An Automated Images to Graphs Pipeline for High Resolution Connectomics

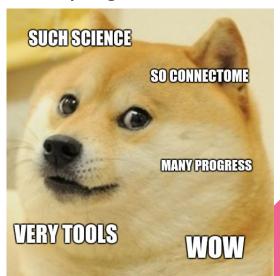
Presented by Brandon Duderstadt

Summary

- Connectomics is a field where research tools are diverse and difficult to interface with each other
 Creation of a pipeline to synchronize tools would be quite helpful
- Additionally, these tools may perform well in isolation, but may perform poorly in conjunction with other tools
 Coordination tools and tune algorithms for best overall performance
- Error metrics for these tools are only interpretable for the individual pieces Implementation of an interpretable pipeline-wide error metric

Opportunity

- Look at all of these great tools in the exciting new field of connectomics!
- I bet we could make a bunch of scientific progress with these.
- How much progress?
- So much progress!



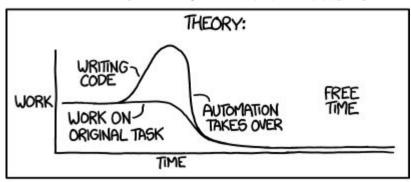
Challenge

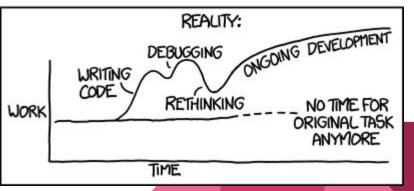
- It's a shame none of these tools work together. That would be so convenient.
 - Imagine trying to build a doghouse without blueprints...
 - With only a hammer and screws...
 - And with wood of all different shapes and sizes...
 - With no saw.

Action

- Create a pipeline to automate the process
 - Fit all of the pieces together
 - Standardize output and process
- But that's too easy
 - Include variable parameter sets
 - Calculate appx. error in final product
 - Distribute computations for faster results
 - Create custom synapse detector

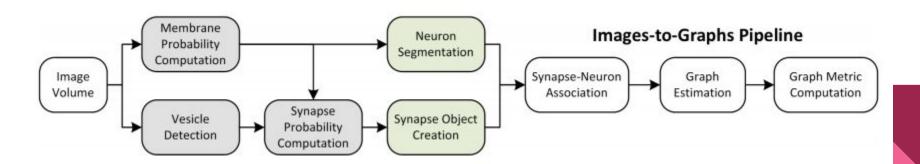
"I SPEND A LOT OF TIME ON THIS TASK. I SHOULD WRITE A PROGRAM AUTOMATING IT!"





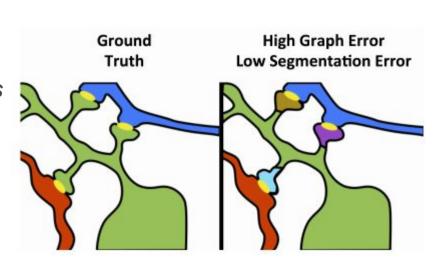
Pipeline Overview

- Begin with cortical tissue image volume
- Membrane Detection -> Neuron Segmentation -> Segment Assembly
- Grey items are computed once
- Green items are parameter based
- White items are computed with every run



Error Measurement

- Effective parameters require a fitness
- This is tricky due to tool diversity
- "As we demonstrate, the optimal results for a subtask may not translate to optimal results for the overall pipeline."
- Topographical vs Segmentation Errors
- Parameter Tuning



Resolution & Future Work

- Pipeline is created and open for use!
- Graph error is a difficult issue, but parameter tuning allows error minimization
- Graphs with some error can still be useful
- Scaffolding for algorithm error optimization is now in place
- Speed & efficiency increase

Discussion

PROS

- Automation
- Standardization
- Error Metric
- Applicable to future work

CONS

- Advances in process may render pipeline obsolete
- Questionable ability for pipeline evolution
- Synapse occurrence may not always imply connectivity

QUESTIONS?

As a total expert on connectomics with TWO (2) whole classes of experience and training. I will now attempt to answer your difficult and insightful questions.

MY HOBBY:
SITTING DOWN WITH GRAD STUDENTS AND TIMING
HOW LONG IT TAKES THEM TO FIGURE OUT THAT
I'M NOT ACTUALLY AN EXPERT IN THEIR FIELD.

