

# Deliverables: 18 Sep 2017

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NOMADS

(**NO**vel **M**ethod of **A**utonomous **D**etection for **S**ynapses)

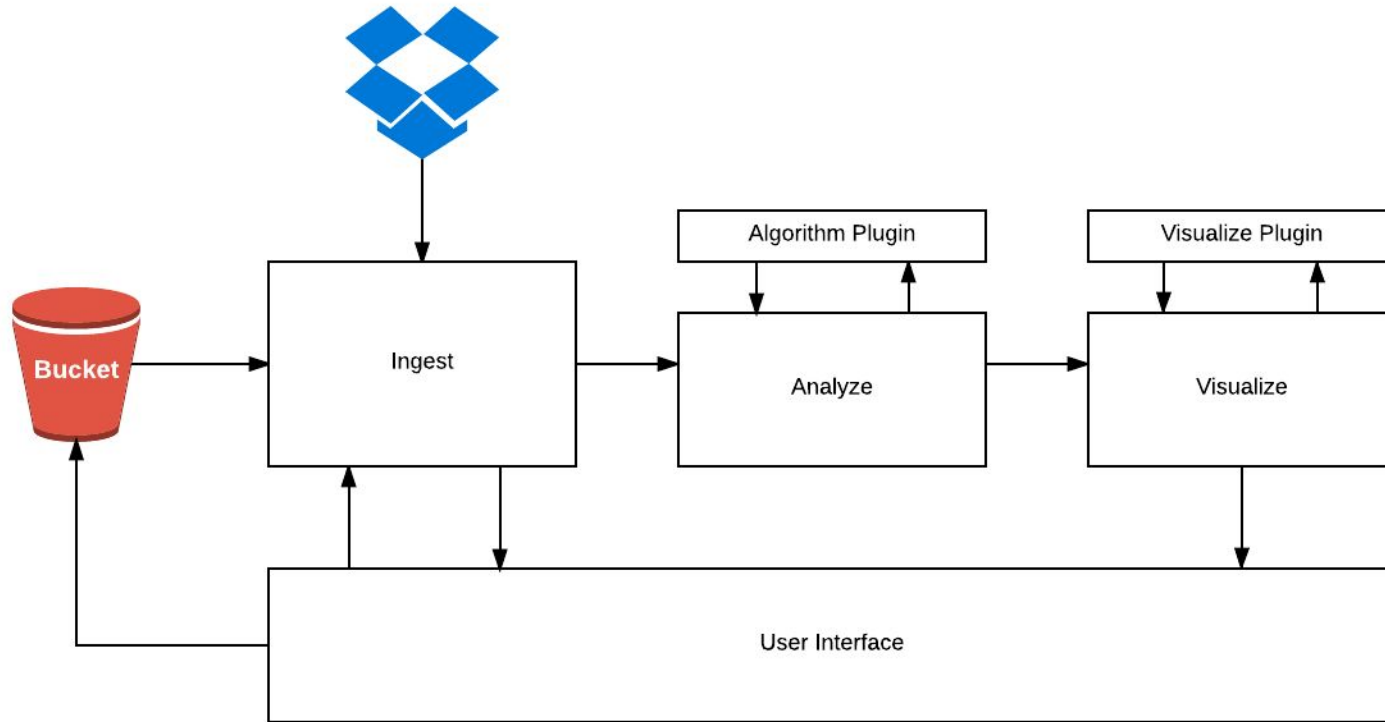
# From Last Week

- Diagram of Overall Project Vision
- Detailed Analysis of Boss Download Speed
- Biomarker Glossary
- Data Sanity Check (Reach)
- Collman Algorithm Implementation (Reach)
- For Sharmini - Enroll in NDDI

