



# Mutimage UCA : Facilitating the management of life sciences imaging data



S.Ben Aïcha - F. Bekkouche - R. Arkowitz - F. Brau





# Project Goals

Facilitate the management, sharing and analysis of biological images on the life sciences community

Organize, coordinate, structure and share resources and expertise to promote and stimulate quantitative image analysis in biology



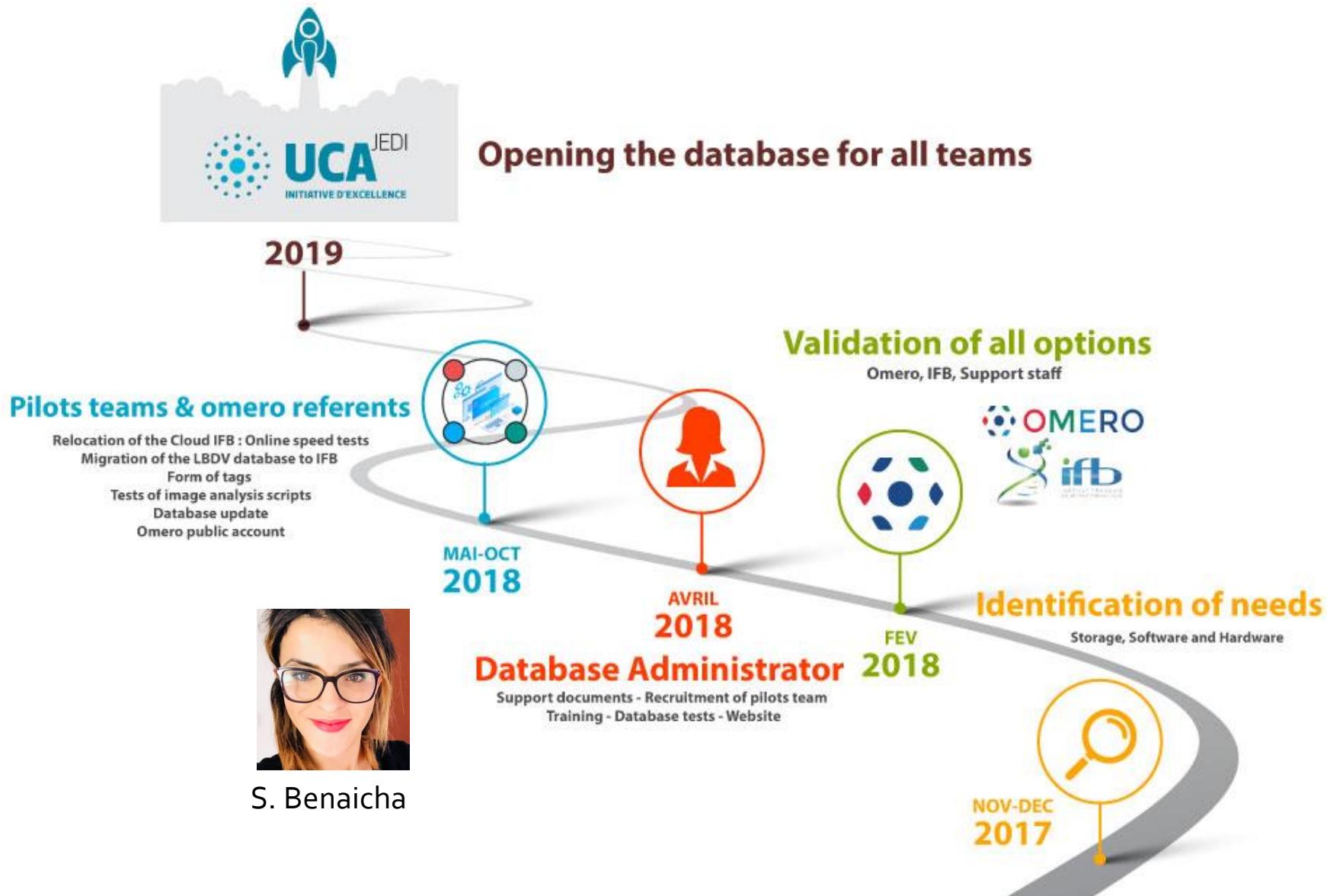
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CÔTE D'AZUR

