

BisQue

Scientific image data management and
analysis in the Cloud

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Center for Bio-Image Informatics, UCSB, 2014

- Textual and 5D graphical annotations
- Biologically meaningful objects and groups

- Large-scale
- Cluster support
- Data parallelization
- Extensible



- 5D large images
- 200+ formats
- Data summarization

- Scalable and distributed
- Everything is web accessible
- Searchable

BisQue

No need to customize

Kosik Lab

3D CLSM analysis
+Metadata
condition, channels
Results:
Nucleus location,
Cell classification,
Count PHF+/PHF-
Cell Density

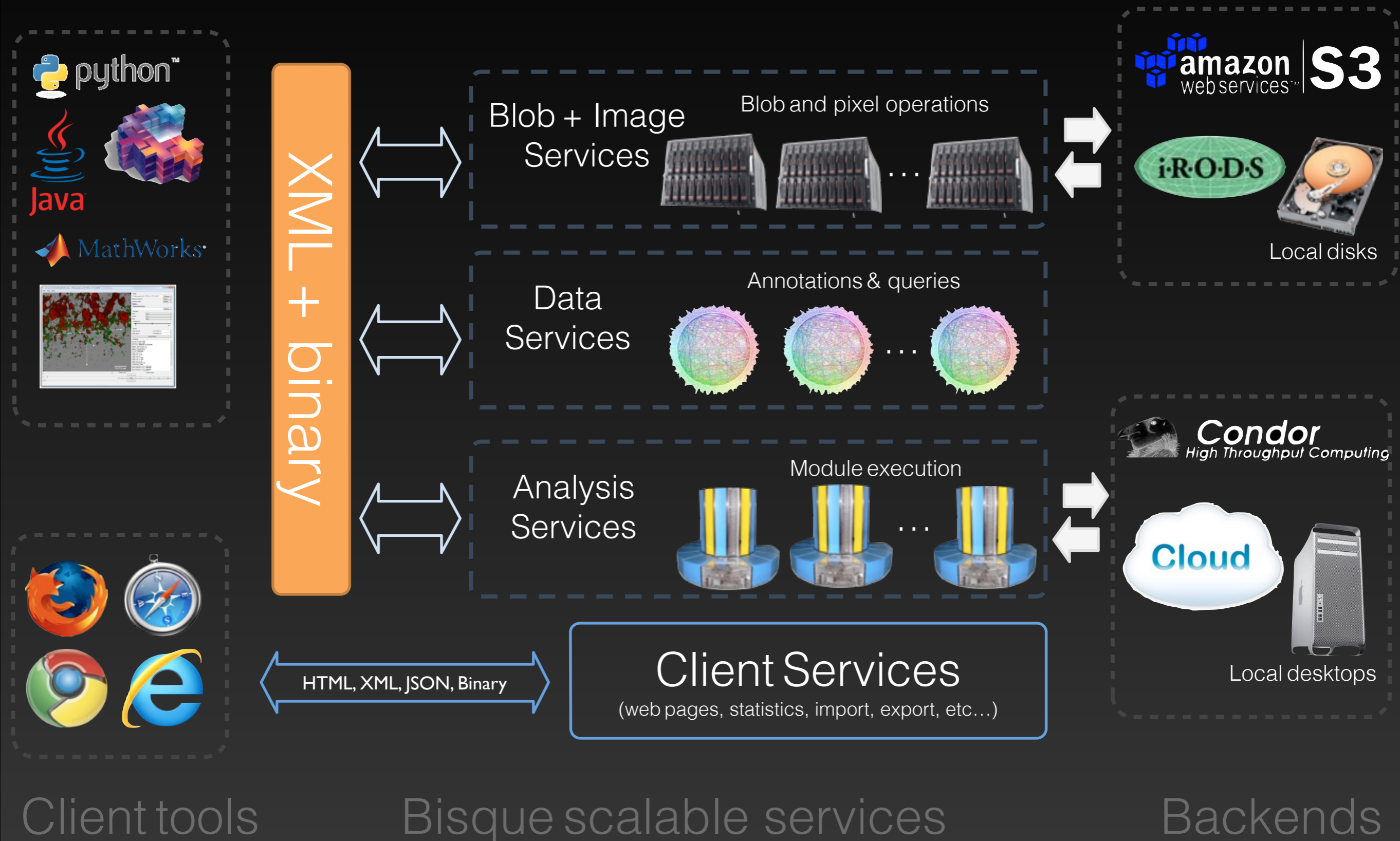
Greenpeace

Video analysis
900GB Video
+Metadata
depth, location
Results:
Object Location,
Object type,
Laser dots,
resolution...

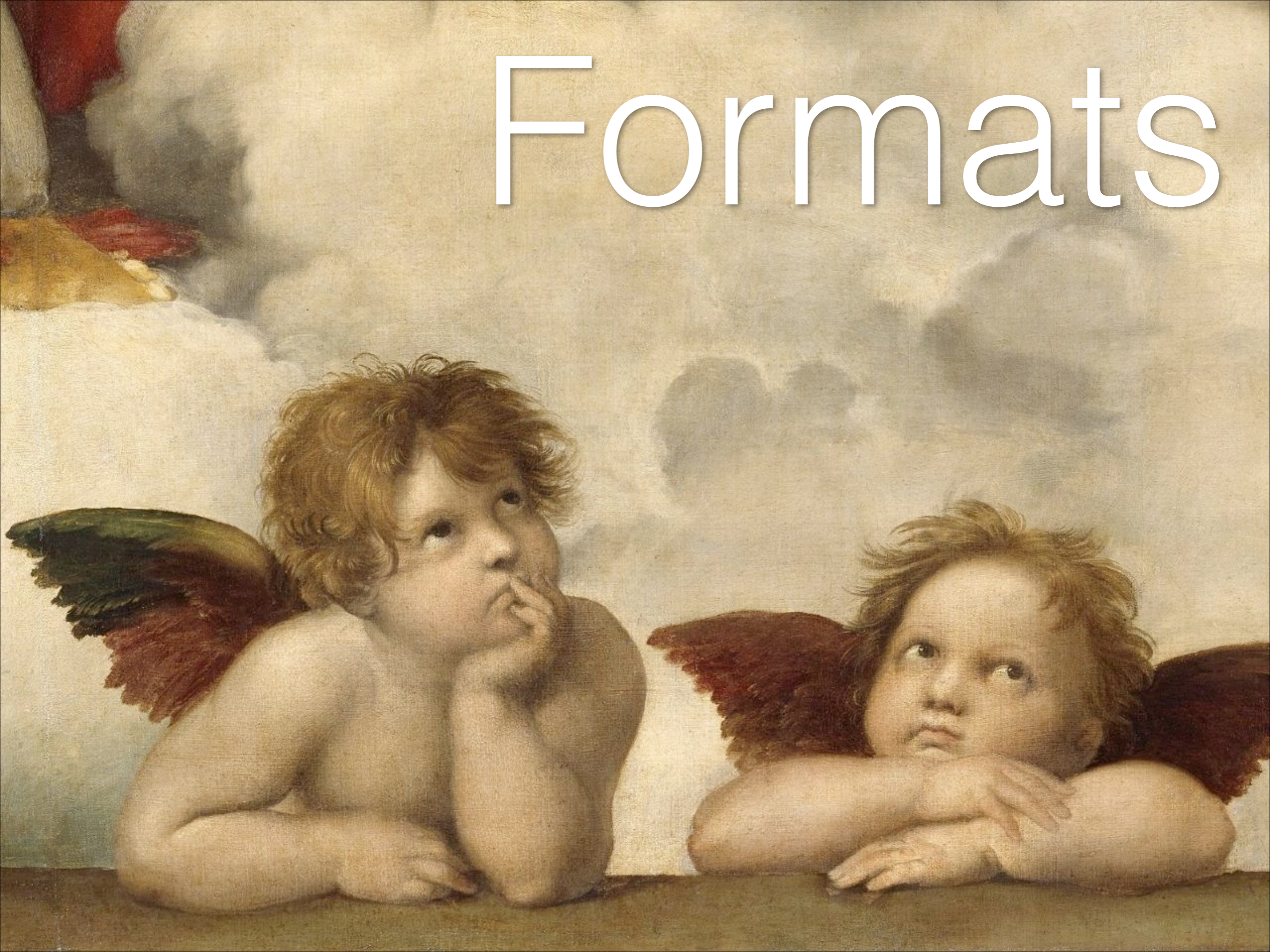
Spalding Lab

HT 4D Analysis
100GB Timeseries
+Metadata
conditions, mutaton...
Results:
Root tips in time,
Seed perimeter,
Root angle,
seed area...