# From Last Week

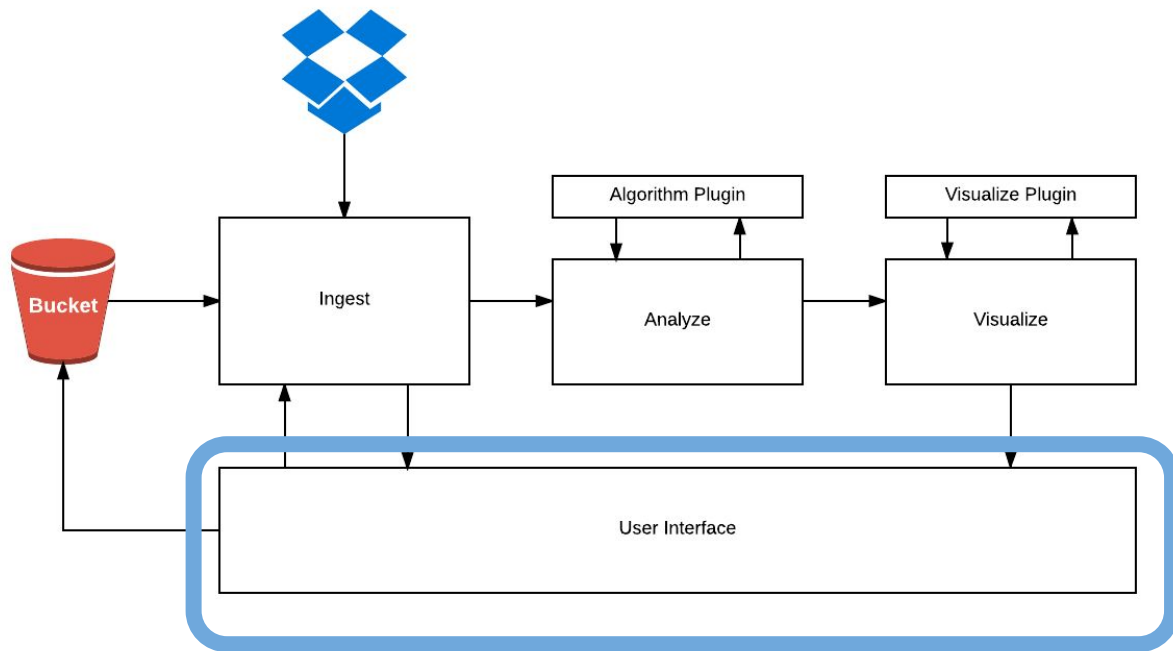
- Diagram of Overall Project Vision ✓
- Detailed Analysis of Boss Download Speed ✓
- Biomarker Glossary ½
- Data Sanity Check (Reach) ½
- Collman Algorithm Implementation (Reach) ✓
- For Sharmini - Enroll in NDDI ½

# Project Vision



# User Interface

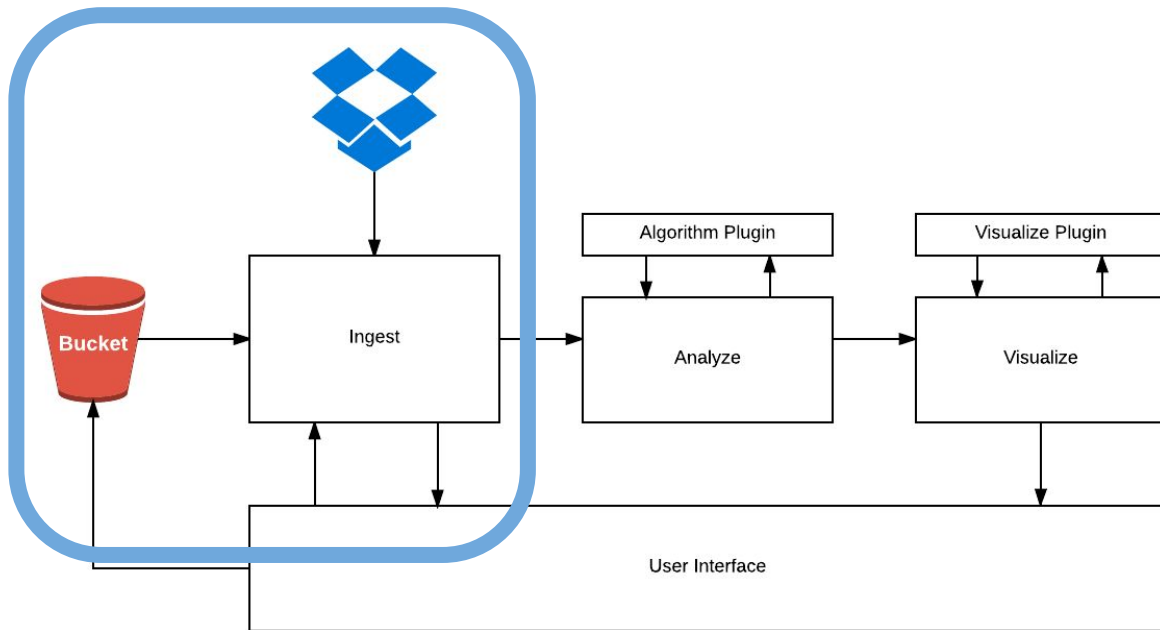
- Web U.I.
- Form Based
- Large Data Ingest
  - One Click
  - S3 Bucket + Upload
- Pipeline Options
  - What algorithm?
  - What visualizations?