# The real goal



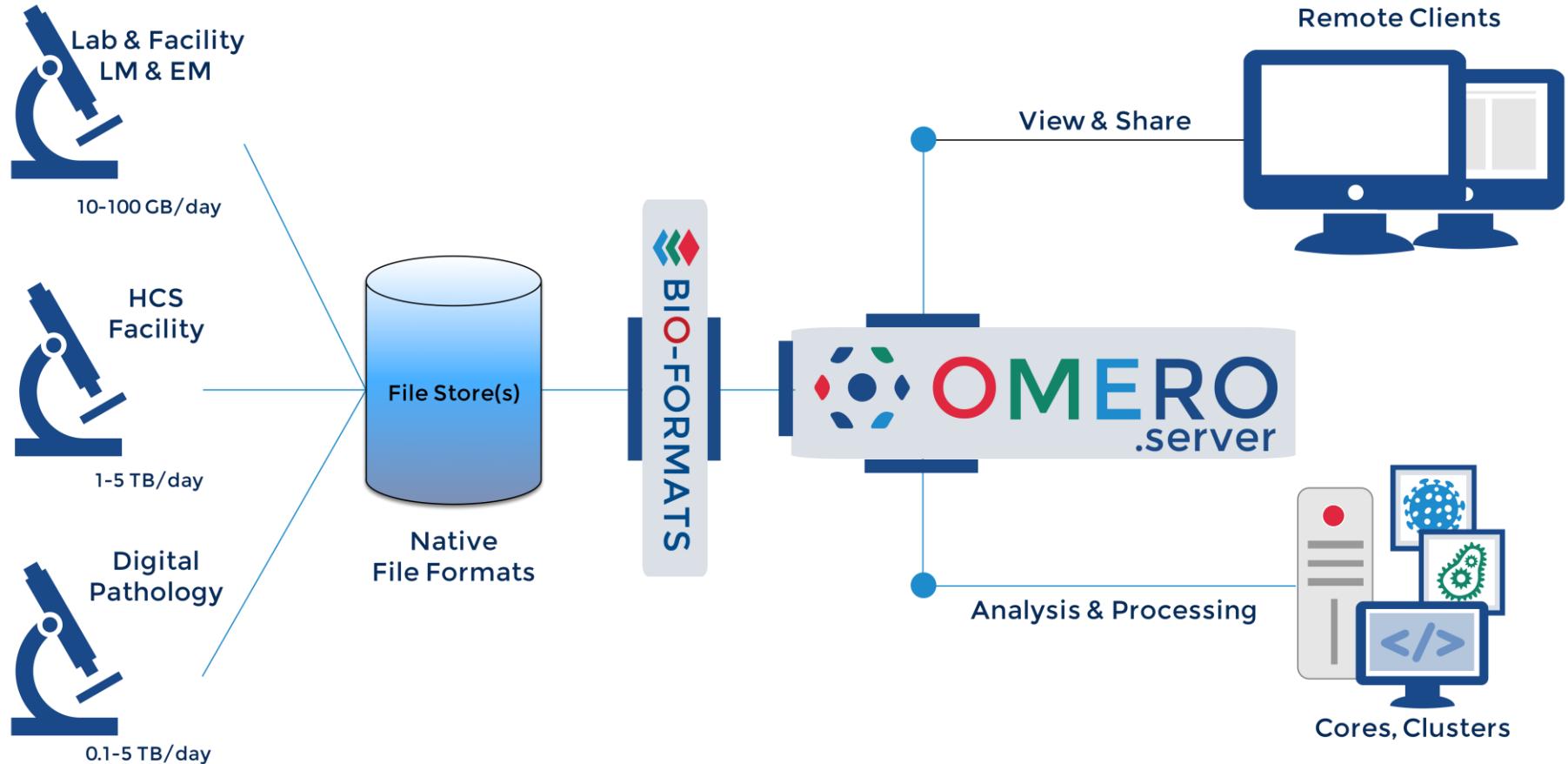


# Building the project





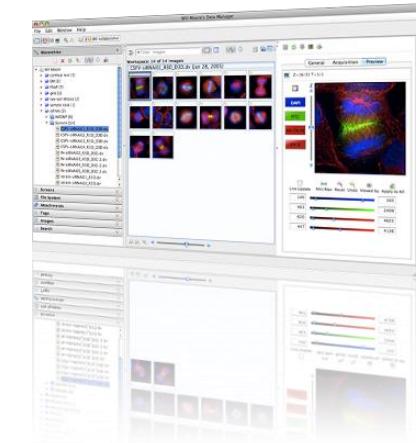
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<https://openmicroscopy.org/omero>



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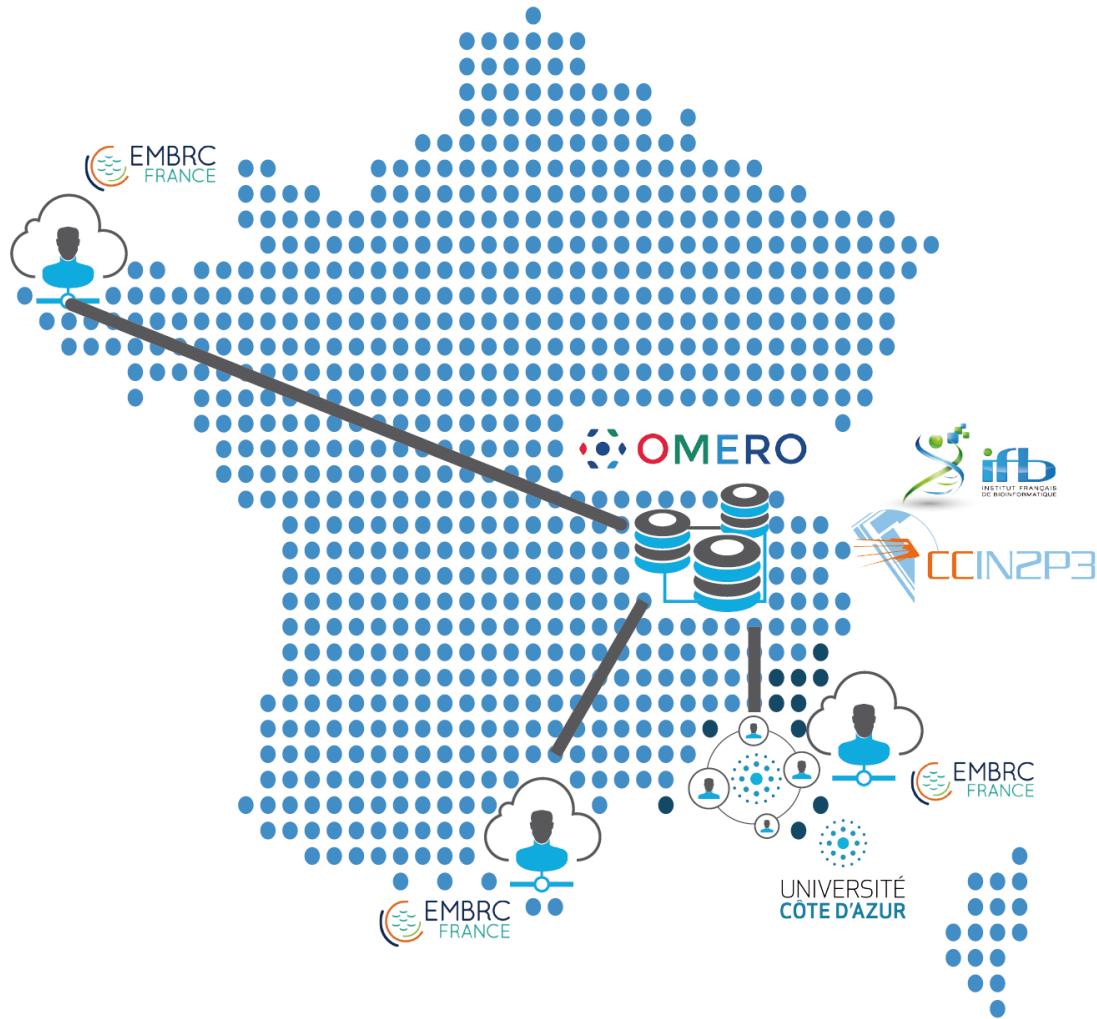


