Swedlow, Colin Blackburn, Jean-Marie Burel, Mark Carroll, Gus Ferguson, Helen Flynn, Kenny Gillen, Roger Leigh, Simon Li, Dominik Lindner, Josh Moore, Will Moore, Balaji Ramalingam, Gabriella Rustici, Aleksandra Tarkowska, Petr Walczysko, Eleanor Williams *and you?!?*
- University of Wisconsin, Madison (LOCI) - Kevin Eliceiri, Curtis Rueden, Mark Hiner
- UT Southwestern – Gaudenz Danuser, Sebastian Besson
- Oxford – Ilan Davis, Douglas Russell
- Cambridge – Rafael Carazo-Salas, Bálint Antal
- CRS4 - Gianluigi Zanetti, Gianmauro Cucurru, Simone Leo, Luca Lianas
- Edinburgh – Richard Baldock, Bil Hill, Jianguo Rao
- Carnegie-Mellon – Robert Murphy, BK Cho, Ivan Cao-Berg
- Imperial – Paul French, Chris Dunsby, Ian Munro, Yuriy Alexandrov
- NIA, NIH – Ilya Goldberg, Chris Coletta
- Pasteur – Spencer Shorte, Sebastien Simard, Julien Jorde
- EMBL-EBI – Gerard Kleywegt, Ardan Patwardhan, Ingvar Lagerstedt, Alvis Brazma
- Glencoe Software – Chris Allan, Joshua Ballanco, Andreas Knab, Melissa Linkert, Chris MacLeod, Josh Moore, Mike Rossner, Emil Rozbicki, Liza Unson, Rebecca Walker, Wilma Woudenberg

**THIS MEETING.....**

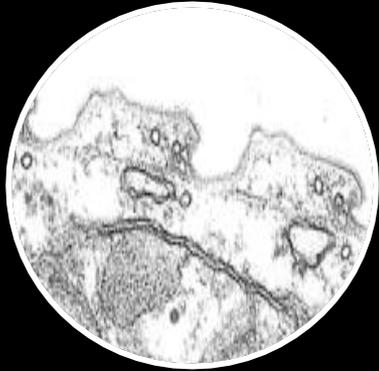
# Meeting Purpose

## 10<sup>th</sup> Annual User's Mtg

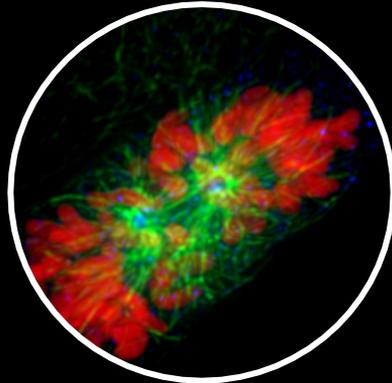
- Attendees
  - OME Consortium
  - Invited Speakers
  - Broad cross-section of users
- *Day 1: Presentations*
  - Lightning Talks
  - Project Overview
  - Users & Guests
- *Day 2: Workshops & Demos*
- Progress Report
- Future development priorities & planning

# THE PROBLEM

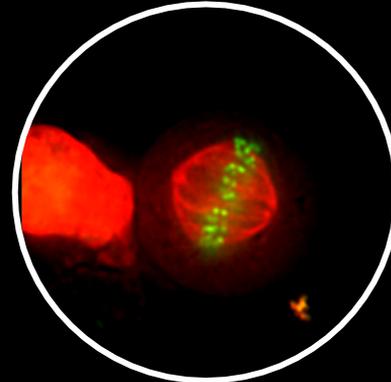
# The Image Problem... is Ubiquitous



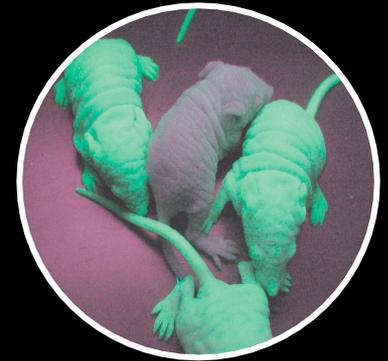
Organelles



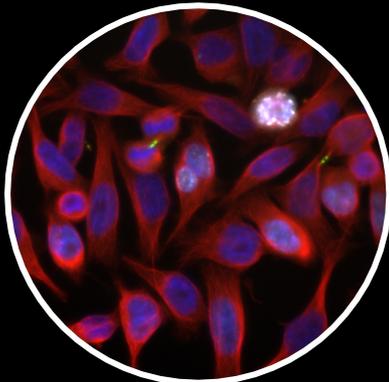
Cells



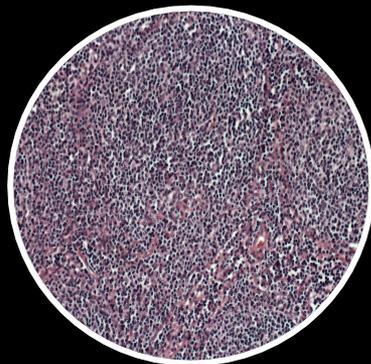
Dynamics



Physiology



Lead Discovery  
Target Validation



Pathology



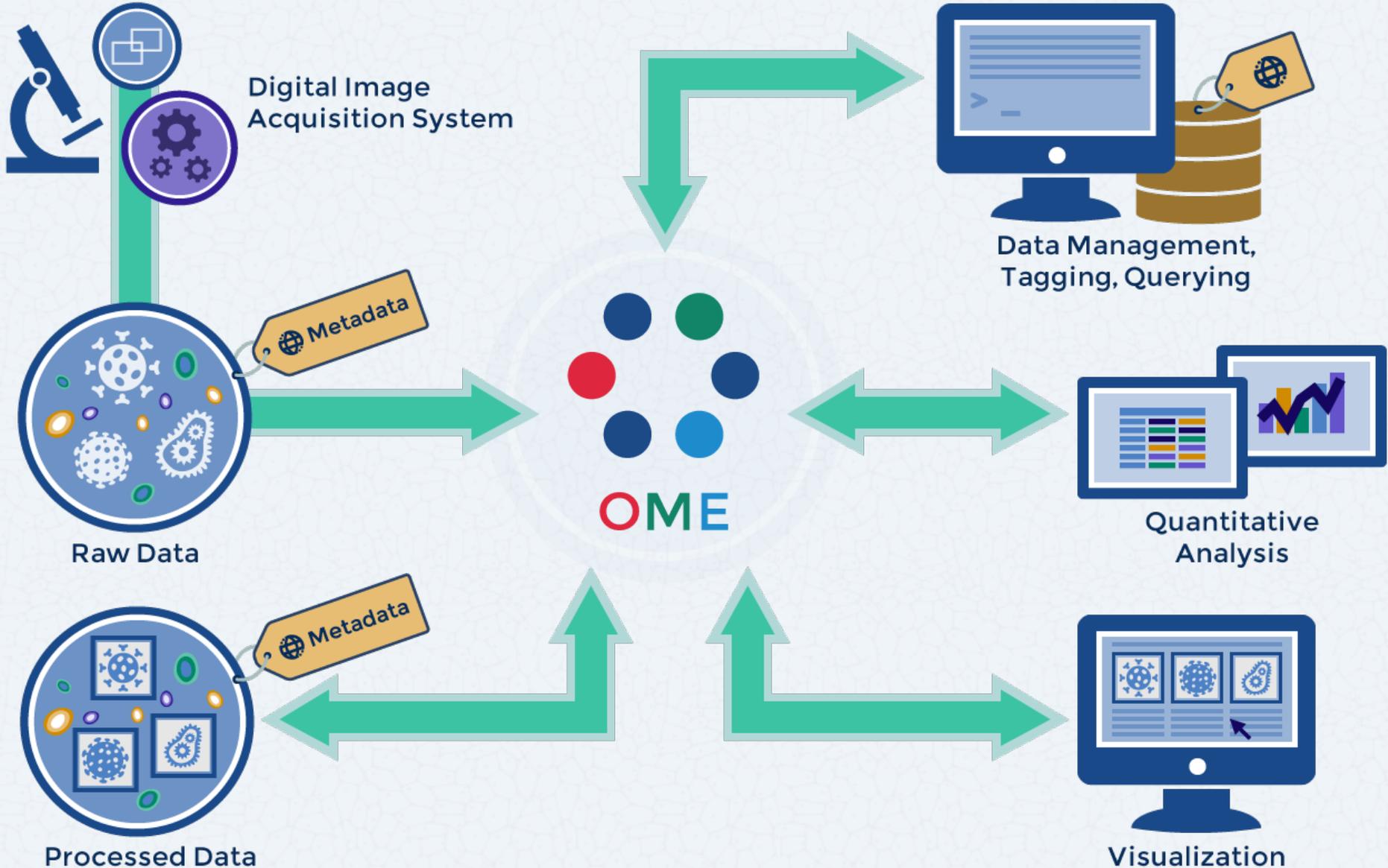
In Vivo

A pretty picture?