Bisque architecture



Formats



240+ Image formats

Bisque supports pluggable image en/de-coders

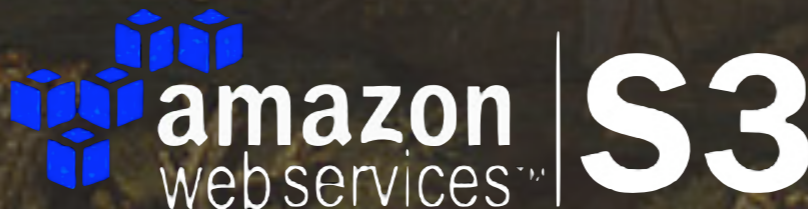
- **Native** -
Most popular 5D images and videos
- **OpenSlide** -
whole-slide images
- **Imaris** - (exclusively for Bisque)
Industry leading microscopy support
- **BioFormats** -
Largest open-source bio-image library

multi-file formats

- Any combination multi-file-multi-image
- In-place support with no re-conversion
- 5D series (e.g. TIFF Z series)
- User correctable geometry/resolution
- Proprietary (e.g. Volocity, Slidebook)
- Import/Export

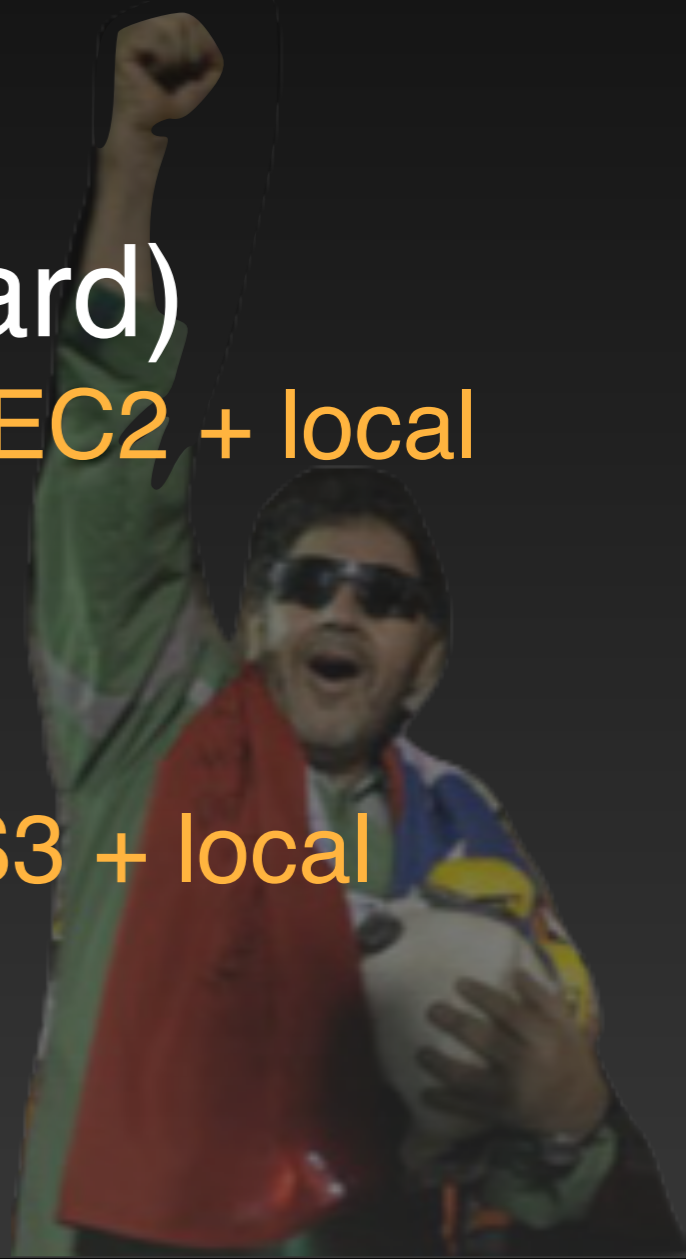
On shoulders of giants

- High-throughput analysis,
- Distributed storage,
- On-demand computation



16 tools, what do you get?

- Click to Bisque (credit card)
 - Rightscale template for Amazon EC2
- Datacenter per dataset (credit card)
 - On-demand condor nodes on Amazon EC2 + local
- 99.999% reliability
 - Distributed data on iRODS + Amazon S3 + local





Bisque UI

- Web based - HTML5
- Query everything
- Integrated image analysis

Bisque: Welcome to the Bi x

bisque.ece.ucsb.edu/client_service/

Bisque + Create Upload Download Analyze Browse Find resources using tags

My recent stuff

Datasets Images Analysis

Showing 1-8 of total 175

nd3d-sim-01.ome.tif

wtheadstack_compressed...

combinedsubtractions.lsm

Rab582 3d det gfapBrdU...

JR341601.ome.tif_filt...

JR341601.ome.tif

FV1000-2008-12-18 12.5...

smith_z_stack_simple.o...

20 t000

Bisque usage

Image uploads - monthly

Month	Image uploads (approx.)
Mar '13	0
Apr '13	0
May '13	0
Jun '13	0
Jul '13	75,000
Aug '13	0
Sep '13	10,000
Oct '13	0
Nov '13	0
Dec '13	0
Jan '14	0
Feb '14	0
Mar '14	70,000

Image Viewer

- Very large images (our PanoJS3)
- 5D support – projections, navigation and movies
- Dynamic tiles – enhancement, channel fusion
- Pixel formats (up to 64bit integer, 80 bit float)
- Mobile support: gestures

Our PanoJS3 is used by many projects including OMERO and Journal Cell

2D viewer/editor

Image enhancements
Channel fusion
Projections

5D graphical annotations

Analysis results

Metadata

Annotations

Graphical

Embedded

Analysis

Map

Create custom

Graphical annotations

Type

circle

ellipse

label

line

point

polygon

polyline

rectangle

square

Cell

Tree of annotations

Visibility

Add

Delete

Color

Stats

Type:Name	Value
<div> <div> <div></div> <div>T</div> </div> <div>NuclearDetector3D</div> </div>	2014-02-22 08:50:20
<div> <div> <div></div> <div>Sepal: Sepal</div> </div> </div>	
<div> <div> <div> <div></div> <div>Cell: Cell</div> </div> <div> <div> <div></div> <div>circle</div> </div> <div> <div></div> <div>polygon</div> </div> </div> </div> </div>	<div>311,242,23,0,0; 322,242,23,0,0;</div> <div>11,363,23,0,0; 71,232,23,0,0; 178,146,...</div>

Bisque: Resource viewer

vidi.ece.ucsb.edu:9090/client_service/view?resource=http://vidi.ece.ucsb.edu:9090/data_service/image/2710

Bisque

Create Upload Download Analyze Browse Find resources using tags

Download Share Visibility: published Delete View: 2D Export Operations

Bisque admin

image: MC00111.svs

Image: 95700x67916 ch: 3/8bits Scale: 1.5625%

Region of interest

Point of interest

1345.6128 um (94208,42240)px (3018273.59,1353302.02)um

Metadata

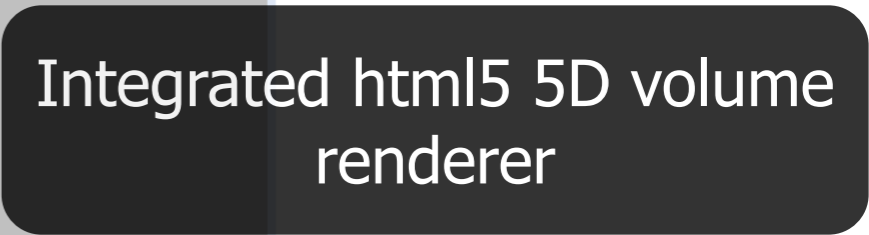
Annotations Graphical Embedded Analysis Map

Export

Name	Value
T channel_0_name	red
T channel_1_name	green
T channel_2_name	blue
T channel_color_0	255,0,0
T channel_color_1	0,255,0
T channel_color_2	0,0,255
custom	
aperio	
T AppMag	40
T Date	09/18/11
T DisplayColor	0
T DSR ID	149.171.101.36
T Exposure Scale	0.000001
T Exposure Time	109
T Filename	MCO CRC Cohort 0111
T Filtered	3
T Focus Offset	-0.000100
T ICC Profile	ScanScope v1
T ImageID	36787
T Left	29.586740
T LineAreaXOffset	0.013821
T LineAreaYOffset	-0.000312
T LineCameraSkew	0.001294
T MPP	0.2503
T OriginalHeight	68016
T OriginalWidth	100000
T ScanScope ID	SS1379
T StripeWidth	1000
T Time	20:11:58
T Time Zone	GMT+10:00
T User	a2d9f8cc-87b9-4fdb-a6e2-ec430f4c9d98

100K x 100K whole slide
annotations stored in Bisque

Textual annotation
Embedded metadata



Bisque: Resource viewer

bisque.ece.ucsb.edu/client_service/view?resource=http://bisque.ece.ucsb.edu/data_service/image/3976100

Bisque

Create Upload Download Analyze Browse Find resources using tags

Download Share Visibility: private Delete View: 2D Export Operations image: Rab582 3d det gfapBrdUmib1z24Live9-14-2007_11-55-58_AM copy.tif

Image: 512x512 ch: 3/16bits Scale: 200%

Measurements:
via graphical objects

10.4409 um (487,289)px (60.53,35.92)um

Metadata

Annotations Graphical Embedded Analysis Map

Create custom

Graphical annotations

Type

circle
ellipse
label
line
point
polygon
polyline
rectangle
square

17-Mar-2009 14:46:57

Tree of annotations

Visibility Add Delete Color Stats

Type:Name	Value
<input checked="" type="checkbox"/> line	283,278,0,0,0; 298,247.5,0,0,0;

Bisque: Resource viewer

bisque.ece.ucsb.edu/client_service/view?resource=http://bisque.ece.ucsb.edu/data_service/image/3976100