**UCA** JEDI  
INITIATIVE D'EXCELLENCE

# Mutualisation UCA EMBRC-Fr



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EMBRC  
FRANCE

UNIVERSITÉ  
CÔTE D'AZUR





# Functional organization

C. Blanchet  
Data base – Storage – Data Security



F. Bekkouche  
*Technical referent Omero*



S. Pisano  
IRCAN

M. Irondelle  
C3M

O. Pierre  
ISA

S.Lachambre  
iBV

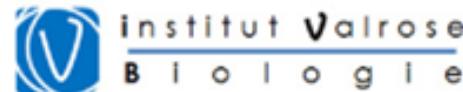
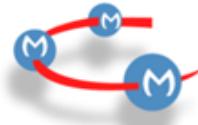
G. Carles  
TIRO-MATOS

F. Brau/S. Abelanet  
IPMC

Sameh.Ben Aicha  
LBDV

E. Debreuve  
Morpheme

Manage groups and users access  
Training & Support





# User accounts management

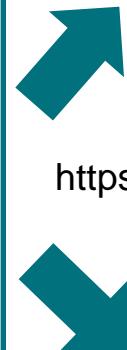


Team 1

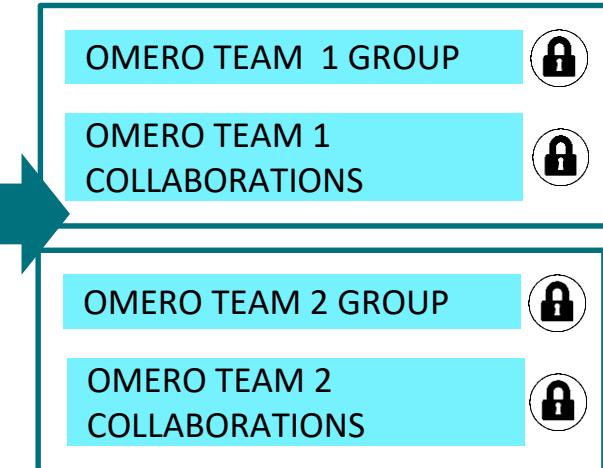
Team 2

Team 3

...



The OMERO.web login page is shown. It has a dropdown menu set to 'omero.4064'. Below it are fields for 'Username:' and 'Password:', both currently empty. A 'Forgot your password?' link and a 'Login' button are at the bottom.



The OMERO.web PUBLIC login page is shown. It has a dropdown menu set to 'omero.4064'. Below it are fields for 'Username:' and 'Password:', both currently empty. A 'Forgot your password?' link and a 'Login' button are at the bottom. The word 'PUBLIC' is prominently displayed in red capital letters above the login fields.



<https://omero.france-bioinformatique.fr/omero/webclient/login/>

# Bio-Formats: support for >150 file formats



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BIO-RAD

OXFORD  
INSTRUMENTS ANDOR

HAMAMATSU  
PHOTON IS OUR BUSINESS

PerkinElmer®

Leica  
MICROSYSTEMS

Nikon

GE

ZEISS

OLYMPUS®

BD  
BD Biosciences

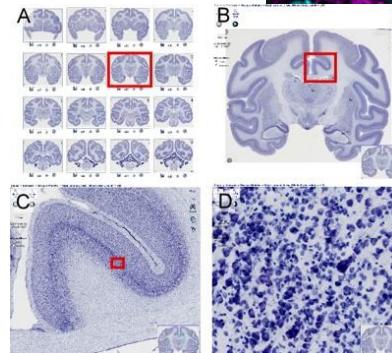
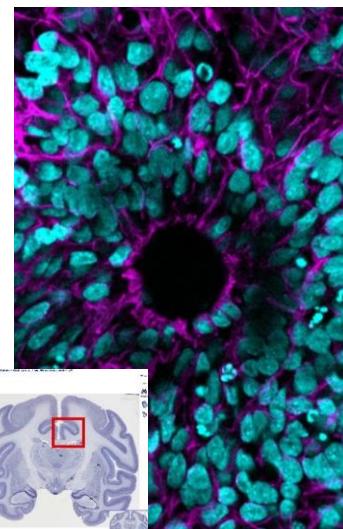
MetaMorph®  
BRUKER

IMARIS 8  
BITPLANE

ThermoFisher  
SCIENTIFIC

OME

ImageJ  
Fiji  
CellProfiler  
OMERO  
MATLAB  
Octave  
R  
Orbit  
...



## Image pyramids

<https://www.openmicroscopy.org/bio-formats>



# Image management

BIO-FORMATS

**Tags**

OMERO Data History

Polly Stack

Disease 3

- 201102142-Fig 1 2
- 201102142-Fig 4 6
- 201102142-Fig 6 8
  - ...eated C3H hepatocytes.ome.tif
  - ...eated C3H hepatocytes.ome.tif
  - ...ted C57BL hepatocytes.ome.tif
  - ...ted C57BL hepatocytes.ome.tif
  - ...eated C3H hepatocytes.ome.tif
  - ...eated C3H hepatocytes.ome.tif
  - ...ted C57BL hepatocytes.ome.tif
  - ...ted C57BL hepatocytes.ome.tif

Structural Analysis 1

Collaboration 3

- ...nd kinetochores (red).ome.tif
- ...nd kinetochores (red).ome.tif
- ...nd kinetochores (red).ome.tif

Orphaned images

Thumbnails

General Acquisition

31918-GAPDH staining of DDC-treated C3H hepatocytes.ome.tif

IMAGE ID: 110

Launch full viewer

GAPDH staining of DDC-treated C3H hepatocytes

Owner: Polly Stack  
Acquisition Date: 2012-06-29 10:57:54  
Imported Date: 2012-06-29 10:52:30  
Dimensions (XY): 1024 x 1024  
Pixels Type: uint8  
Z-sections/Timepoints: 1 x 1  
Channels: 0, 1, 2

**Tags**

RATING  
No ratings

TAGS

ATTACHMENTS

OTHERS:



