

LIMPID/BisQue

A scalable infrastructure for
reproducible, image driven data science

Principal Investigators

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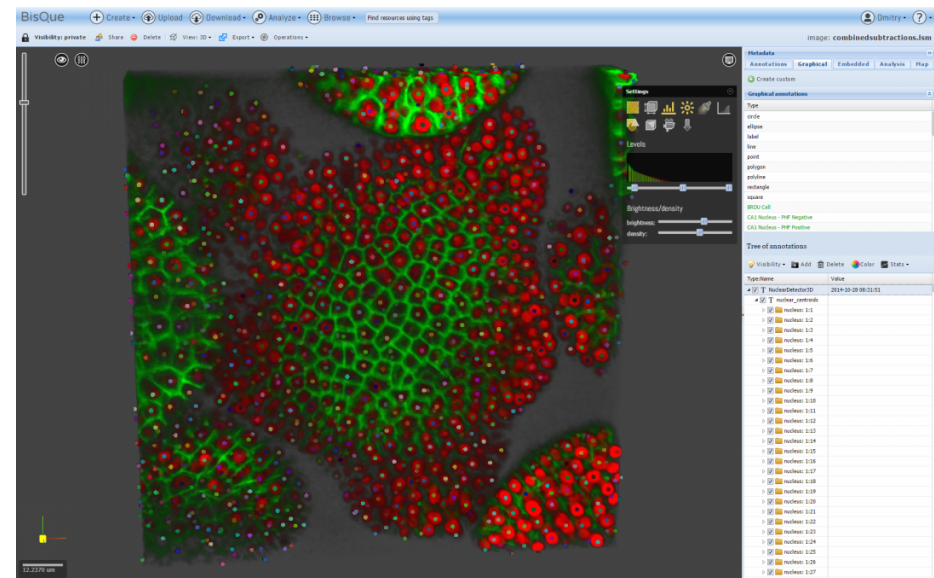
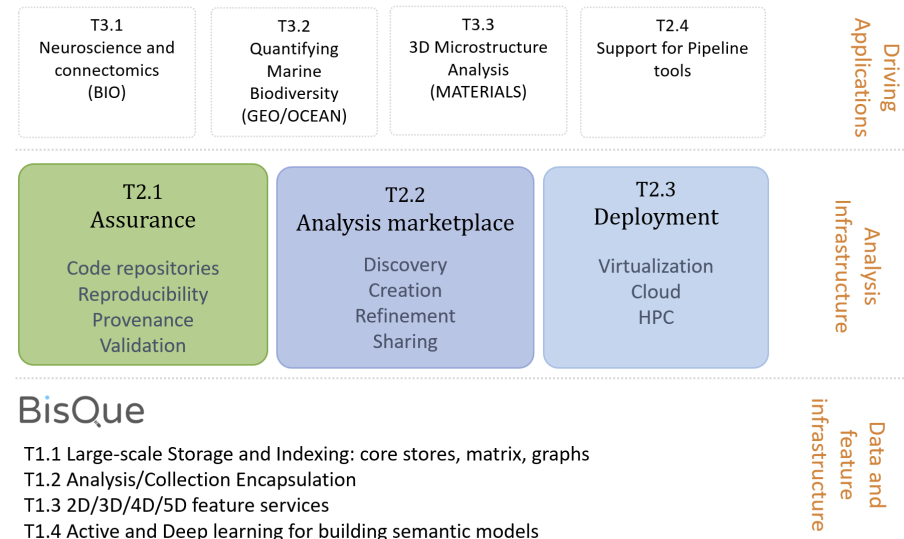
Applications

McClean Echlin, Materials
Marat Latypov, Materials
Anandasankar Ray, Neuroscience
William Smith, Neuroscience

NSF #1664172 (EAGER, 2016-18)
NSF #1650972 (SSI, 2017-22)