# Ingest

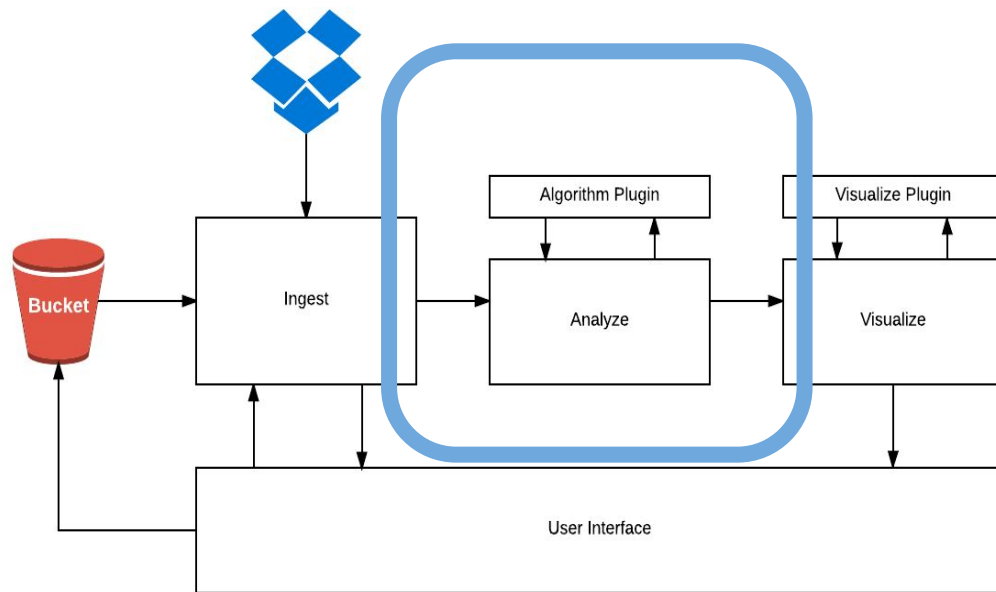
- Based on User Input

- S3 Bucket + Upload
- Dropbox
- Boss/NDStore?



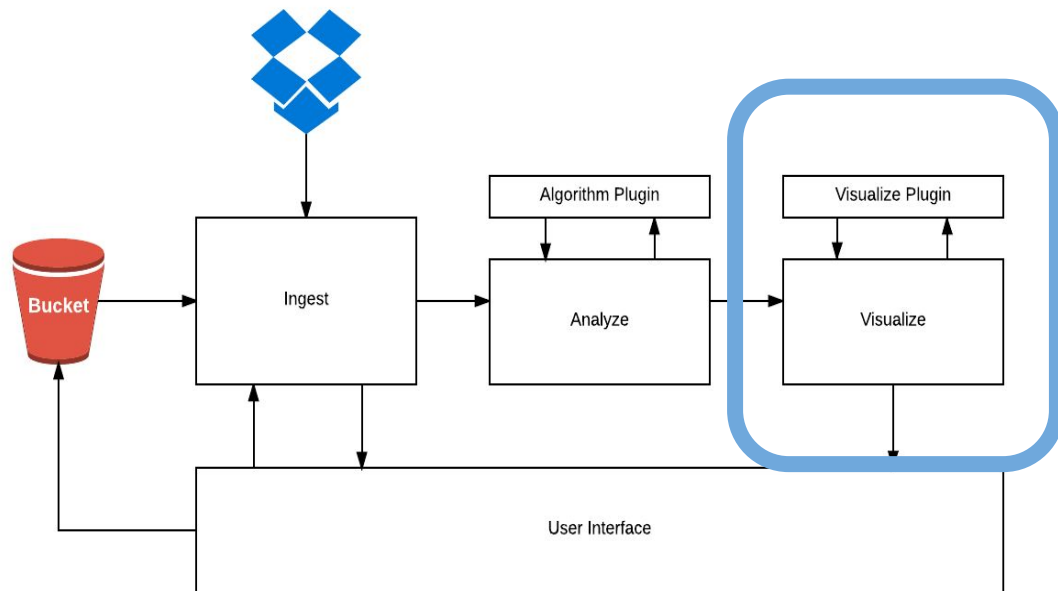
# Analyze

- **Biomarker Clustering**
  - Given bright spot in EM, what biomarkers are in AT?
  - Attempt to characterize synapse types
- **Detection**
  - Given AT/EM, where are the synapses at?
- **Hotswapping**
  - Standard way to swap in & compare other algorithms



# Visualize

- NDWebTools-esque result viewer
- Downloadable Raw Outputs
  - Work with FIJI
  - User Specified in form
- Charts
  - “Hotswappable” as visualize plugins
  - T-SNE for clusters





# Boss Download Speeds

- Fastest way to download an entire would be to download using 2048x2048x16 chunks - would take about ~36 secs.
- Downloading an entire channel at once takes about ~54 secs.