# Image management

BIO-FORMATS

**Search engine**

**Tags**

**Metadata**

**Tags**

**Attach files**

OMERO Data History

Tags

Polly Stack

Disease 3

- 201102142-Fig 1 2
- 201102142-Fig 4 6
- 201102142-Fig 6 8
  - ...eated C3H hepatocytes.ome.tif
  - ...eated C3H hepatocytes.ome.tif
  - ...ted C57BL hepatocytes.ome.tif
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  - ...eated C3H hepatocytes.ome.tif
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Structural Analysis 1

Collaboration 3

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Orphaned images

Thumbnails

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Pixels Type: uint8  
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RATING  
No ratings

TAGS

ATTACHMENTS

OTHERS:



# Image management

BIO-FORMATS

**Search engine**

**Scripts**

**Tags**

**Metadata**

**Tags**

**Attach files**

**OMERO Data History**

**Explore**

**Polly Stack**

- Disease 3
  - 201102142-Fig 1 2
  - 201102142-Fig 4 6
  - 201102142-Fig 6 8
    - ...eated C3H hepatocytes.ome.tif
    - ...eated C3H hepatocytes.ome.tif
    - ...ted C57BL hepatocytes.ome.tif
    - ...ted C57BL hepatocytes.ome.tif
    - ...eated C3H hepatocytes.ome.tif
    - ...eated C3H hepatocytes.ome.tif
    - ...ted C57BL hepatocytes.ome.tif
    - ...ted C57BL hepatocytes.ome.tif
- Structural Analysis 1
  - Collaboration 3
    - ...nd kinetochores (red).ome.tif
    - ...nd kinetochores (red).ome.tif
    - ...nd kinetochores (red).ome.tif
- Orphaned images

Thumbnails

General Acquisition

IMAGE ID: 110

Launch full viewer

GAPDH staining of DDC-treated C3H hepatocytes

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Imported Date: 2012-06-29 10:52:30  
Dimensions (XY): 1024 x 1024  
Pixels Type: uint8  
Z-sections/Timepoints: 1 x 1  
Channels: 0, 1, 2

RATING  
No ratings

TAGS

ATTACHMENTS

OTHERS:

The screenshot displays the Bio-Formats software interface for image management. At the top, there's a navigation bar with tabs for OMERO, Data, and History. Below this is a toolbar with various icons for file operations like New, Open, Save, and Delete. A large blue box labeled 'Tags' is overlaid on the left side of the interface, covering the file tree and some of the image thumbnails. In the center, there's a grid of image thumbnails showing fluorescence microscopy images of cells. To the right of the thumbnails is a detailed metadata panel for a specific image (ID 110), which includes fields for owner (Polly Stack), acquisition date (2012-06-29 10:57:54), imported date (2012-06-29 10:52:30), dimensions (1024 x 1024), pixel type (uint8), z-sections/timepoints (1 x 1), and channels (0, 1, 2). Another blue box labeled 'Tags' is overlaid on the bottom right of the metadata panel. At the very bottom, there are additional blue boxes labeled 'Scripts', 'Search engine', 'Metadata', 'Attach files', and 'Others'.



# OMERO iViewer

Open multi-selected images

OMERO File ▾ ROIs ▾ Help ▾ scram\_06.r3d

Info Settings ROIs [3]

Save Undo Redo  Show Comments

Comment 10

Z  T 1

Show Z T C Comment

5.04 μm

Multi-D Scroll Z projection



# OMERO iViewer

Open multi-selected images

The screenshot shows the OMERO iViewer interface with two main panels displaying microscopy images. Both panels have a vertical stack of thumbnail images on the left and a central image area with a white outline of a cell. In the first panel, a blue box highlights the text "Multi-D Scroll Z projection". A red box highlights the vertical scroll bar on the far left. In the second panel, a red box highlights the vertical scroll bar. White arrows point to specific features within the cell outlines. The top navigation bar includes "File", "ROIs", and "Help" dropdowns, and the title "scram\_06.r3d". The right side of the interface features a toolbar with "Info", "Settings", and "ROIs [3]" tabs. The "Settings" tab is active, showing buttons for "Save", "Undo", "Redo", and "Show Comments". It also includes a "Comment" input field, a "Z" button, and various measurement tools like a square, circle, and triangle. A table lists three regions of interest (ROIs) with columns for Show, Z, T, C, and Comment. The first ROI has values 21, 1, 1, 1, and Comment 10. The second ROI has values 1, 1, 1, 1, and Comment 10. The third ROI has values 1, 1, 1, 1, and Comment 10.

Multi-D Scroll  
Z projection

5.04 μm

Info      Settings      ROIs [3]

Save   Undo   Redo   Show Comments

Z   Comment   10

Show   Z   T   C   Comment

Show	Z	T	C	Comment
<input checked="" type="checkbox"/>	21	1	1	Comment 10
<input checked="" type="checkbox"/>	1	1	1	Comment 10
<input checked="" type="checkbox"/>	1	1	1	Comment 10