LIMPID/BisQue Overview

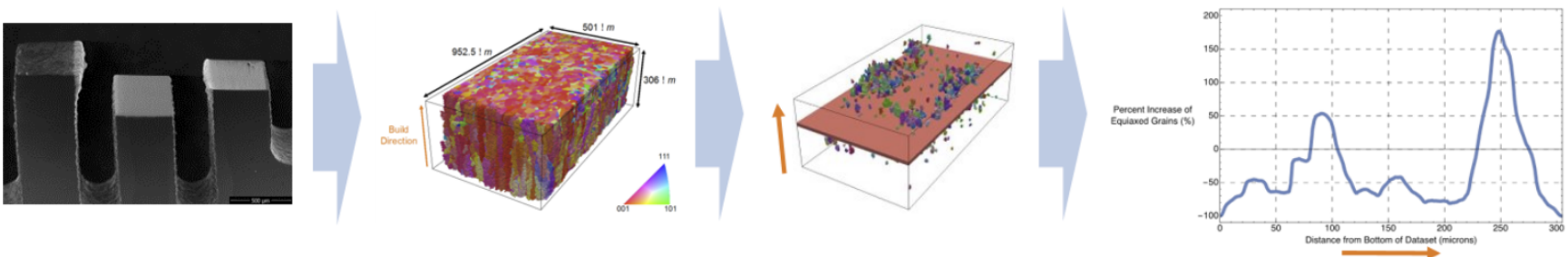
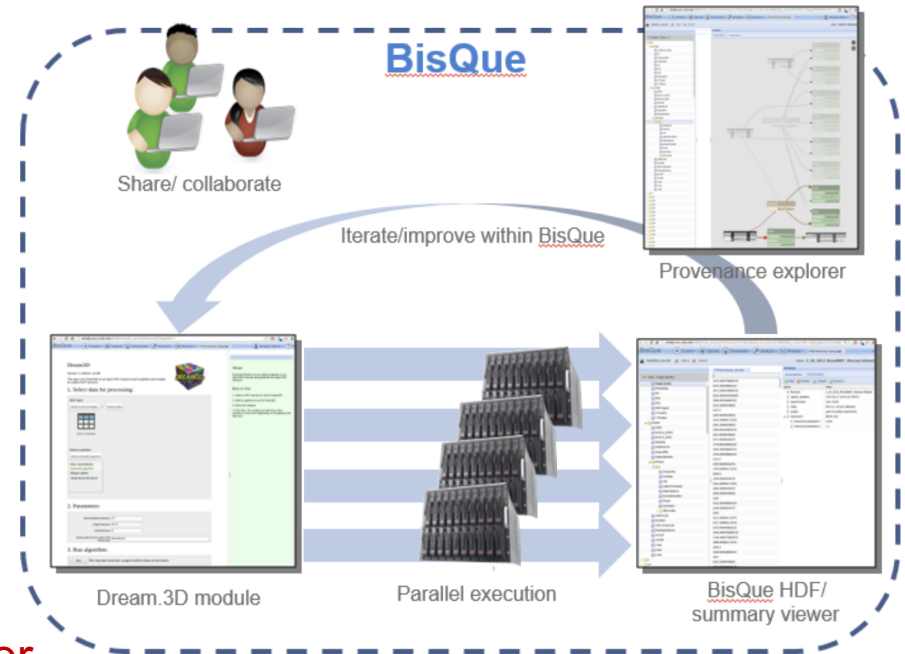
- LIMPID is built on **cloud-based analysis platform BisQue**
- **Management, analysis, and sharing** of images and metadata for large-scale data science
- **Flexible and scalable query system** across network of multimodal data items
- **Module system** for scalable integration of analysis tasks over images and metadata
- 200+ life science image and video formats
- **Analysis marketplace:** easy sharing and discovery of analysis modules
- *More information on BisQue:*
<http://bioimage.ucsb.edu/bisque>



Dream.3D Integration

(Use Case: Materials Structure Analytics)