A measurement?

A resource?

# ...Towards Image Informatics



# OME : What We Do



**OME-XML**

**OME-TIFF**

**BIO-FORMATS**

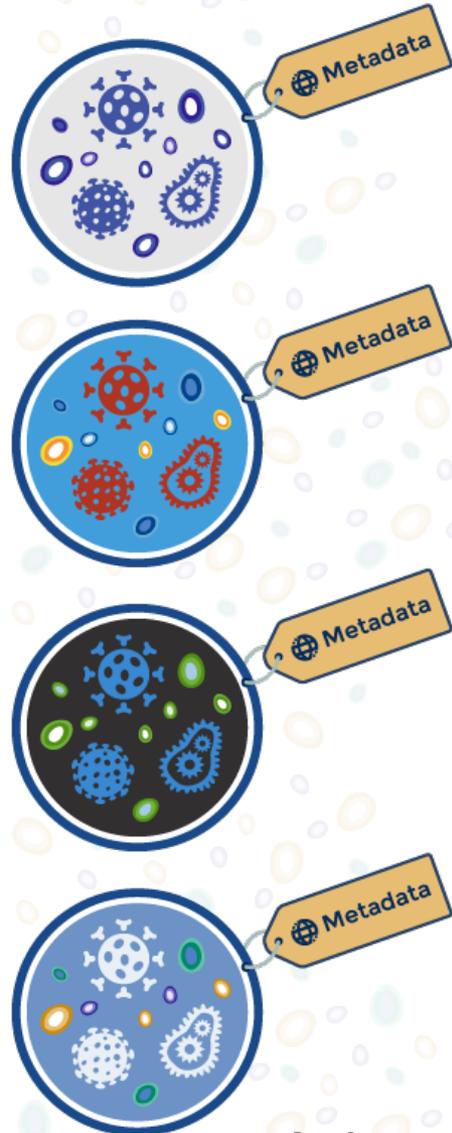
Open, exchangeable file formats



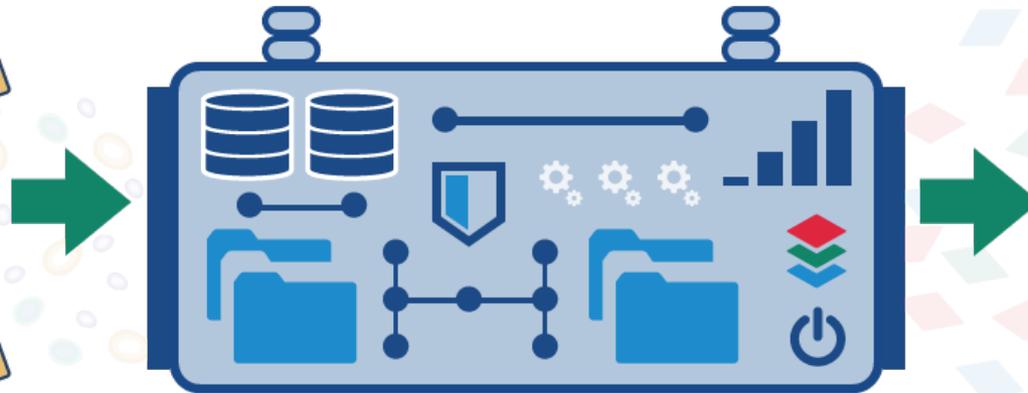
Open Image  
Management Software

# BIO-FORMATS: Proprietary File Conversion

Raw Data



## BIO-FORMATS



594,900+ FILES



28,600+ DATASETS



1.7 TERABYTES

Processed Data



→ Apps using Bio-Formats were started >1.5M times so far in 2015 ←



# BIO-FORMATS, OME-TIFF & $\mu$ -Manager: openFLIM-HCA



GitHub

This repository Search

Explore Features Enterprise Blog

Sign up

Sign in

imperial-photonics / openFLIM-HCA

Watch 4

Star 0

Fork 0

## Home

Fredxxx edited this page 13 days ago · 15 revisions

# openFLIM-HCA $\mu$ Manager Plugin

Welcome to the wiki for the openFLIM-HCA plugin for  [\$\mu\$ Manager](#). This is an open-source project that aims to allow experimenters to conduct their own HCA-FLIM studies.

This wiki is intended to be the main source of documentation both for **users** with access to an FLIM HCA platform, and for **developers** keen to implement their own FLIM HCA solutions.

- [Getting started](#)
- [Software reference](#)
- [Hardware reference](#)
- [Experiment guide](#)
- [About FLIM-HCA](#)
- [Links](#)

Pages 19

Find a Page...

Home

1 [Software reference](#)

1.1 [Getting started](#)

1.2 [The FLIM panel](#)

1.3 [The XYZ panel](#)

1.4 [The light path panel](#)

1.5 [The ProSetting Panel](#)

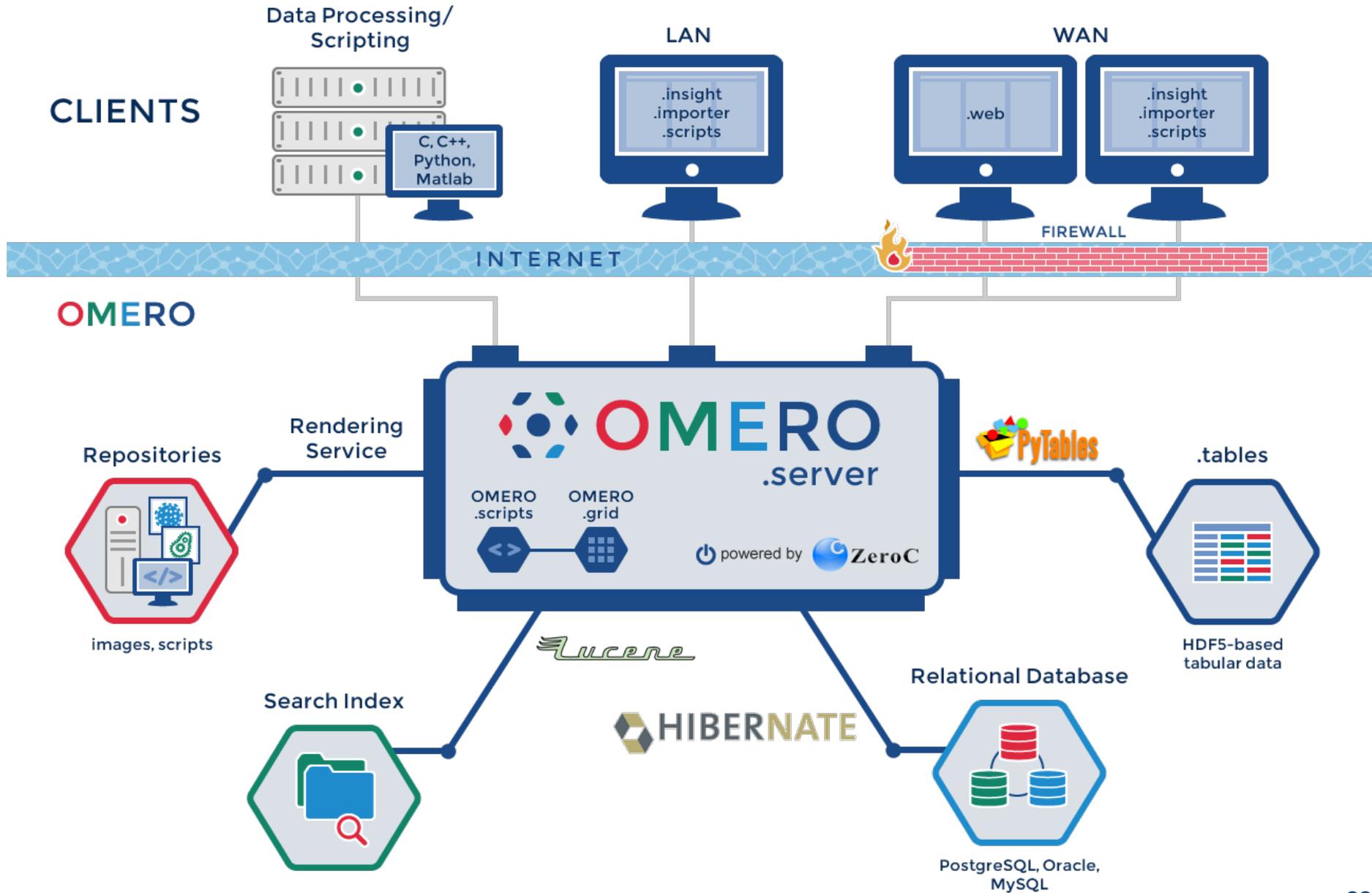
1.6 [The sequenced acquisition panel](#)