Bisque

Create Analyze Browse Find resources using tags

Download Share Visibility: private Delete View: 2D Export Operations

image: Rab582 3d det gfapBrdUmib1z24Live9-14-2007_11-55-58_AM copy.tif

Measurements:

area estimation via segmentation

Pixel Counter

Regional counts

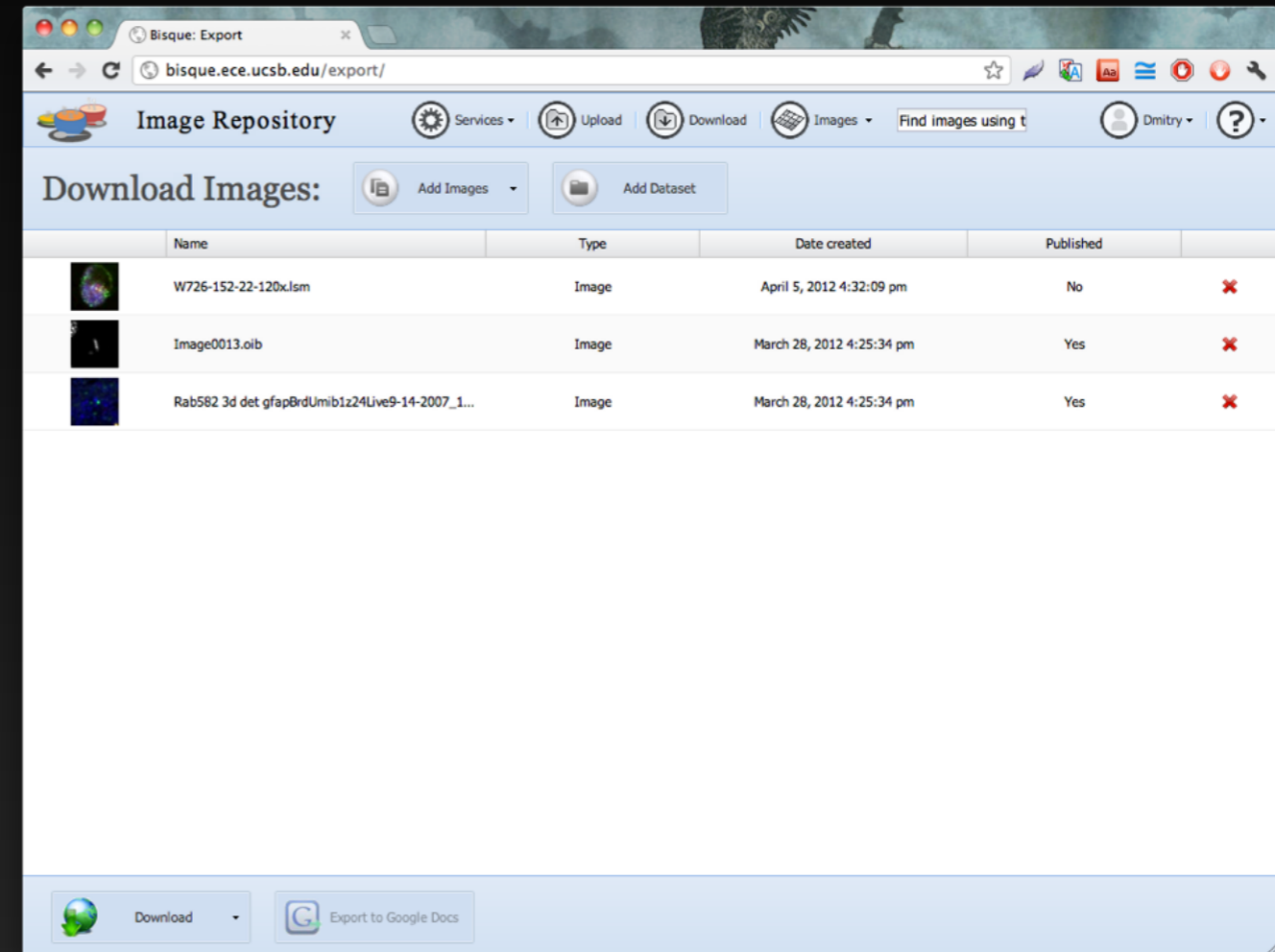
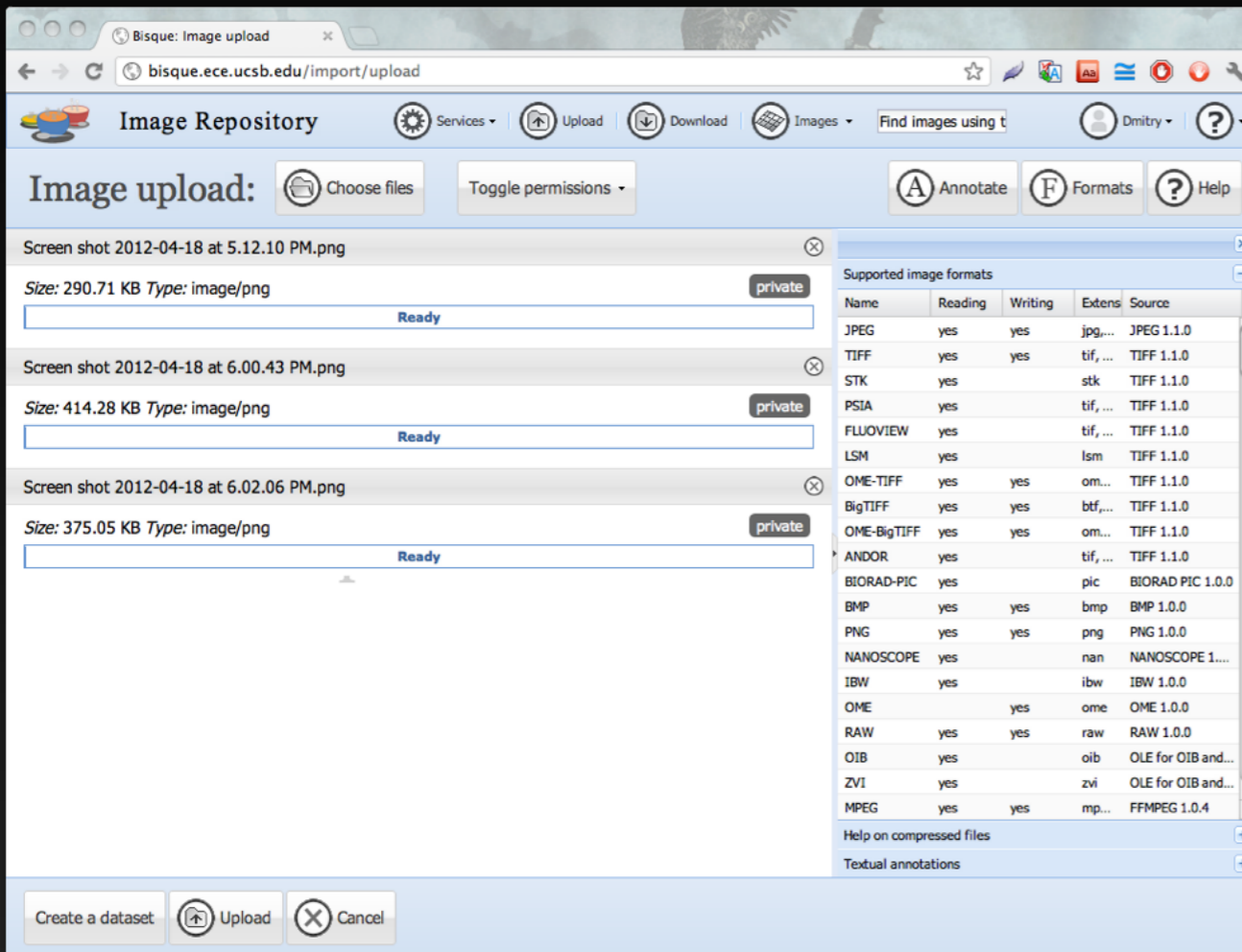
Close

Regional Counts

Click on any part of the image to segment out a region. The pixel count will be displayed below.

centroid	pixels	microns ²
363,106	986	60.93

Import and Export



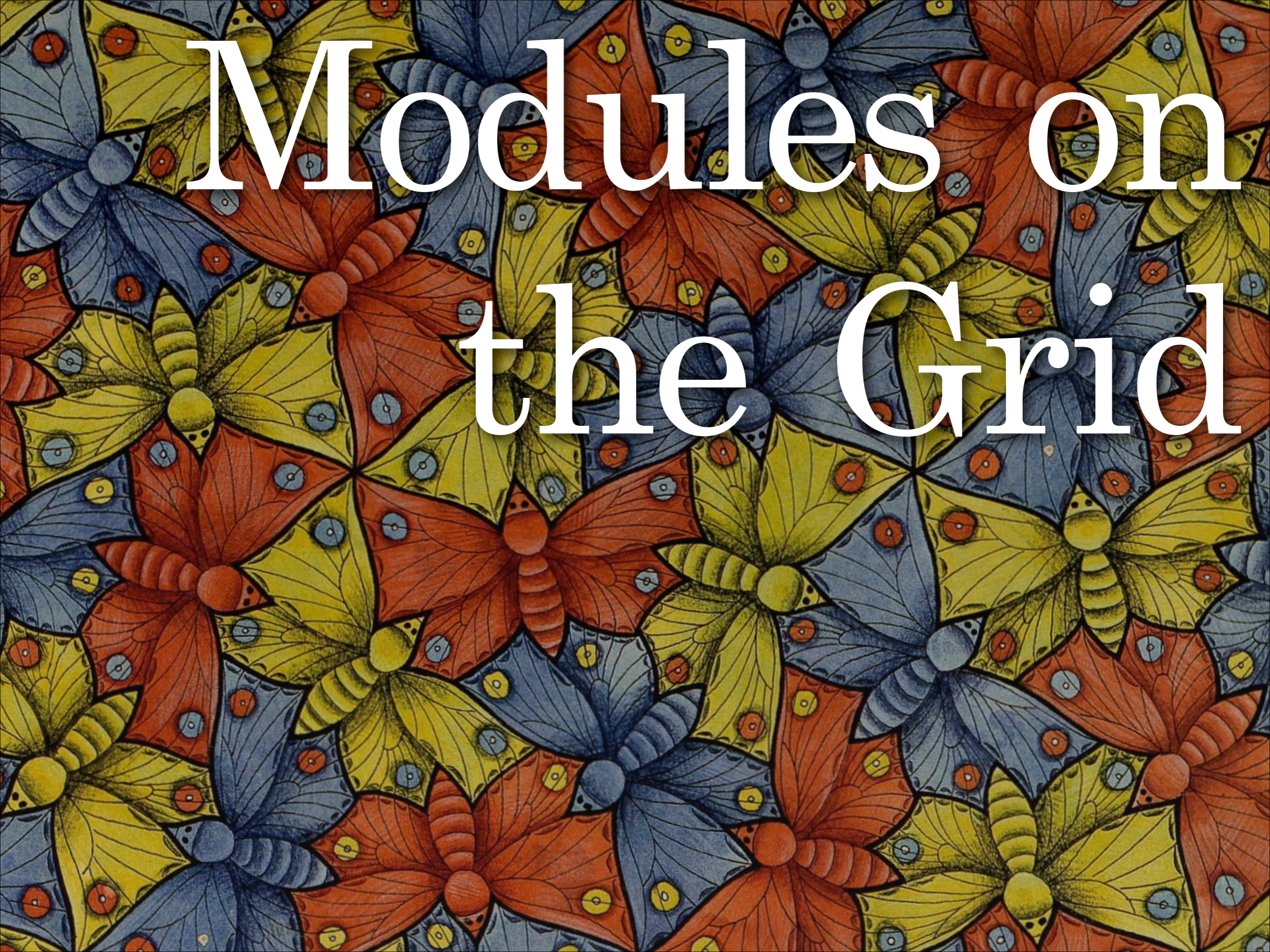
- support for many large files, directories
- plugin free HTML5 uploads
- drag/drop
- import annotations
- processing compressed files
- compose 5D from multiple files