Draw and customize ROIs  
Measure  
Extract new images

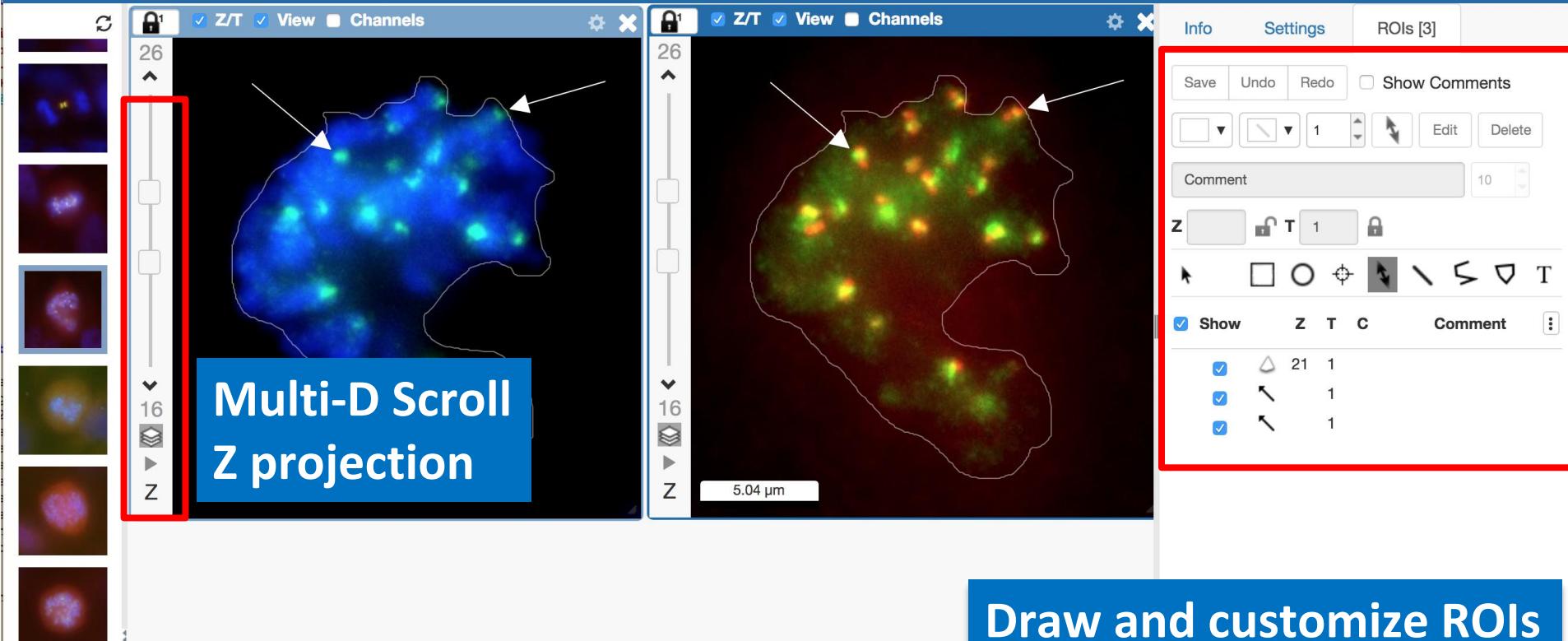


# OMERO iViewer

Open multi-selected images

OMERO File ▾ ROIs ▾ Help ▾

scram\_06.r3d



Zoom and rotate  
Global contrast adjustments

Draw and customize ROIs  
Measure  
Extract new images

# OMERO Figure : forget the other softwares



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OMERO File ▾ Edit ▾ Help ▾ Can't Save Porter et al., Bod1: 2007 Add Image Delete Export PDF

Legends

DNA AuroraB tubulin merged

Control siRNA Bod1 siRNA

Easy alignment to grid

Control siRNA Bod1 siRNA

DAPI HEC1 MCAK

Multi-channel display

P-MCAK ACA DAPI

Zoom

MCAK is not efficiently phosphorylated in Bod1siRNA cells. Data from JCB

125 Edit Figure Legend

The screenshot displays the OMERO software interface. At the top, there's a menu bar with 'File', 'Edit', 'Help', and a status message 'Can't Save'. Below the menu is a title bar 'Porter et al., Bod1: 2007'. To the right of the title are buttons for 'Add Image', 'Delete', and 'Export PDF'. On the far left, a blue box labeled 'Legends' contains four rows of microscopy images: 'Control siRNA' and 'Bod1 siRNA' for DNA, AuroraB, and tubulin channels, followed by a 'merged' row. A large blue box labeled 'Easy alignment to grid' covers the merged row. In the center, there's a 'Multi-channel display' section with two rows of images: 'Control siRNA' and 'Bod1 siRNA' for DAPI, HEC1, and MCAK channels. A second blue box labeled 'Multi-channel display' covers this section. On the right side, there's a 'Zoom' control panel with sliders for 'T' (190, 337, 298) and 'Z' (1/1, 60, 31, 3021), a 'Zoom' slider set to 100%, and a 'Labels' tab in the top right corner. At the bottom, a text box states 'MCAK is not efficiently phosphorylated in Bod1siRNA cells. Data from JCB'. The bottom navigation bar includes a slider for '125' and a button for 'Edit Figure Legend'.

# OMERO Figure : forget the other softwares

OMERO File ▾ Edit ▾ Help ▾ Can't Save Porter et al., Bod1: 2007 Add Image Delete Export PDF

Legends

Easy alignment to grid

Control siRNA

DNA AuroraB tubulin merged

Bod1 siRNA

Control siRNA Bod1 siRNA

DAPI HEC1 MCAK

Multi-channel display

Control siRNA Bod1 siRNA

P-MCAK ACA DAPI

Zoom

No intermediate images

MCAK is not efficiently phosphorylated in Bod1siRNA cells. Data from JCB

125

Edit Figure Legend

The screenshot shows the OMERO software interface. At the top, there's a menu bar with 'OMERO', 'File', 'Edit', 'Help', and a status message 'Can't Save Porter et al., Bod1: 2007'. Below the menu is a toolbar with 'Add Image', 'Delete', and 'Export PDF'. A blue box labeled 'Legends' is on the left. In the center, there's a grid of fluorescence images for 'Control siRNA' and 'Bod1 siRNA' cells, showing channels for DNA (blue), AuroraB (green), and tubulin (magenta). A blue box highlights the text 'Easy alignment to grid'. To the right, a large image shows a merged view of these channels. Another blue box highlights 'Export pdf figures & images'. On the far right, there's a zoom control panel with sliders for 'T' (190, 337, 298) and 'Z' (60, 51, 3021), and a 'Zoom' button. A blue box highlights 'Multi-channel display'. At the bottom, another blue box highlights 'No intermediate images'. A note at the bottom states 'MCAK is not efficiently phosphorylated in Bod1siRNA cells. Data from JCB'. The bottom navigation bar includes a slider set to '125' and an 'Edit Figure Legend' button.

# OMERO Figure : forget the other softwares

The screenshot displays the OMERO software interface with several features highlighted:

- Legends**: A legend panel on the left shows four rows of images: Control siRNA (DNA, AuroraB, tubulin, merged), Bod1 siRNA (DNA, AuroraB, tubulin, merged), Control siRNA (DAPI, HEC1, MCAK), and Bod1 siRNA (DAPI, HEC1, MCAK). A blue box highlights the "Easy alignment to grid" feature.
- Export pdf figures & images**: A blue box highlights the "Export PDF" button in the top right menu bar.
- Multi-channel display**: A blue box highlights the main image viewer showing a merged channel of DNA (blue), Phospho-H3 (green), and AuroraB (red).
- Zoom**: A blue box highlights the zoom controls at the bottom right, which include a zoom slider from 100% to 3021, a T (translate) slider from 190 to 600, and a Z (depth) slider from 337 to 3021.
- No intermediate images**: A blue box highlights the text "MCAK is not efficiently phosphorylated in Bod1siRNA cells. Data from JCB" at the bottom center.
- Easy scaling**: A blue box highlights the zoomed-in views of the multi-channel images.