1.6.1 [XYZ positions setup for HCA sequence](#)

1.6.2 [Filter setup for HCA sequence](#)



# The OMERO Platform





# OMERO & BIO-FORMATS: OMERO.insight Java Client

The screenshot displays the OMERO.insight Java Client interface. The left sidebar shows a project tree for 'Jason Swedlow' with a sub-project 'Analysis MCAK - GFP-MCAK cell line [8]'. The main workspace shows a grid of 11 images. The right panel displays 'Image details' for image ID 3840635, including acquisition and annotation information. A 'Measurement Tool' window is open in the foreground, showing a table of ROI data.

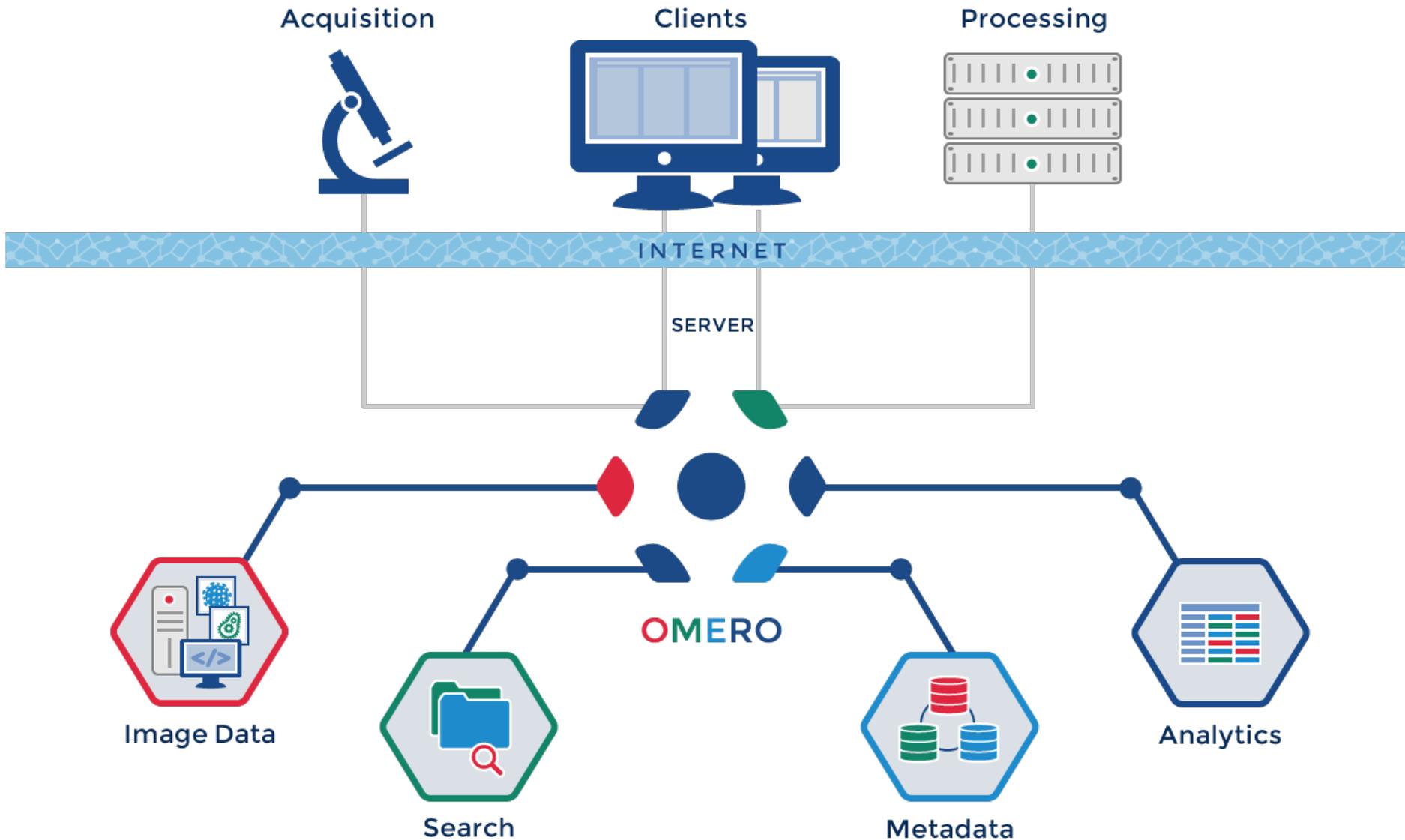
ROI	id	T	Z	Type	Text	Visible
	5662	[1,1]	[1,60]	<input type="checkbox"/>		<input checked="" type="checkbox"/>
	5663	[1,1]	[1,60]	<input type="checkbox"/>		<input checked="" type="checkbox"/>
	5664	[1,1]	[1,60]	<input type="checkbox"/>		<input checked="" type="checkbox"/>