- on the fly streaming export
- dataset export
- export annotations
- Formats: Zip, Tar, GZip, BZip

- Share images, analysis and annotation with collaborators (Google style)
- Publish data for public access and search
- No headache about formats and location
- Simple import of 5D images and annotations
- Export sets of images and metadata
- Protected private data



Modules on the Grid



Modules run on the cluster

- Scheduling with Condor grid
- Amazon EC2
- Low cost on-demand pricing (EC2)

Automatic parallelization for datasets

Easy to create

- Wrapping binaries
- APIs: Matlab, Python, Java, C++
- Pipelines: ImageJ, CellProfiler
- Automated and rich web UI

Bisque infrastructure

- Basic image access
- Storage and retrieval of textual and graphical annotations
- Basic statistics and plotting

Bisque WebApp

Detector of nuclei in 3D/4D

Version: 3 Authors: Dmitry Fedorov and Boguslaw Obara
This application detects centroids of nuclei in 3D/4D fluorescence images.

1. Select data for processing:

Select input resource:
Select an Image or Select a set of images or even Upload local images

Planes View Annotations: Navigate Select Polygon

Image: 2048x2048 Z:33 ch: 3/8bits Scale: 25%

106.8870 um

Inputs

Help and workflow

This is a public Beta 3, so please, bear with us if something breaks and don't hesitate to leave your comments, bug reports and suggestions to us!

This application allows you to select local file or an image in the Bisque database and automatically identify nuclear centroid locations in 3D/4D fluorescent microscopic imagery. These locations can be exported as text, excel file, XML or as a 3D visualization. Simple statistics can be calculated. Furthermore, additional analysis is possible.

We start with a laser scanning confocal 3D stack that has a nuclear stain channel. In 3D the stack looks like this:

Single plain of the stack looks like this:

2. Parameters:

Nuclear channel: 1: Red
Nuclear diameter: 5
Voxel resolution X: 0.3181 Y: 0.3181 Z: 0 microns

Input parameters

3. Run algorithm:

Run This may take some time, progress will be shown on the button.

4. Results:

The module ran in 7 minutes 50 seconds

Detection summary

Name	Value
count	1

Detected nuclear centroids

Planes View

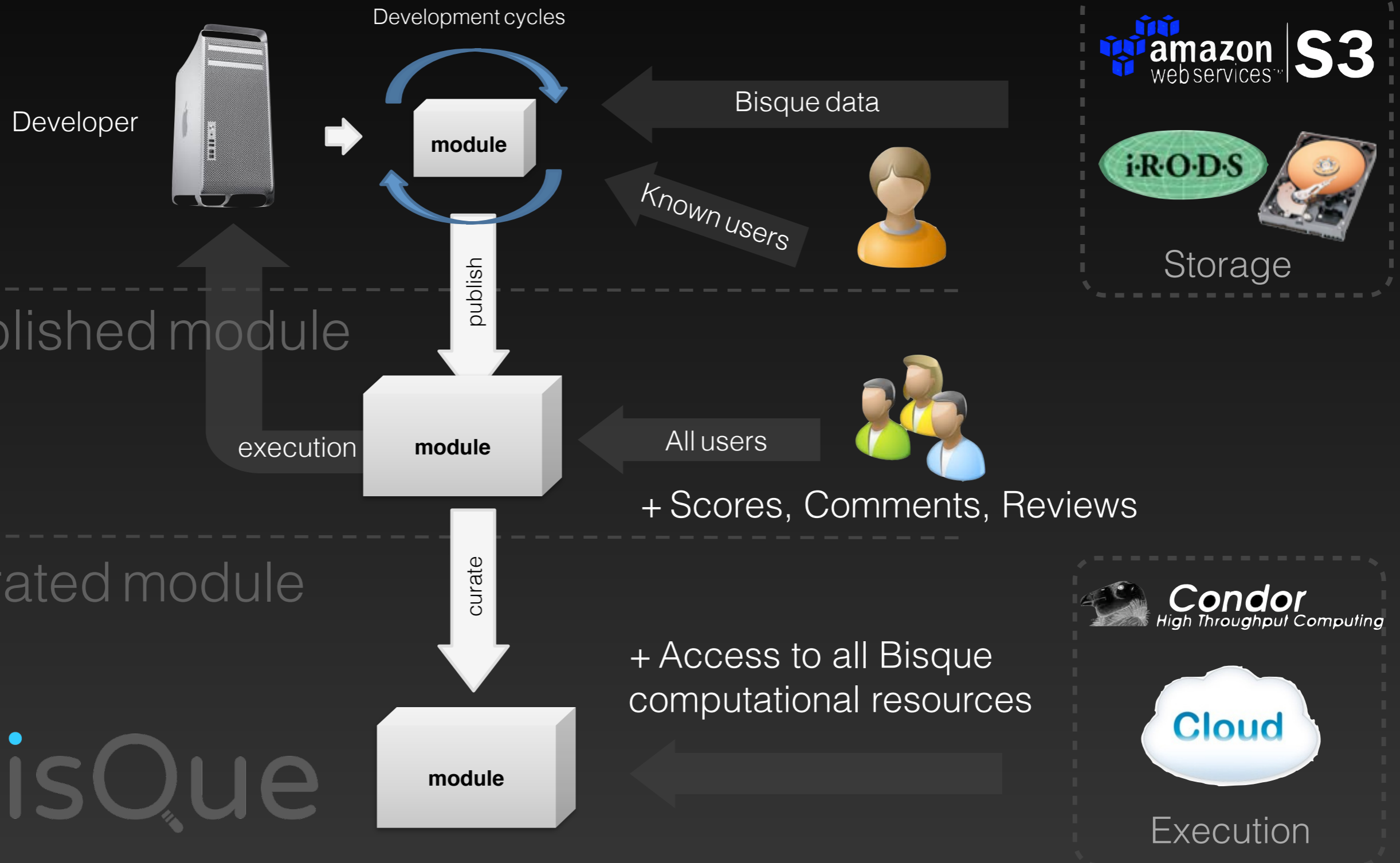
Image: 2048x2048 Z:33 ch: 3/8bits Scale: 25%

106.8870 um

Outputs

Module lifetime

Private module



Running in Bisque



ImageJ pipelines

- Pipelines (scripts) runnable through Bisque
- Support for clusters and datasets
- Results stored and visualized in Bisque



CellProfiler pipelines

- Existing pipelines runnable through Bisque
- Support for clusters and datasets
- Results stored and visualized in Bisque
- Future:
 - Customizable pipelines