# OMERO Figure : forget the other softwares

The screenshot shows the OMERO software interface with several features highlighted:

- Legends**: A blue box on the left side.
- Easy alignment to grid**: A blue box highlighting the alignment tools for multiple images.
- Get ROIs of Zoom**: A blue box highlighting the region of interest (ROI) selection and zooming features.
- Easy scaling**: A blue box highlighting the scaling tools.
- Export pdf figures & images**: A blue box on the right side.
- Multi-channel display**: A blue box highlighting the multi-channel image display.
- Zoom**: A blue box highlighting the zoom controls.
- No intermediate images**: A blue box highlighting the direct figure assembly feature.
- Easily assemble your scientific figures !**: A large green box at the bottom right.

At the bottom of the interface, a status bar displays the text: "MCAK is not efficiently phosphorylated in Bod1siRNA cells. Data from JCB".



# Image Analysis

## Local Desk

ImageJ/Fiji



Matlab



Python



Cellprofiler



Orbit



WEB



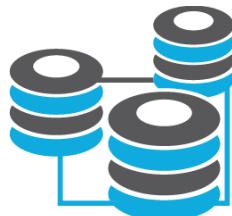
## Galleries & Data bases



Marimba

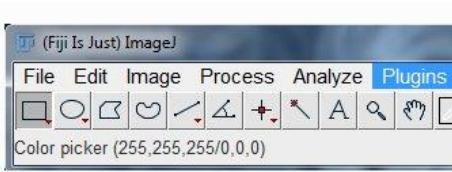


 OMERO

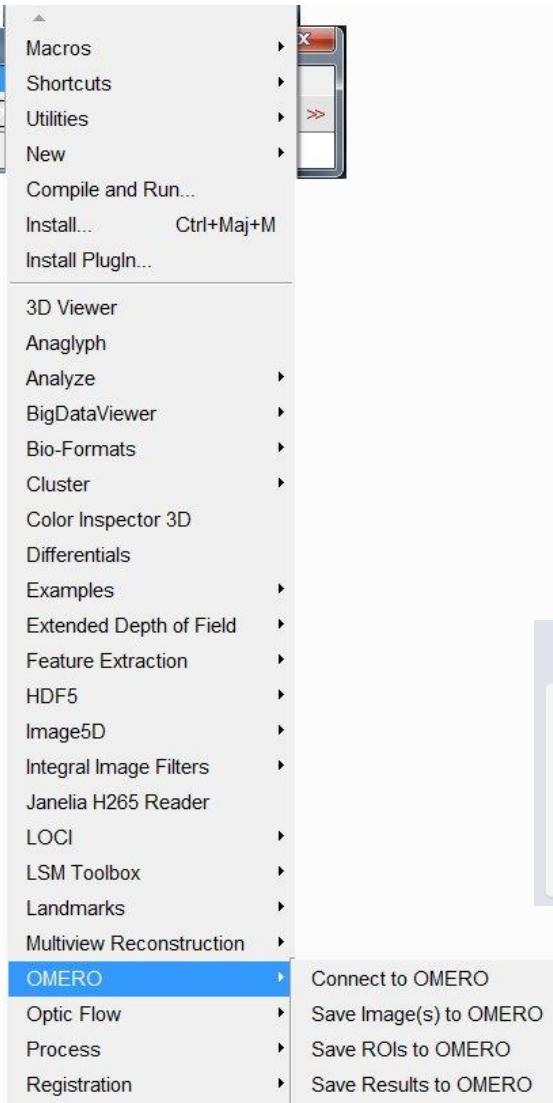




# Image Analysis



## Local Desk



**OMERO**



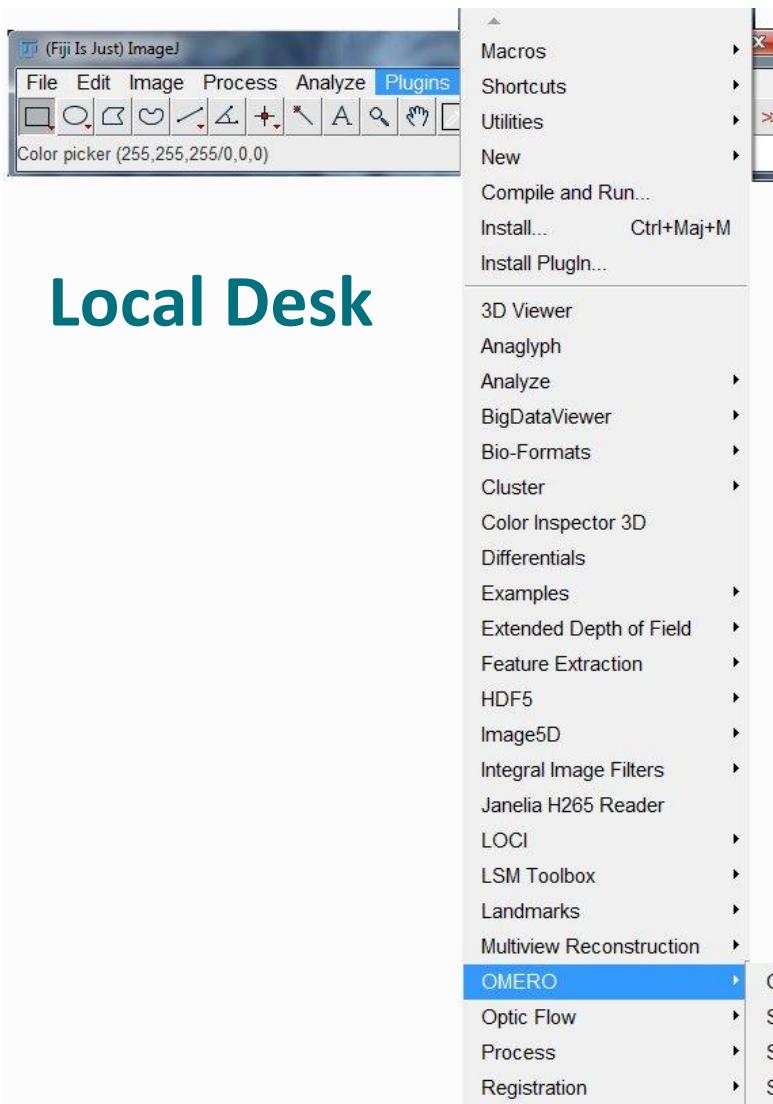
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DE BIOINFORMATIQUE