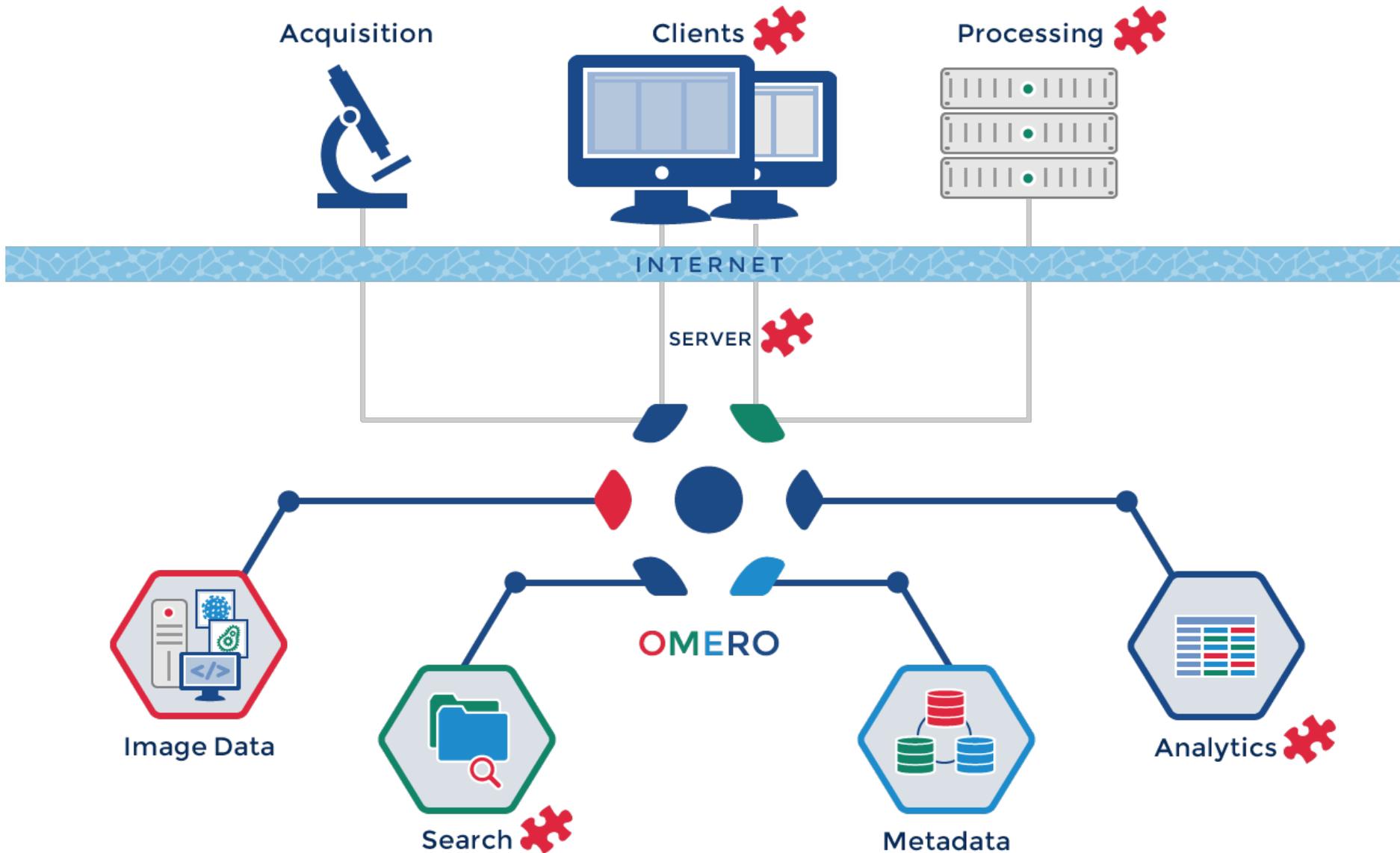
# OMERO & BIO-FORMATS: OMERO.web Client

The screenshot displays the OMERO.web Client interface. At the top, the OMERO logo and navigation tabs (Data, History) are visible. The user is identified as Jason Swedlow. The main area shows a list of images on the left, with the selected image being '101216-siHP1g\_LPM\_M\_total 11'. The central viewer displays a microscopy image with a 'Viewing Options' panel on the left, including controls for Max Intensity, Split Channel, Quality, Zoom (50%), and Line Plot. The 'Rendering Details' panel shows channels for DAPI (white), FITC (green), RD-TR-PE (red), and CY-5 (blue). The 'Current Image' panel shows Z: 32/60 | T: 1/1. The 'ROI Count: 3' panel is also visible. The right side of the interface shows the 'General' tab with metadata for the image: 'Z:/aferrand/HP1/101026-siRNA/101026-siCTL-GFPMCAK\_SKIP\_ACA\_02\_07\_R3D\_D3D.dv', IMAGE ID: 3840635, Owner: Alexia Ferrand, Acquisition Date: 2010-10-26 13:55:25, Imported Date: 2010-10-26 14:06:41, Dimensions (XY): 512 x 512, Pixels Type: uint16, Pixels Size (XYZ) (µm): 0.1001 x 0.1001 x 0.2000, Z-sections/Timepoints: 60 x 1, Channels: DAPI, FITC, RD-TR-PE, CY-5. A comment from Alexia Ferrand at 2010-11-09 14:16:33 is also visible.

# The OMERO Platform



# The *Extensible* OMERO Platform



 Plugins Welcome



# OMERO & BIO-FORMATS: Extensible Platforms

GitHub  [Explore](#) [Features](#) [Enterprise](#) [Blog](#) [Sign up](#) [Sign in](#)

[Contributions](#) [Repositories](#) [Public activity](#) [Follow](#)



**Daniel Matthews**  
drmatthews

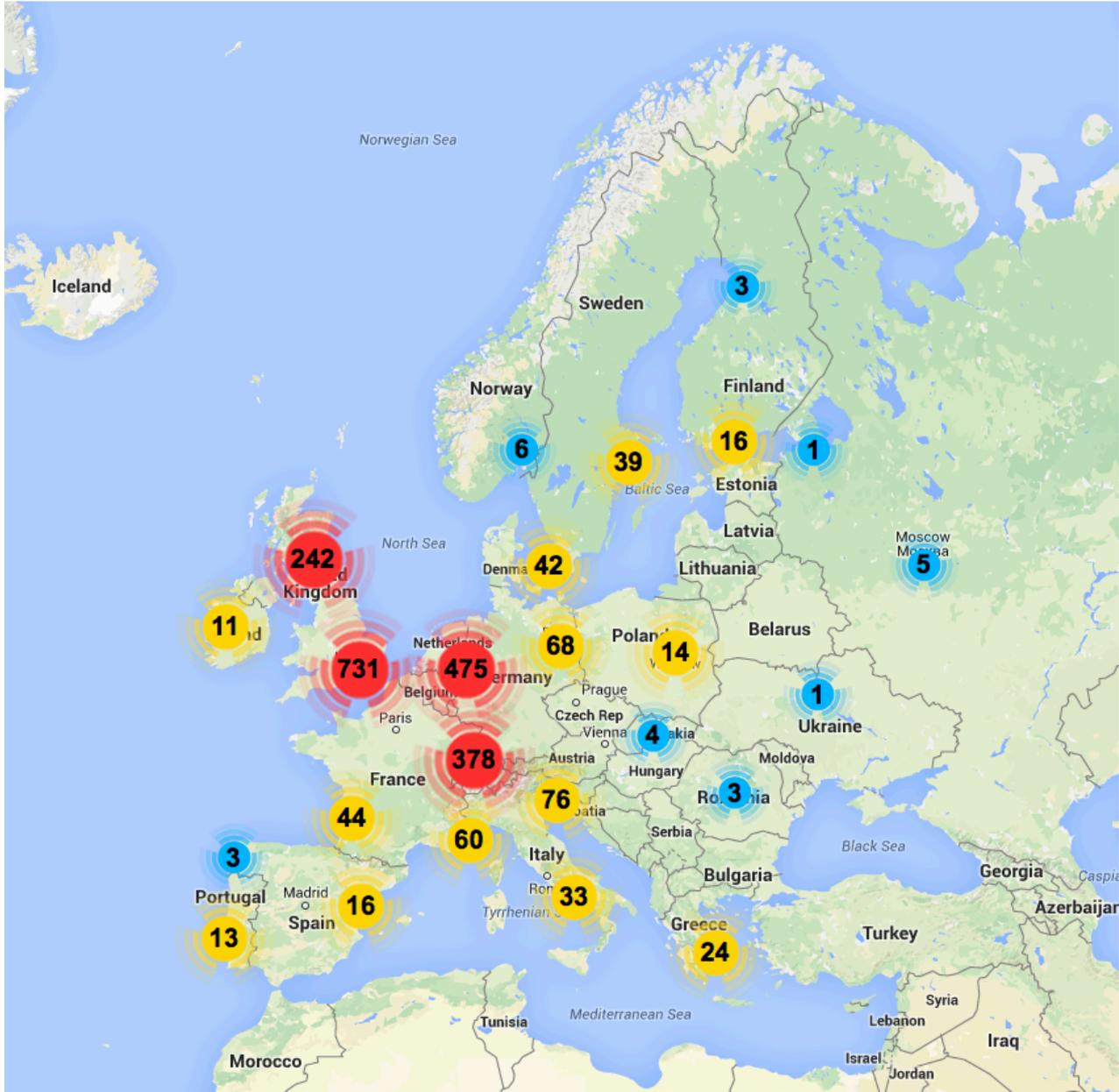
Joined on 18 Jun 2012

Popular repositories	
<a href="#">bioformats</a> Bio-Formats is a Java library for reading and ...	0 ★
<a href="#">SIMcheck</a> SIMcheck: ImageJ tools for assessing Structur...	0 ★
<a href="#">CellProfiler</a> CellProfiler is open-source cellular image anal...	0 ★
<a href="#">simple-STORM</a> Data processing software for dSTORM super-r...	0 ★
<a href="#">CUDA_SIMrecon</a> Image reconstruction for structured-illuminatio...	0 ★

Repositories contributed to	
<a href="#">QBI-Microscopy/SlideCrop</a> Python app, with wxPython GUI, for automate...	0 ★
<a href="#">QBI-Microscopy/omero-scripts-...</a> Utility functions and classes for omero-user-sc...	0 ★
<a href="#">QBI-Microscopy/Fiji-scripts</a> Jython scripts for Fiji-ImageJ	0 ★
<a href="#">QBI-Microscopy/Matlab-localisa...</a> a collection of scripts for parsing and analysin...	0 ★
<a href="#">QBI-Microscopy/omero-auto-upl...</a>	0 ★



# OMERO: EU Unique IPs: 2015YTD



# OUR PROGRESS

# OME : 2011-2016

- More data types (FLIM, LSFM, 3DEM, DigPath)
- Support complex, multi-dimensional, heterogeneous data (OME 5)
- Alternative image data storage methods (e.g., HDF5...)
- Validation of interfaces for analysis (esp. Matlab, Python, etc.)
- Integration of multi-parameter image-based search
- Data sharing & publication
- Shared Analysis Resources