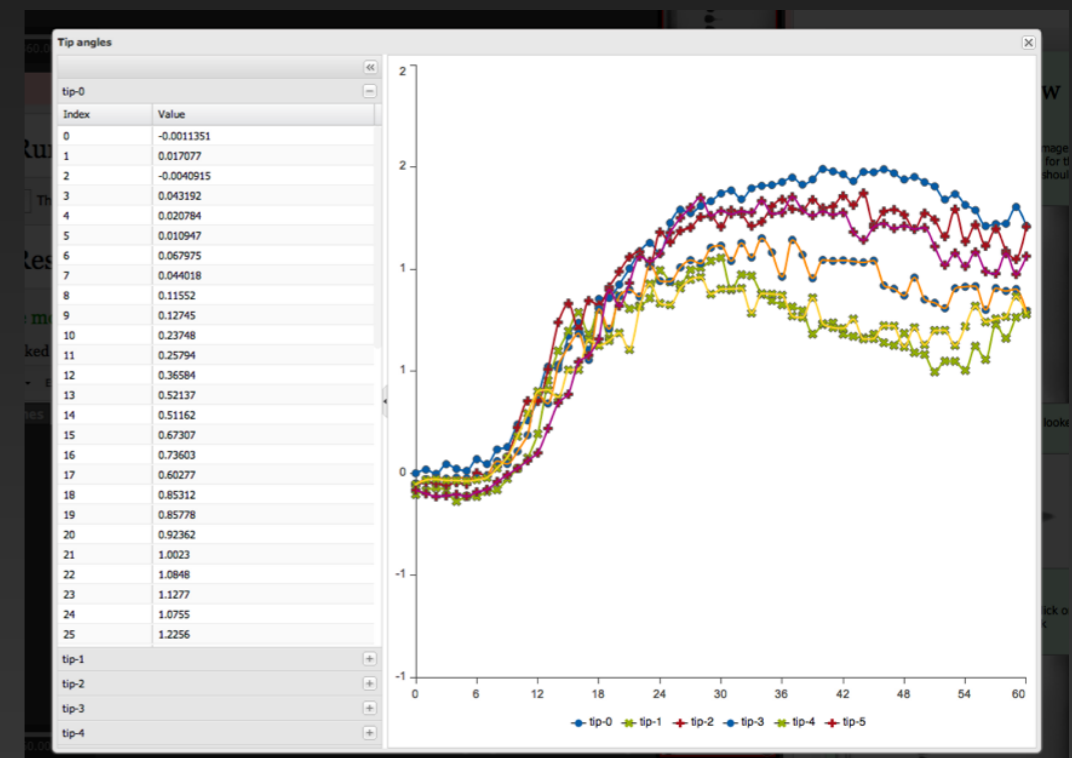
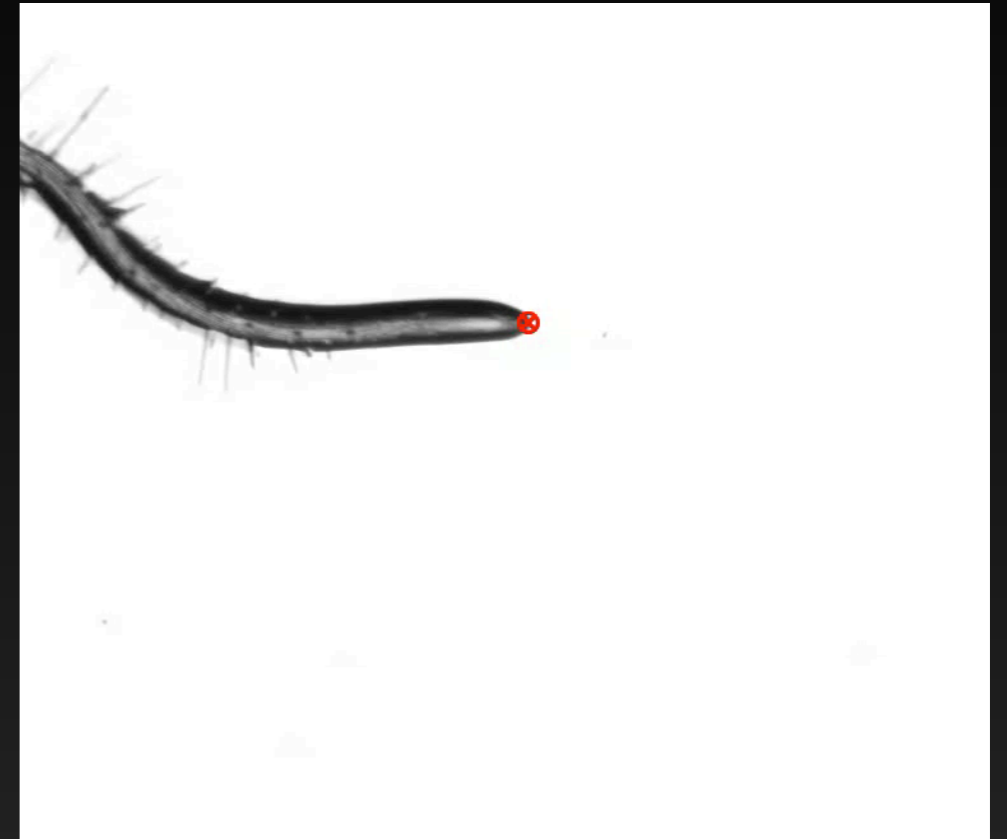




Our analysis

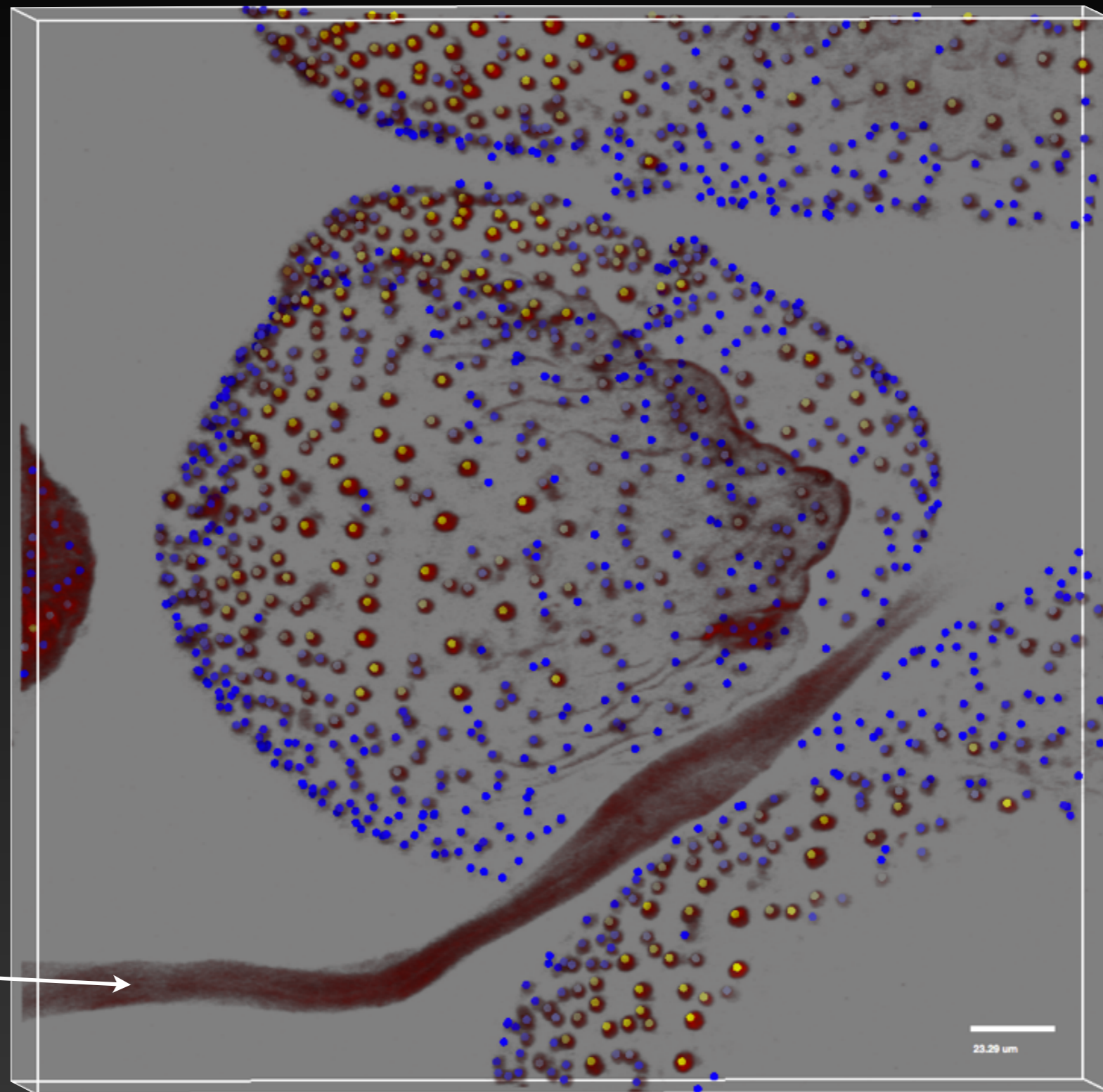
BQPhytomorph

1. Whole seed-size analysis
2. Root tip angle / Multi-tip and growth-rate
3. Root branching



Collaboration with Spalding Lab, University of Wisconsin and iPlant

Detection of nuclei in 3D/4D



Centroids color coded by confidence from blue to yellow (highest)

Image contributed by Adrienne Roeder, Cornell

Rendered with bioView3D

Nuclear channel
pre-filtered with
membrane
channel

Collaboration with Meyerowitz lab @ Caltech and Roeder Lab @ Cornell

Detection of nuclei in 3D/4D

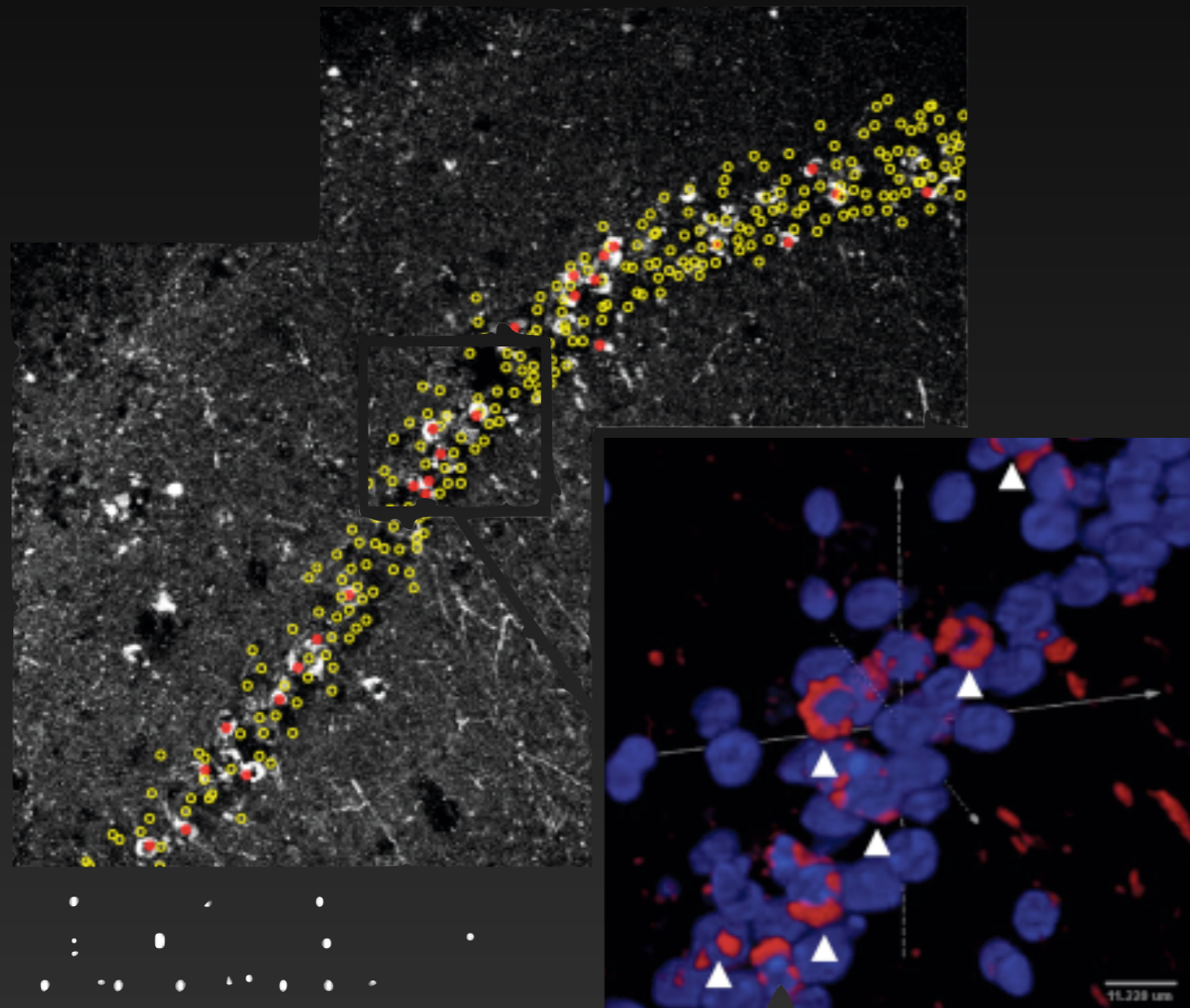
- Detects nuclei in up to 4D
- Supports GPU acceleration*
- Filtering with uncorellated channels
- Works on images of any size
- Each centroid has confidance