# Image Analysis

## Macro / Python / Groovy

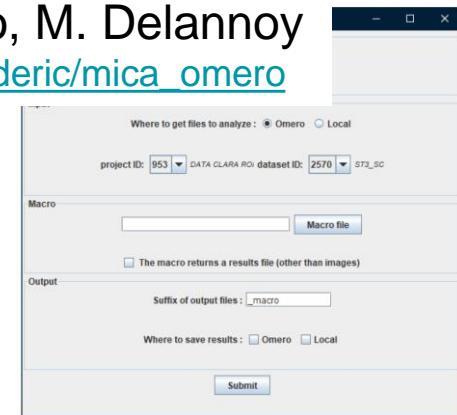
### Local Desk



### GUI to batch Macros

C. Belle, M. Carvalho, M. Delannoy

[https://github.com/brau-frederic/mica\\_omero](https://github.com/brau-frederic/mica_omero)



# OMERO





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# Image Analysis

## Virtual Desk

ImageJ/Fiji

Matlab

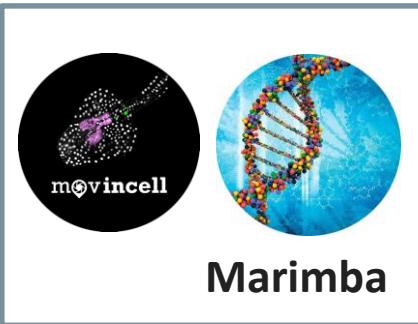
Python

Cellprofiler

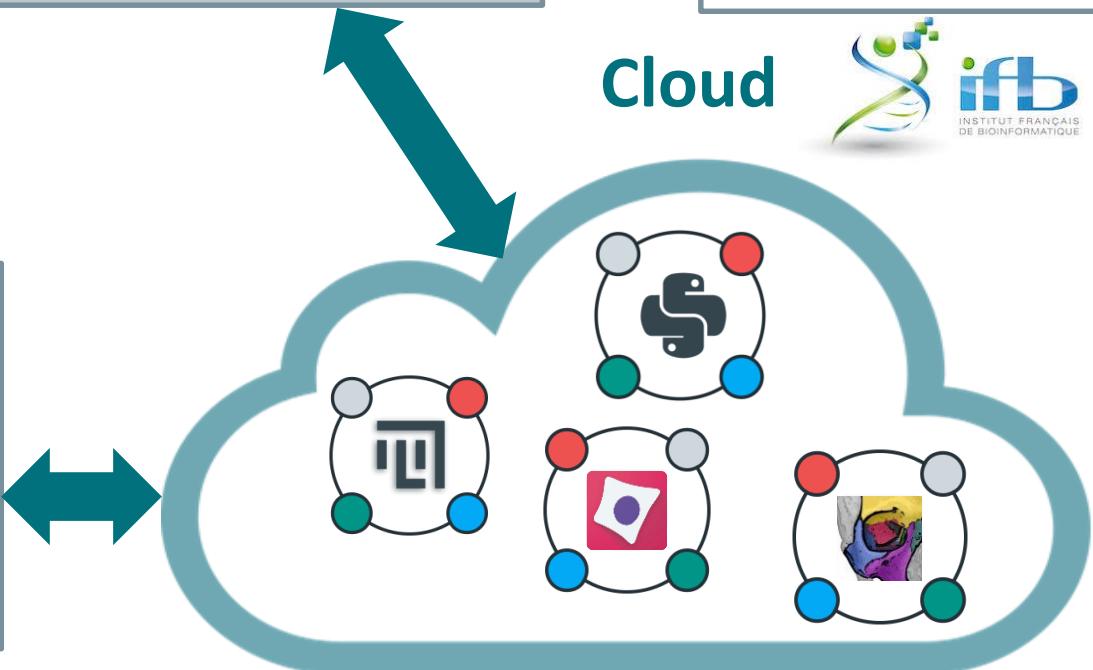
Orbit

WEB

## Galleries & Data bases



## Cloud





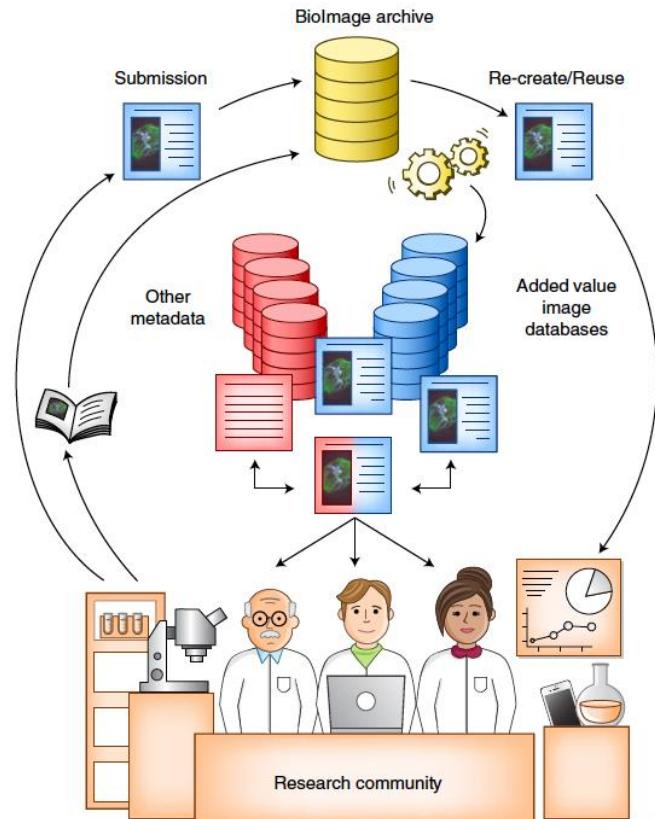
# FAIR Management

## Metadata integration : format compatible IDR

- **Findable**— Original Image data linked to Paper by DOI
- **Accessible**— Metadata retrievable, open API
- **Interoperable**— Uses common defined vocabularies
- **Reusable**— Licensed (CC-BY 4.0), provenance included