# OMERO & BIO-FORMATS: Data Import & Access

Lab & Facility  
LM & EM



1-100 GB/day



HCS



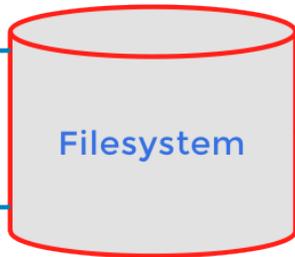
0.1 - 5 TB/day



Digital  
Pathology



0.01 - 1.2 TB/day



OMERO  
.server

Remote  
Clients  
(aware of Repo/.fs differences)

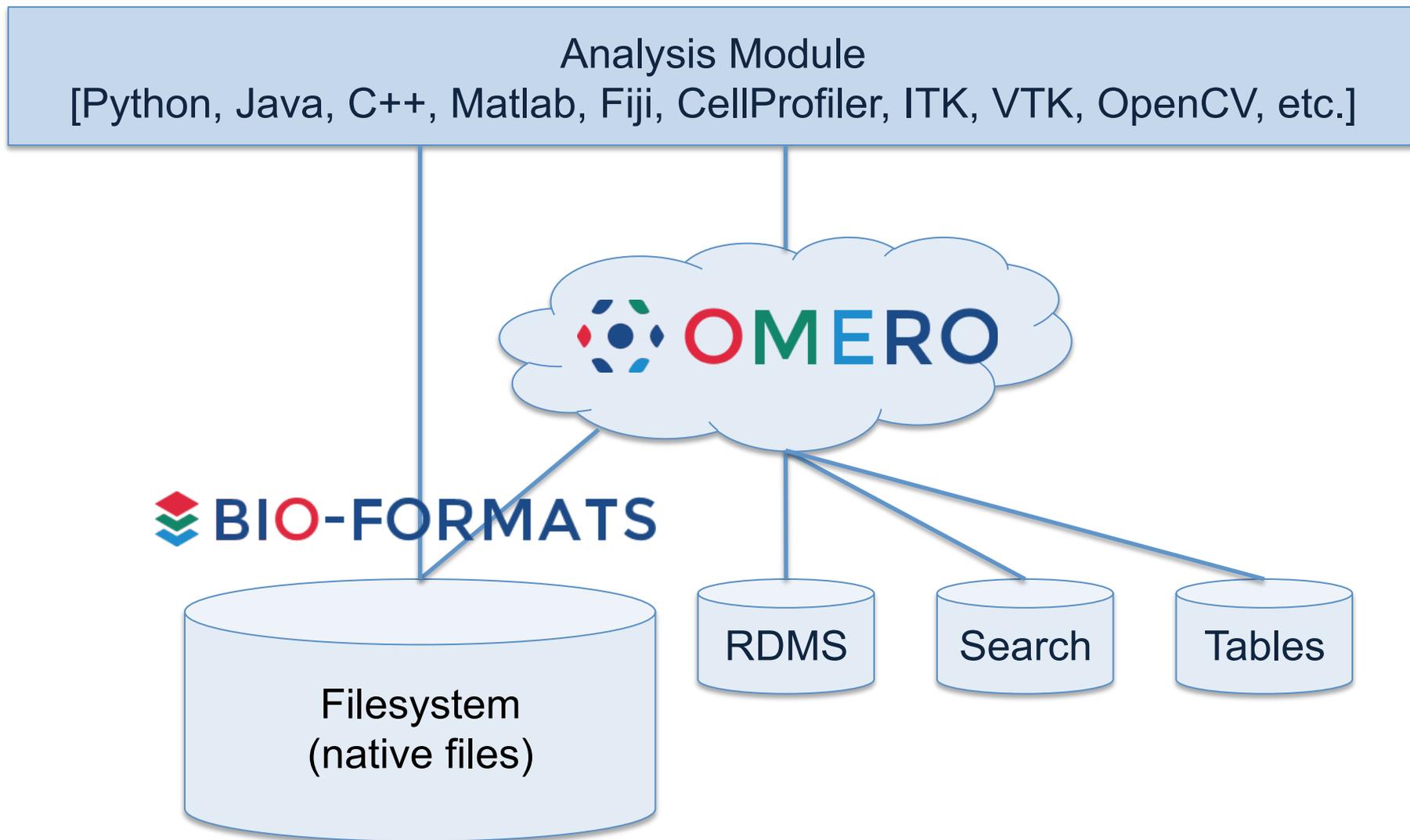
Rendering  
Service



images, scripts

OMERO-5.0 includes OMERO.fs, released Feb/2014 (Google: "OMERO 5.0")

# OMERO & BIO-FORMATS 5.0: Distributed Data Access



# OME 5.0 – 5.1: 2014/2015

## OME 5.1.x ...

- *Performance*
- *Bio-Formats/C++*
- *Improved support for “new” imaging modalities (SRM, LSFM, OPT, ...)*
- *API Updates*
  - Units
  - Map Annotations
  - New detectors
  - Rendering Settings
- *Extended Metadata support*
  - ROIs, Features, etc.
  - Graphs: (trajectories, provenance, ...)
- *Data Sharing & Publication*
  - R-W groups
  - Lab, Institute, National/Worldwide Repositories
- *Substantial UX and UI Updates*
  - Consistent UX
  - Configurable Web UI

# OME 5.1: Map Annotations

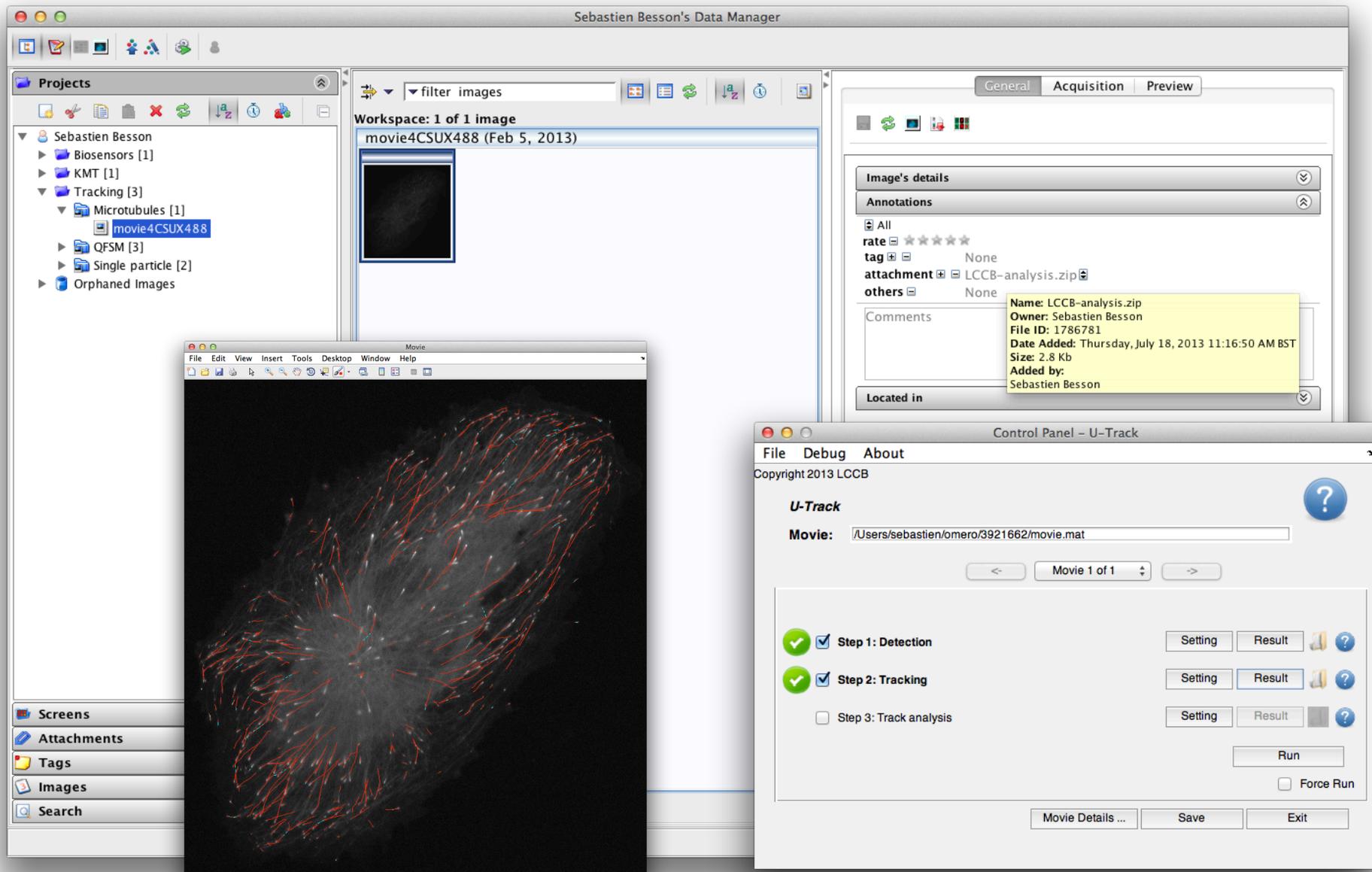
The screenshot displays the OME 5.1 interface for an image titled "1.6M P-TRE\_22\_R3D\_D3D\_VOL.dv". The "Annotations" section is highlighted, showing a table of key-value pairs. A blue arrow points to the table, with the text "Ordered, non-unique, key-value pairs" next to it.