** GPU support partially sponsored by Nvidia with hardware donation*

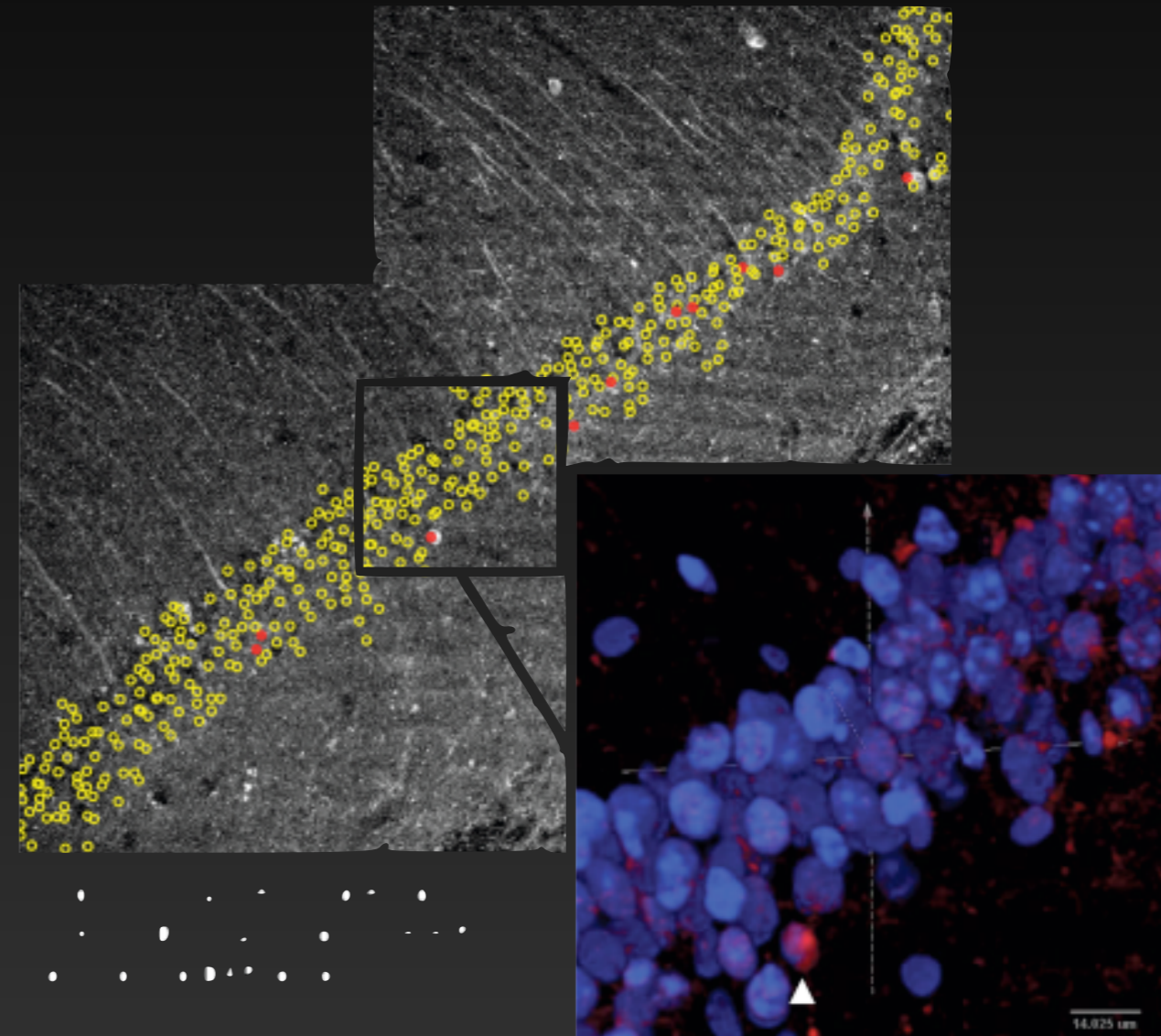


Cell classification

Control

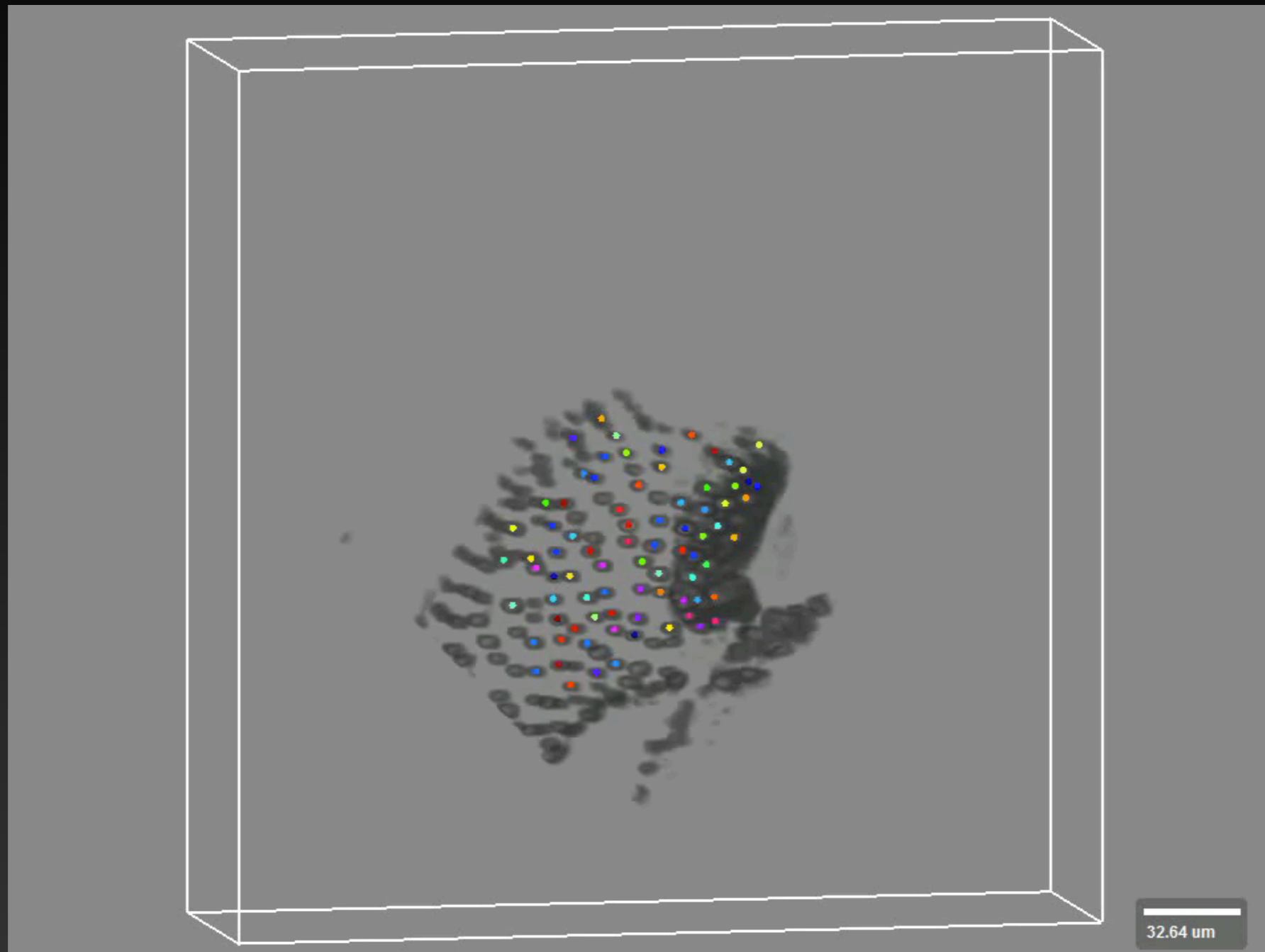


Treatment



Classified nuclei overlaid on PHF Z projection: Red – PHF positive, Yellow – not PHF positive