[https://nbviewer.jupyter.org/github/NeuroDataDesign/nomads-f17s18/blob/master/source/jlc/Boss\\_Chunk\\_Analysis/Boss\\_Chunk\\_Analysis.ipynb](https://nbviewer.jupyter.org/github/NeuroDataDesign/nomads-f17s18/blob/master/source/jlc/Boss_Chunk_Analysis/Boss_Chunk_Analysis.ipynb)

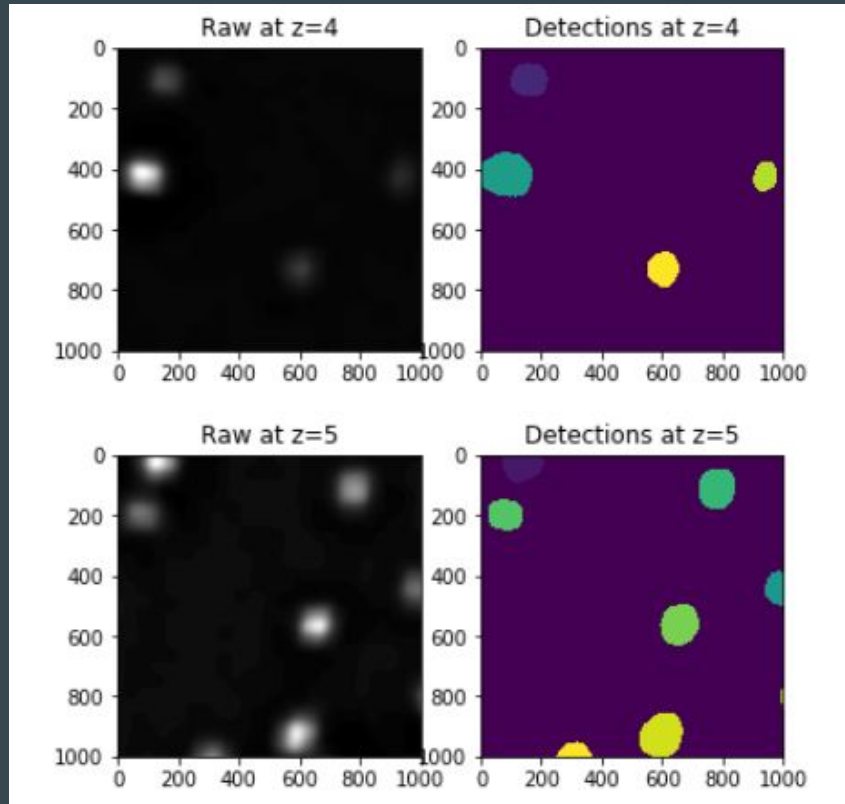
# Biomarker Glossary

- Locations of biomarkers in Collman paper researched- separated into pre/postsynaptic and excitatory/inhibitory synapses
- Many unanswered questions- work will continue next week

<https://github.com/NeuroDataDesign/nomads-f17s18/blob/master/source/dmannan/biomarkers/Colleman%20write%20up.docx>

# Collman Alg

- [https://github.com/NeuroDataDesign/nomads-f17s18/blob/master/source/bstadt/Collman/Collman\\_Algorithm.ipynb](https://github.com/NeuroDataDesign/nomads-f17s18/blob/master/source/bstadt/Collman/Collman_Algorithm.ipynb)



# For Sharmini - Enroll in NDDI

- Enrollment already confirmed with Dr. Vogelstein - just need to make it official with JHU registration

----- Forwarded message -----

From: **Cathy Jancuk** <[cjancuk@jhu.edu](mailto:cjancuk@jhu.edu)>

Date: Fri, Sep 15, 2017 at 4:41 PM

Subject: Active Approval for your course got turned off at 4:30

To: jovo Vogelstein <[jovo@jhu.edu](mailto:jovo@jhu.edu)>

That is the deadline of electronic adds at the 2-week mark. Tell the student to show up with an add/drop form in my office on Monday. She'll have to get a couple of signatures so she can add this late.

Cathy

# For Next Week

- Learn to read EM data- identify cell structures without relying on biomarkers
- Complete data-sanity check
- Check to see if biomarkers' location dependent on regions of the brain
- Collman SVM Implementation
- Collman Basic and Collman SVM Histogram Stats
- Clustering & T-SNE on Collman Synapses (Reach)