Annotations:

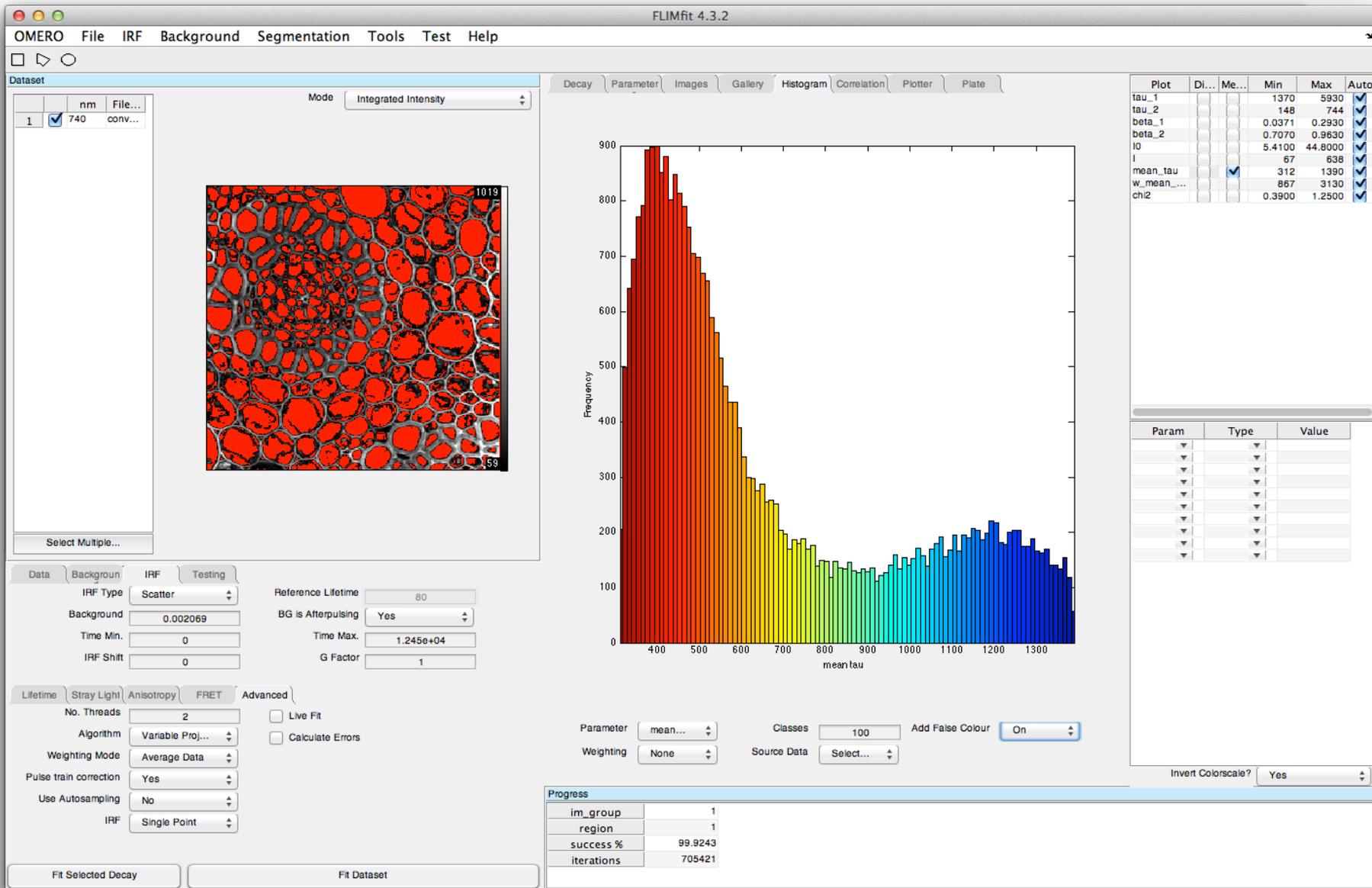
Key	Value	
Temperature	37.0	x
Cell Line	HT-29	x
Concentration - Sodium	150.0	x
Concentration - Potassium	4.3	x
Add key	Add value	
Concentration - Potassium	4.3	
Concentration - Potassium	4.8	
Antibodies	Sheep	