Cell tracking

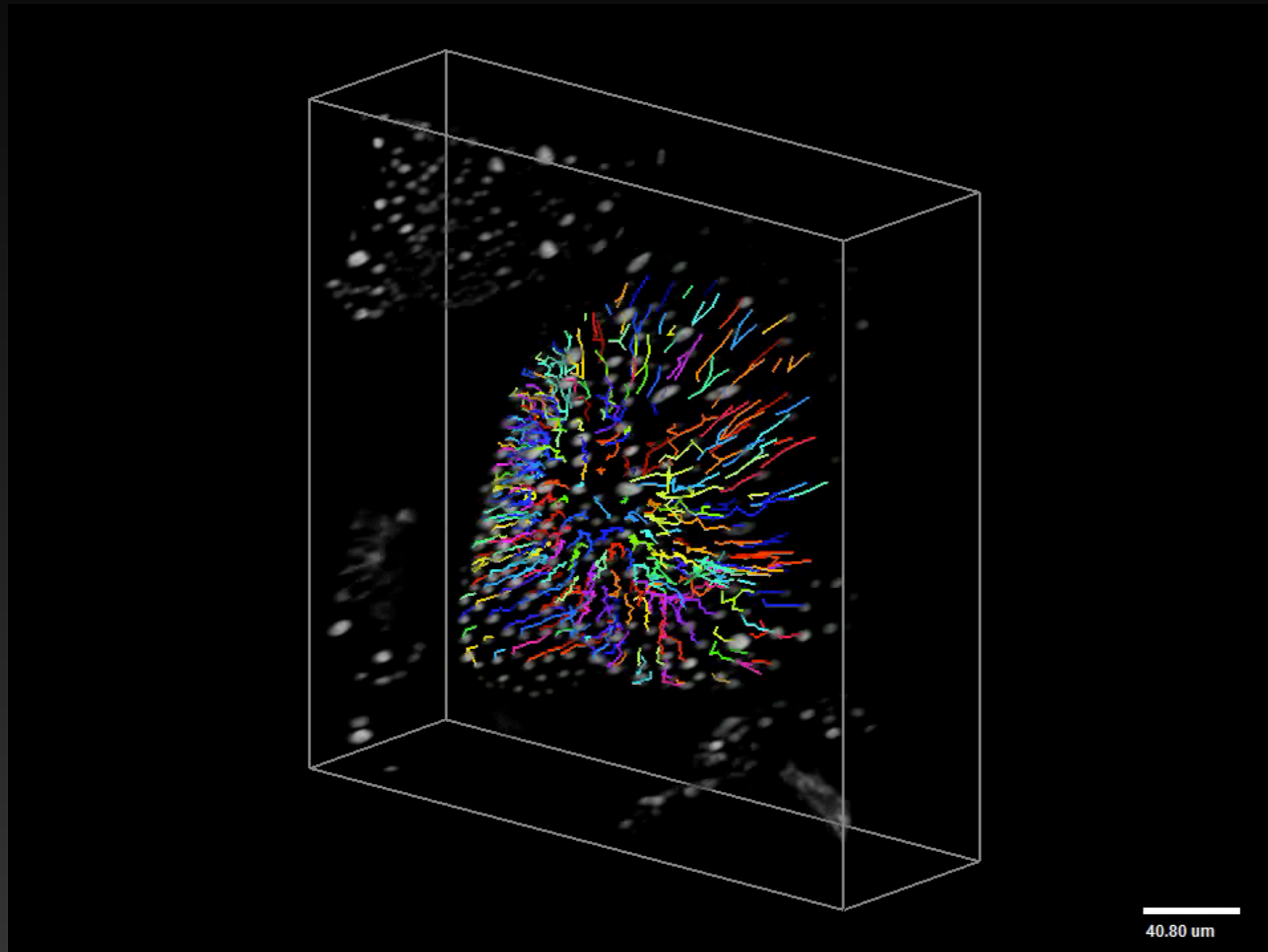


Centroids color
coded by lifeline

4D dataset
contributed by
Adrienne Roeder,
Cornell

Movie rendered
with bioView3D

Cell tracking



Lifelines projected
onto last time
point

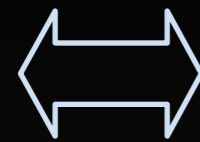
4D dataset
contributed by
Adrienne Roeder,
Cornell

Movie rendered
with bioView3D

The background is an abstract composition of organic, textured shapes. On the left, there's a bright, almost white area that transitions into a vibrant green. This green area then blends into a deep, dark blue that covers the right side of the image. The textures are reminiscent of water ripples, marbled paper, or perhaps a microscopic view of certain minerals. The overall effect is dynamic and visually rich.

Indexing and feature descriptors

XML + binary



Query Services

Query dialects and extensions: XQuery, SPARQL, BisQue,...

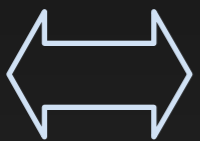


Index Services



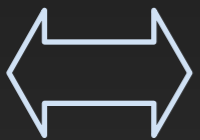
Feature Services

Extractors: Shape, Color, Location, Texture, etc...



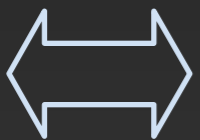
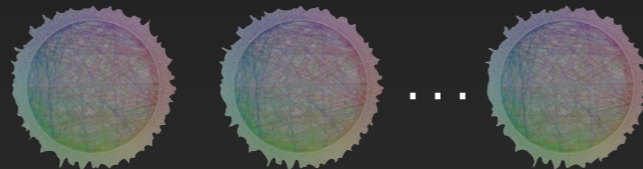
Blob + Image Services

Blob and pixel operations



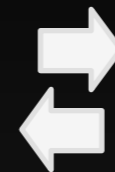
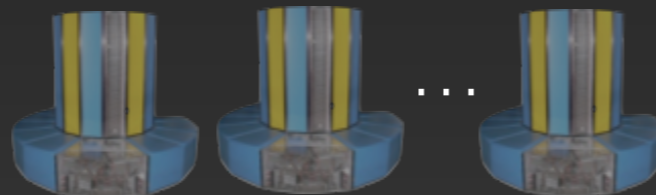
Data Services

Annotations & queries



Analysis Services

Module execution



Full text

Geometry



Image similarity

Graph



Near duplicates

Bisque scalable services

Indices

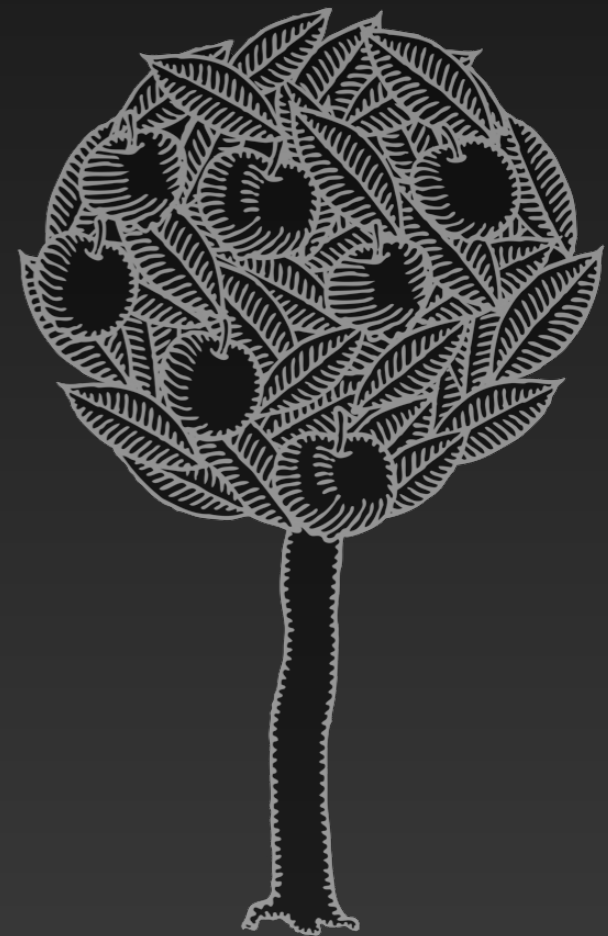
Feature service

- Uniform access to feature descriptor computations
- Distributed computation
- Efficiently cached
- Feature descriptor libraries:
 - MPEG7 (6: DCD, CSD, CLD, RSD, EHD, HTD)
 - OpenCV (4: SIFT, SURF, ORB, BRISK)
 - Mahotas (5: DCD, CSD, CLD, RSD, EHD)
 - WNDCharm (100+: Radon, Chebyshev, Gabor, Histograms, Moments, Tamura, Edge stats, Otsu objects, Zernike, Haralick, ...)