An online, public resource that stores, integrates and serves image datasets from published scientific studies





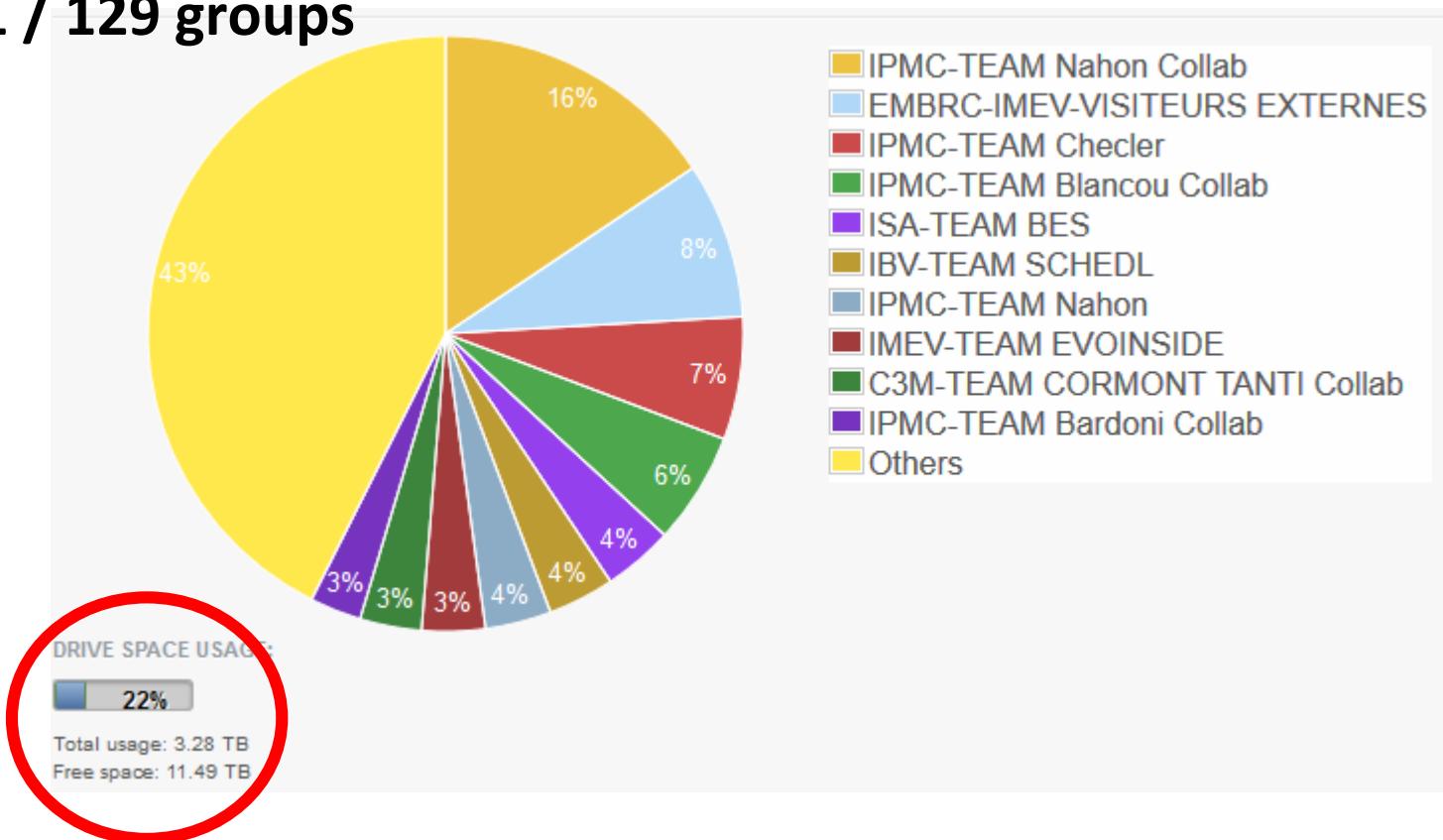
# Usage report

June 2019 / today

292 / 355 users

111 / 129 groups

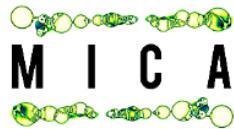
⇒ 3,2To / 15 To ⇒ + de 21 000 images



Flexible storage volume according to needs



# Communication & support



MICROSCOPIE  
IMAGERIE  
CÔTE D'AZUR

Website



Protocols  
Links  
FAQ  
Data  
Management  
Plan (DMP)



Diffusion list



[omero-users@univ-cotedazur.fr](mailto:omero-users@univ-cotedazur.fr)  
Informations  
Notifications  
Maintenances



[omero-support@univ-cotedazur.fr](mailto:omero-support@univ-cotedazur.fr)  
Users  
questions



Policy



DMP  
for Optimized  
Sharing and  
Interoperability of  
Research Data



Website



Guides  
Tutorials  
User Forum



# Communication / trainings

**Trainings** 11 teams

+ > 50 people individually

1 advanced session for referents

**1 CNRS session – Next in April**

**Launching seminars**

6 institute meetings & 1 CdE

1 Scientific day

**Users workshop/ Training Omero**

Dundee 2019

Montpellier 2020



April 2019 OMERO users workshop

Two days workshop on how to use OMERO held in Dundee.

# Thank you for your attention



## Académie 4



S. Benaicha  
C. Baron



S.Ben Aïcha - F. Bekkouche - R. Arkowitz - F. Brau



C. Blanchet



# DEMOCRATISONS LA SUPER-RESOLUTION

M I C A  
MICROSCOPIE  
IMAGERIE  
CÔTE D'AZUR