Ordered, non-unique, key-value pairs

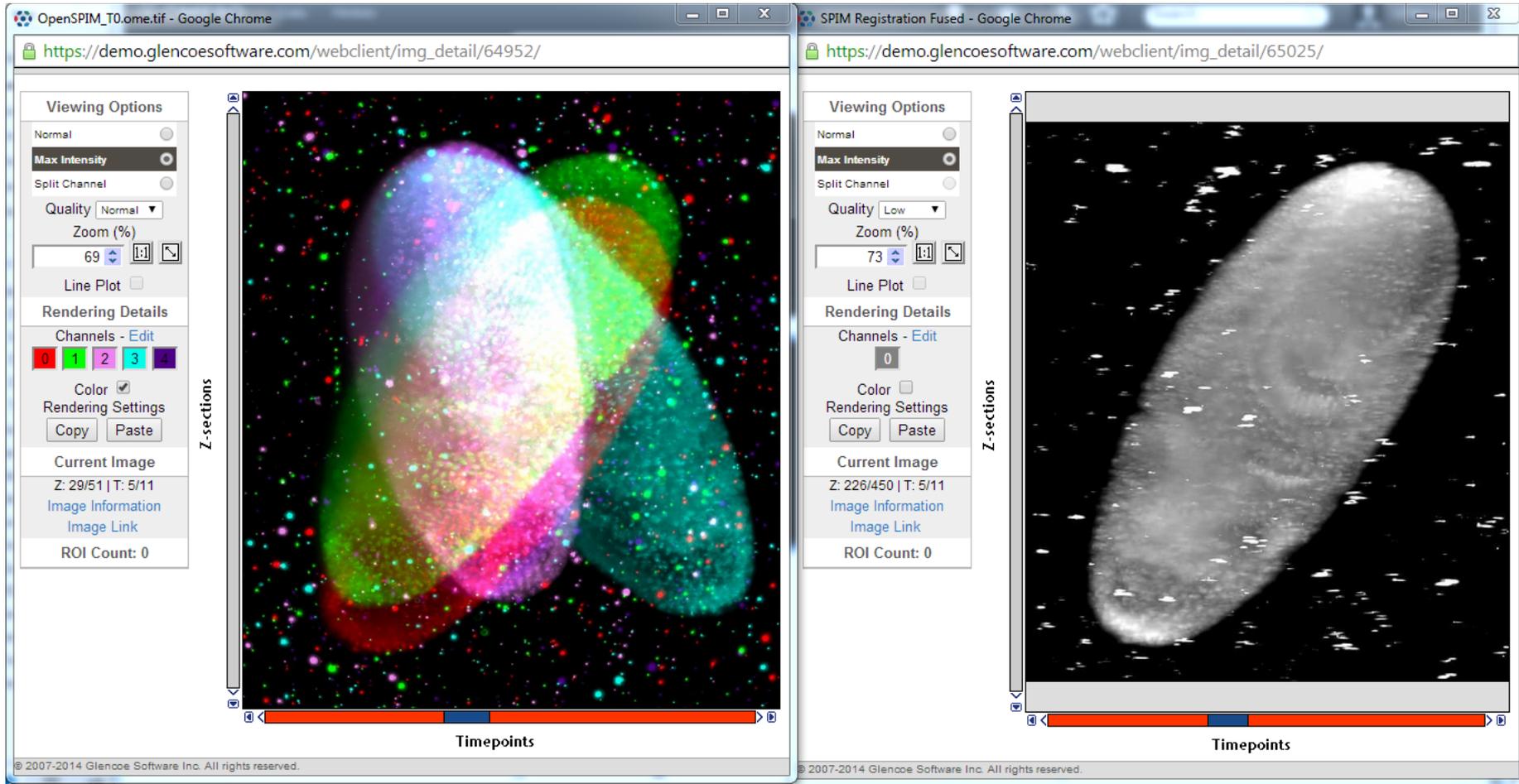
# OMERO & u-track



# OMERO & FLIMfit



# OMERO & Bio-Formats: LSFM MV Reconstruction

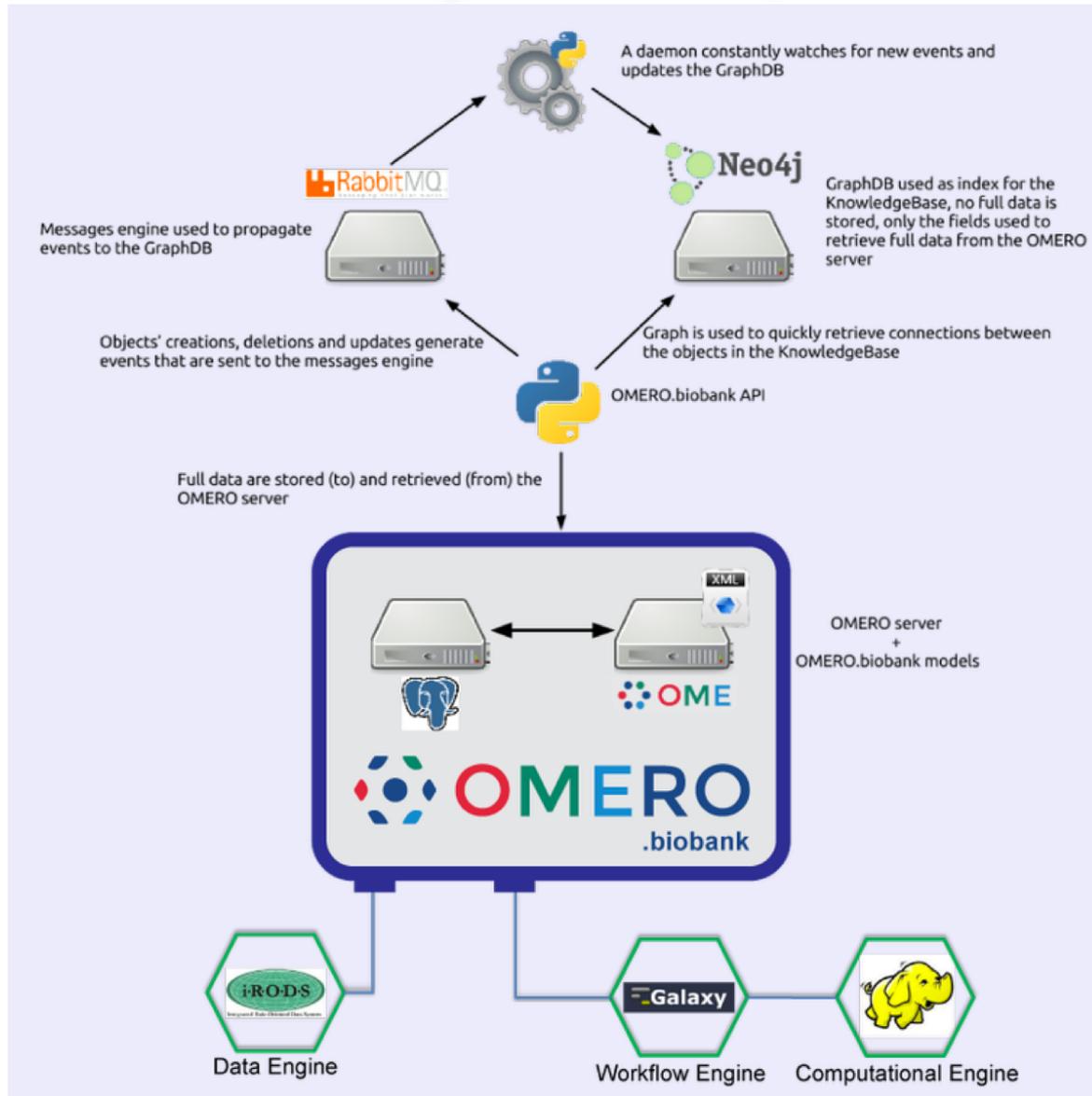


Emil Rozbicki & Chris Allan, Glencoe Software

Inspired by Preibisch et al. (2010) Nature Meth, 7: 418-419; [http://fiji.sc/SPIM\\_Registration](http://fiji.sc/SPIM_Registration) 



# OMERO.biobank: Enabling Meta-Compute



# OMERO & Auto-Tagging

The screenshot shows the OMERO webclient interface. The browser address bar displays `https://omero1.bioch.ox.ac.uk/webclient/userdata/`. The interface includes a navigation menu with 'Data', 'History', 'Admin', and 'Tag Search'. The user is logged in as 'Douglas Russell'. The main area shows a file tree on the left with folders like 'Demo 3', 'CSFV 12', 'INCEP 6', 'scrambled 6', 'P1 1', 'TestCopy1 1', 'TestLothar 1', and 'Orphaned images'. The central table displays a list of files with columns for tags: 'siRNAi', 'home', 'example', 'dpwrussell', 'data', 'INCEP', 'FromWill', 'Downloads', 'CSFV', 'r3d', and 'D3D'. Each cell contains a dropdown menu and a checkbox. The 'siRNAi' column is highlighted in yellow. The right sidebar shows the 'General' tab for the selected file, displaying 'DATASET ID: 4201', 'Owner: Douglas Russell', and 'Creation Date: 2014-04-25 16:02:30'. There are also sections for 'RATING', 'TAGS', 'ATTACHMENTS', and 'COMMENT'.

File Name	siRNAi	home	example	dpwrussell	data	INCEP	FromWill	Downloads	CSFV	r3d	D3D
...siRNAi/CSFV/CSFV_01.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_02.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_03.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_04.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_05.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_06.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_07.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_08.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_09.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_10.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_11.r3d_D3D.dv	<input checked="" type="checkbox"/>										
...siRNAi/CSFV/CSFV_12.r3d_D3D.dv	<input checked="" type="checkbox"/>										
INCEP 6	<input checked="" type="checkbox"/>										
scrambled 6	<input checked="" type="checkbox"/>										
P1 1	<input checked="" type="checkbox"/>										
TestCopy1 1	<input checked="" type="checkbox"/>										
TestLothar 1	<input checked="" type="checkbox"/>										
Orphaned images	<input checked="" type="checkbox"/>										

# OMERO.figure

## OMERO.figure

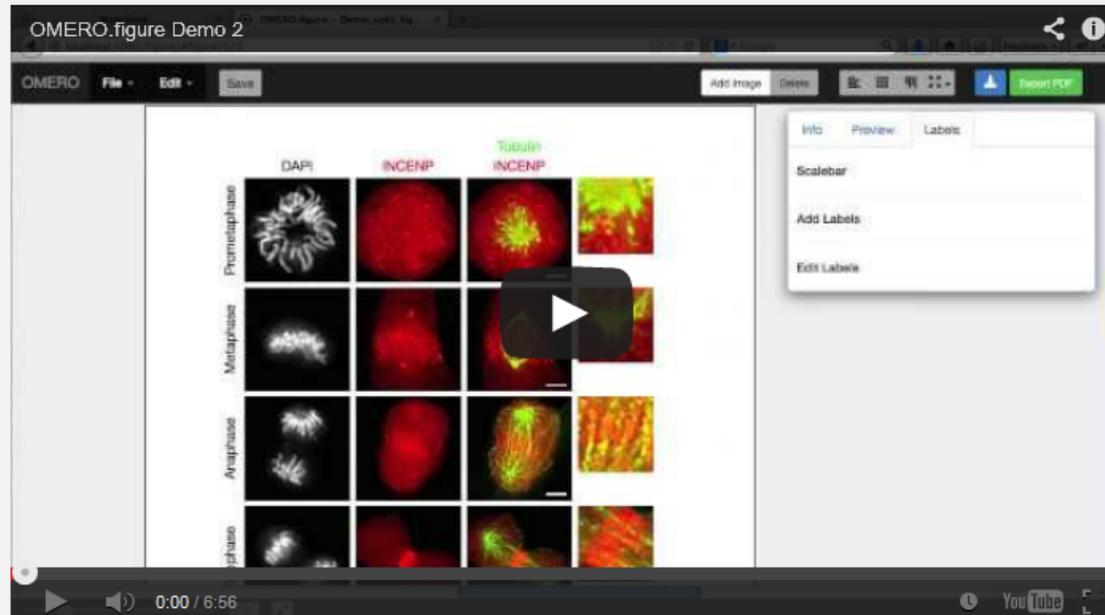
Try the DEMO (beta!)