Index services

- Multi-modal case-based index structures
(custom clustering/indexing, distance measures, etc...)
- Use feature services for descriptor computations
- Designed to answer specific searches:
 - Full text
 - Graph
 - Geometry
 - Image similarity



Query service

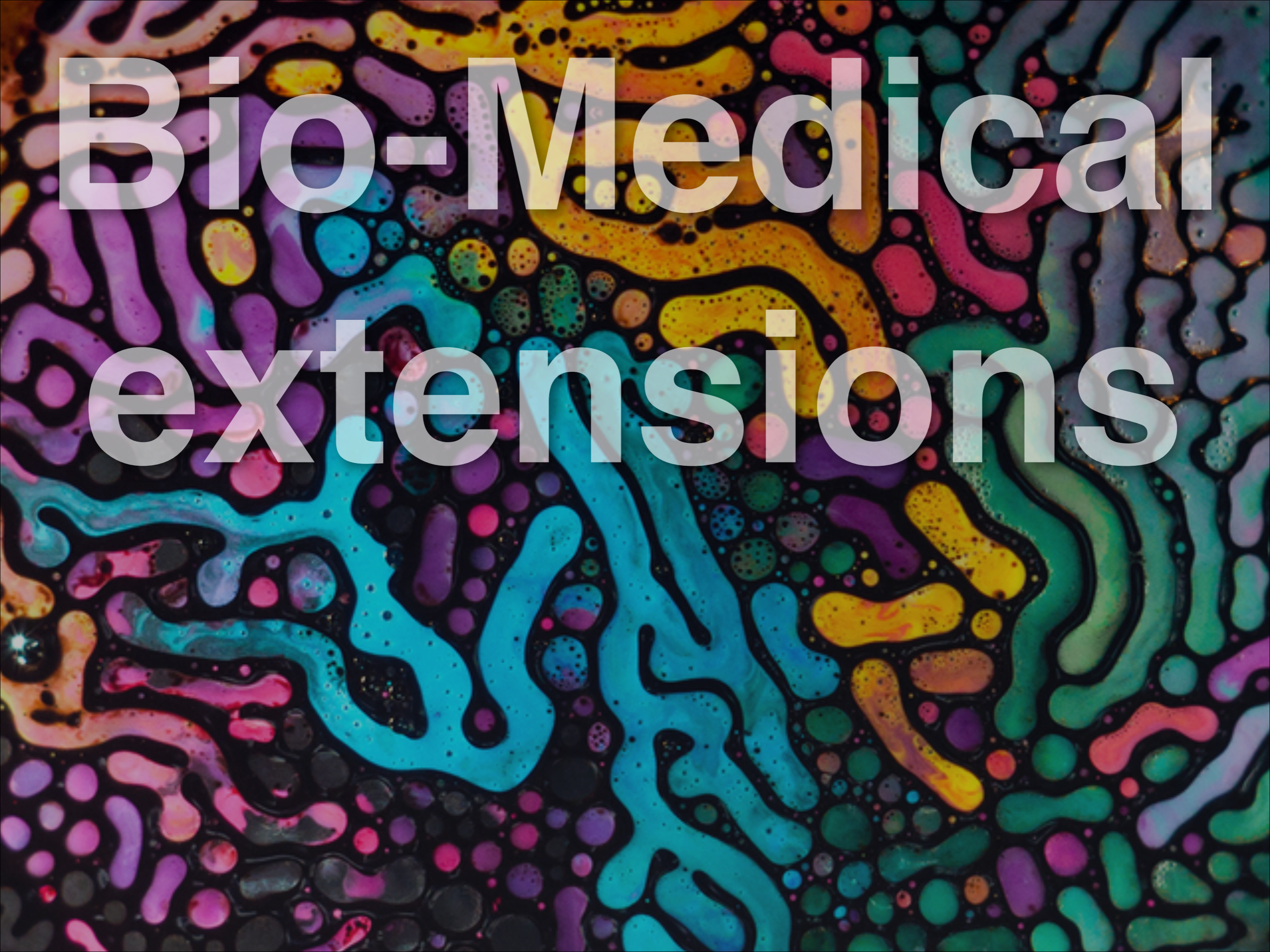
- Aggregates multi-modal index services

*ex: return images with compound X and cell density $> 40\mu\text{m}$
and nuclear texture similar to Y*

- Understands query dialects:

XQuery, SPARQL, BisQue

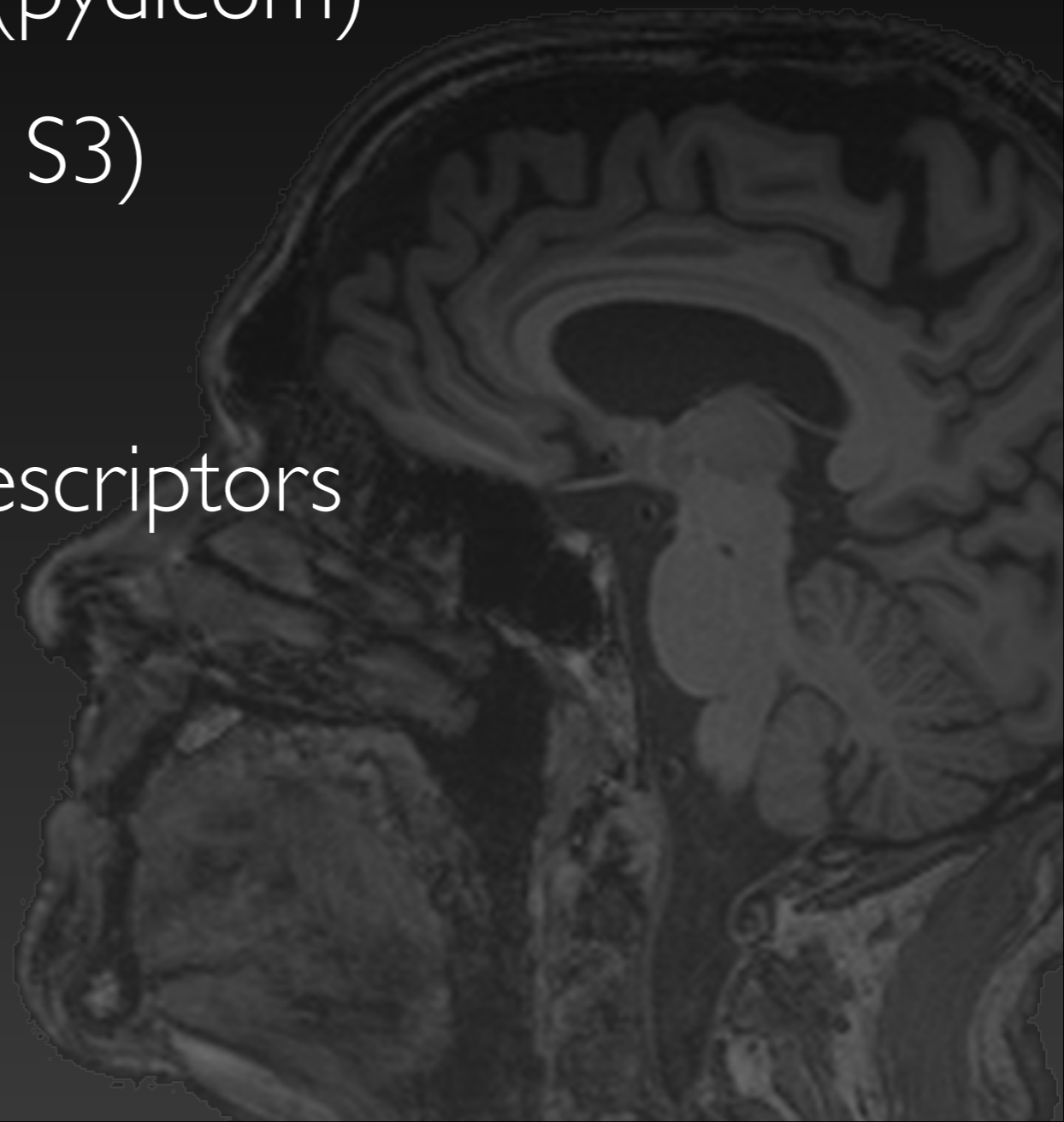




Bio-Medical extensions

Bio-medical extensions

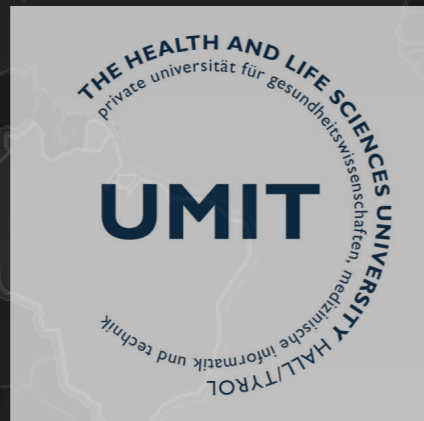
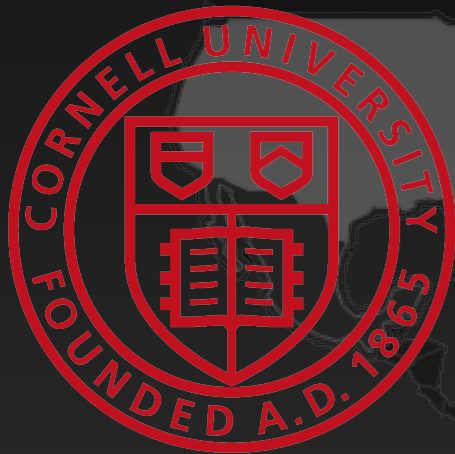
- Improved DICOM support (pydicom)
- Encrypted storage (Amazon S3)
- Graph-based queries
- Extended medical feature descriptors
- Anonymization techniques
- Specific analysis modules



1000+ Bisque users



iPlant
Collaborative™



Cold
Spring
Harbor
Laboratory

Bisque sponsors



iPlant
Collaborative™



NVIDIA®

RIGHT SCALE®
CLOUD MANAGEMENT



Hungry for Bisque

A background image showing a hand holding a small bowl, and another hand holding a small round object, possibly a fruit or vegetable, over the bowl. The image is dark and slightly blurred, with the hands and bowl being the main focus.

- Use a public Bisque:
 - iPlant: <http://iplantcollaborative.org>
 - UCSB: <http://bisque.ece.ucsb.edu>
- Get your own (source and documentation):
<http://bioimage.ucsb.edu/bisque>
- Run on Amazon AWS:
RightScale.com: “Bisque Server Template”

Get in the Kitchen

- <http://biodev.ece.ucsb.edu/projects/bisque>
 - Developer wiki and bug reports
 - Browse source
 - Get source and installation instructions
- Join our mailing list:
<http://groups.google.com/bisque-bioimage>
- Chat with the developers (Google Talk or Jabber):
bisque@partychapp.appspotchat.com