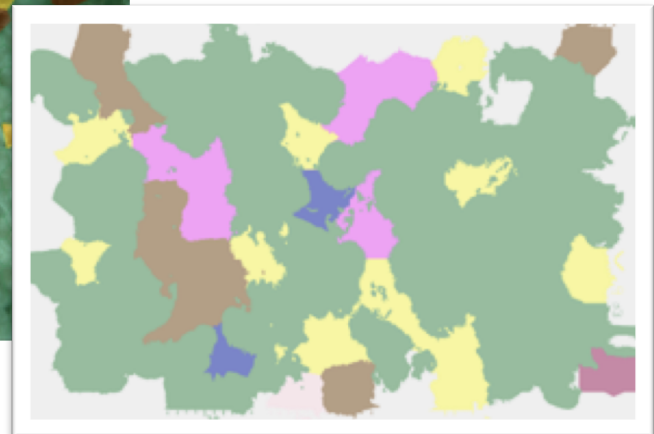
- Large acquired Materials Science datasets analyzed in desktop apps such as Dream.3D
- **Dream.3D pipelines and execution integrated in BisQue**
- Datasets and pipelines can be **shared with collaborators**
- Parameter variations are **parallelized** over compute grid
- Large datasets **viewable in web browser**
- **Provenance tracking** for improved repeatability



Deep Learning Integration

(Use Case: Quantifying Marine Biodiversity)

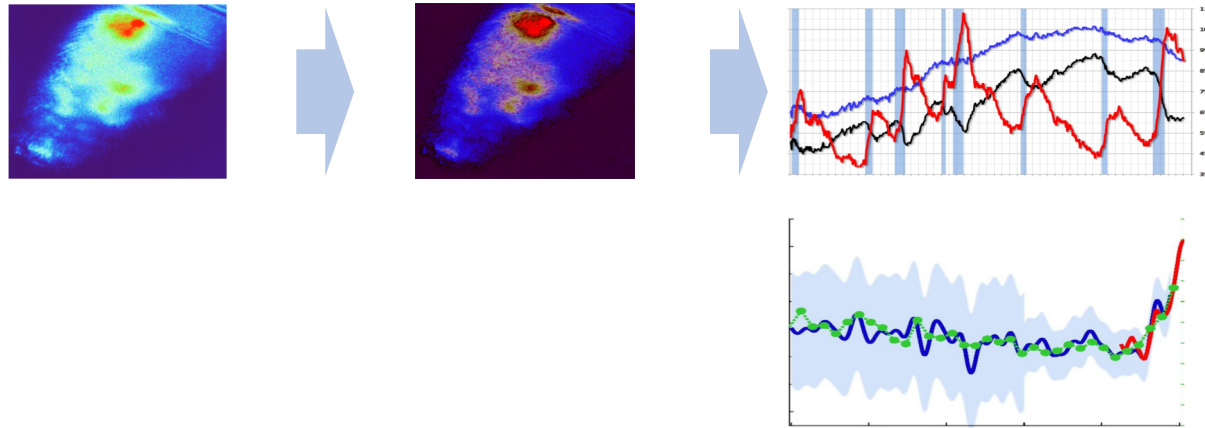
- **Goal:** Automate annotations without engineer in reduced time for specific datasets
- Metadata driven training set selection
- Automatic parallelization
- Integrated validation (model selection)
- Multiple imaging modalities: photographs, satellites, microscopes



Deep learning method development

(Use Case: live Calcium imaging in neurons)

Application: Quantify and map changes in fluorescent signal in 4D data acquired from live imaging of neuronal activity in insect nervous system



- Requires registration of images to compensate for movement.
- Automated identification of neurons from background based on shape, position and fluorescent changes.
- In a post processing phase quantify fluorescent changes over time for each neuron identified.
- Create plots of runs for evaluation as supervised parameters are altered, to create summarized data.