[Home](#) | [Blog](#) | [User Manual](#) |

tar.gz

.zip

### Fast figures from your OMERO images



### Smart figures with metadata

OMERO is a server platform for managing biological images. See the [Open Microscopy Site](#) for more info. OMERO.figure combines OMERO's powerful image rendering and metadata to provide a tool for rapid figure creation. Each panel of the figure becomes a multi-dimensional image viewer allowing you to zoom and pan adjust rendering



# OMERO & BIO-FORMATS: Pathology Instruction/Dundee VM

The screenshot displays the OMERO web client interface. At the top, there is a navigation bar with 'Data' and 'History' menus, a search bar, and a user profile for 'Jason Swedlow'. Below this is a 'Virtual Microscope Anatomy' header. The left sidebar shows a tree view of the file system under 'Anatomy', with '14\_Tendon.svs' selected. The main area features a 'Filter Images' search bar and a grid of image thumbnails. The right sidebar provides metadata for the selected image, '14\_Tendon.svs' (IMAGE ID: 46), including acquisition and import dates, dimensions, and channel information. A detailed view window is open over the selected image, showing 'Viewing Options' (Quality: Normal), 'Rendering Details' (Channels: 0, 1, 2; Color: checked), and 'Current Image' (Z: 1/1 | T: 1/1). The detailed view includes a 'Z-sections' slider and a 'Timepoints' slider, with a 'Scale: 25.00%' indicator.

Property	Value
Owner:	Anatomy
Acquisition Date:	2011-08-26 20:08:22
Imported Date:	2013-07-30 14:23:16
Dimensions (XY):	42257 x 43037
Pixels Type:	uint8
Pixels Size (XYZ) (µm):	0.0000 x 0.0000
Z-sections/Timepoints:	1 x 1
Channels:	0, 1, 2

# OMERO Help

## Getting Started with OMERO.insight v. 5.0.2

[Download PDF](#)



### User Help

[User Help Home Page](#)

#### ▼ Quickstart User Guides

[Getting Started v. 5.0.2](#)

[Getting Started v. 4.4.11](#)

[Using ImageJ with OMERO](#)

[Try the OMERO Demo Server](#)

#### ▼ Workflow User Guides

[Sharing Data](#)

#### ▶ OMERO.insight

#### ▶ OMERO.web

#### ▼ Other OMERO Applications

[OMERO.figure](#)

[Virtual Microscope](#)

[OMERO.dropbox](#)

[OMERO.editor](#)

#### ▼ More

[Guides for Previous Versions](#)

[Resources](#)

[Contact Us](#)

[Main OME Website](#)

© 2000-2014

The Open Microscopy Environment

OMERO stores image data on a central server. You can use the OMERO.insight client to upload, view and download data from any personal computer.

Your institution's OMERO server administrator will provide you with the server address to use when connecting from OMERO.insight. They may have set up a user name and password for you too. Alternatively your username and password may be your standard login for the institution's networked computers.

### Installing

1. Download OMERO.insight client at: <http://downloads.openmicroscopy.org/latest/omero5>



### OMERO 5.0.2 Downloads

[Clients](#) | [Plugins](#) | [Additional](#) | [Servers](#) | [Virtual Appliance](#) | [API](#) | [Code](#) | [Artifacts](#) | [Legacy](#)

- Information on this release of OMERO is in the [release announcement](#)
- Full documentation is available as [web documentation](#) or [PDF documentation](#) and there are user guides for the clients on our [Help website](#)
- A standard OMERO user just needs to download the client package with the same major version as their institutional server e.g. 5.0 clients with the 5.0 server

### OMERO client downloads

Clients	Size	File Name	Checksum
Windows	70.29 MB	OMERO.clients-5.0.2-icc35-b21.win.zip	3fb156 ( <a href="#">MD5</a> )
Mac OS X	205.68 MB	OMERO.clients-5.0.2-icc35-b21.mac.zip	55f177 ( <a href="#">MD5</a> )
Linux	68.44 MB	OMERO.clients-5.0.2-icc35-b21.linux.zip	a6e1a6 ( <a href="#">MD5</a> )

- Each client package includes [OMERO.insight](#), [OMERO.importer](#) and [OMERO.editor](#) and requires Java Version 1.6 or higher. OMERO.web is part of the server package, so individual users do not need to install it locally.



# OME-based Public Data Resources

- Public Data Resource
  - Harvard LINCS
  - JCB DataViewer
  - Stowers ODR
  - SSBD, Riken
  - EMDataBank– 3D tomograms
  - SYSGRO– *S. pombe* phenotypic screens
  - Liverpool CCI
  - Imperial CISBIC FLIM
  - Dundee Virtual Microscope
  - .....

# 2015 Highlights

- Bio-Formats and OMERO 5.1
  - Model & API Updates
  - Performance
  - Extended Metadata
  - UI Improvements
  - Commercial contributions and participation
    - Specifications, suggestions from several industrial partners
    - Intelligent Imaging, Richard Myers, SlideBook Reader
- Bio-Formats/C++
  - Native OME-TIFF Reader & Writer
- Major releases of OMERO.figure and FLIMfit
- Security: SecVuln Patches and Process
- New modalities
  - LSM: Multiview reconstruction
  - OPT: “Spinny Fish”
- Learning & Teaching
- Growing, Active Community
  - Feedback on Forums
  - Social Media: 2x jump in Twitter followers

# “Super OMERO”



# THE FUNDING

££££££££

- Extension of Strategic Award, co-funded by Wellcome Trust & BBSRC
  - ~£600k
  - Funds Bio-Formats and OMERO through mid-2016
- BBSRC “Big Data” Award
  - £1.79M partnership between Dundee OME, Cambridge & EBI
    - Partnership with Elixir
  - Build & deploy a next generation image data repo
    - >50 TB of GW HCS datasets at start, growing to >100 TB
  - Includes virtual analysis resource
  - More info: “Euro-BioImaging Elixir Data”
- H2020
  - CORBEL, MULTIMOT, (INFRADEV-2)

# 2015/2016 PRIORITIES?

# Some words to think about...

- New Modalities (MS, Raman, X-ray, etc.)
- Multi-modal/Correlative
- Federation:
  - SSO
  - Multiple server Auth
- Import/Export
- Ontologies
- Bio-Formats Decoupling
- OMERO Gateway
- REST API
- C++ API
- Client Architecture
- Archiving
- ...

# OME Consortium

- Dundee – Jason Swedlow, Colin Blackburn, Jean-Marie Burel, Mark Carroll, Gus Ferguson, Helen Flynn, Kenny Gillen, Roger Leigh, Simon Li, Dominik Lindner, Josh Moore, Will Moore, Balaji Ramalingam, Gabriella Rustici, Aleksandra Tarkowska, Petr Walczysko, Eleanor Williams *and you?!?*
- University of Wisconsin, Madison (LOCI) - Kevin Eliceiri, Curtis Rueden, Mark Hiner
- UT Southwestern – Gaudenz Danuser, Sebastian Besson
- Oxford – Ilan Davis, Douglas Russell
- Cambridge – Rafael Carazo-Salas, Bálint Antal
- CRS4 - Gianluigi Zanetti, Gianmauro Cucurru, Simone Leo, Luca Lianas
- Edinburgh – Richard Baldock, Bil Hill, Jianguo Rao
- Carnegie-Mellon – Robert Murphy, BK Cho, Ivan Cao-Berg
- Imperial – Paul French, Chris Dunsby, Ian Munro, Yuriy Alexandrov
- NIA, NIH – Ilya Goldberg, Chris Coletta
- Pasteur – Spencer Shorte, Sebastien Simard, Julien Jorde
- EMBL-EBI – Gerard Kleywegt, Ardan Patwardhan, Ingvar Lagerstedt, Alvis Brazma
- Glencoe Software – Chris Allan, Joshua Ballanco, Andreas Knab, Melissa Linkert, Chris MacLeod, Josh Moore, Mike Rossner, Emil Rozbicki, Liza Unson, Rebecca Walker, Wilma Woudenberg