### NOMADS

•••

Neurodata's Opensource Method for Autonomous Detection of Synapses

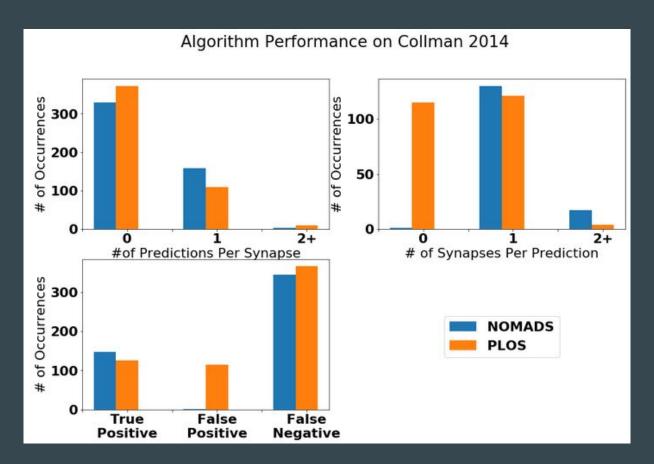
## LAST WEEK

- PLoS vs NOMADS on Collman14
  - DoD: Collman Plot w/ Comparison
- NOMADS on super new Collman Data
  - o DoD: Collman Plot w/ Comparison
- Just look at it"<sup>TM</sup> of False Negatives
  - DoD: Notebook with the analysis
- Synaptome Code for calculating features
  - DoD: Notebook demonstrating usage

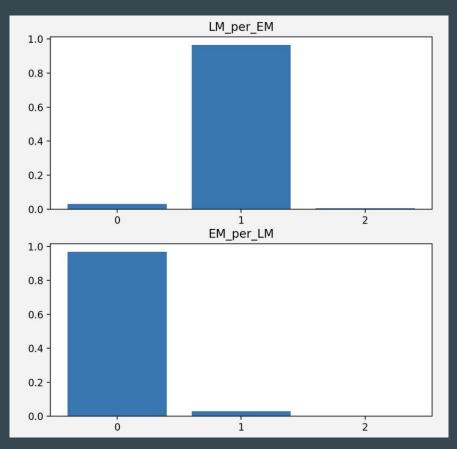
## LAST WEEK

- PLoS vs NOMADS on Collman14
  - o DoD: Collman Plot w/ Comparison
- NOMADS on super new Collman Data ☑
  - o DoD: Collman Plot w/ Comparison
- Just look at it"™ of False Negatives ✓
  - o DoD: Notebook with the analysis
- Synaptome Code for calculating features \( \mathbb{Z} \)
  - DoD: Notebook demonstrating usage

#### NOMADS Bests PLoS on Collman 14



#### NOMADS Merges Everything on *Super New* Collman

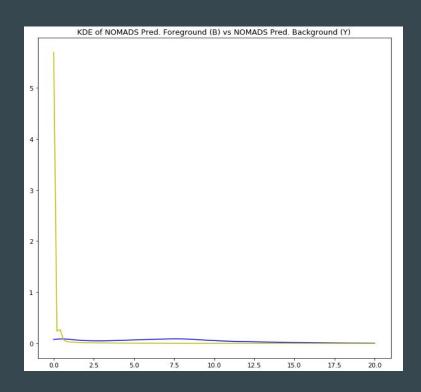


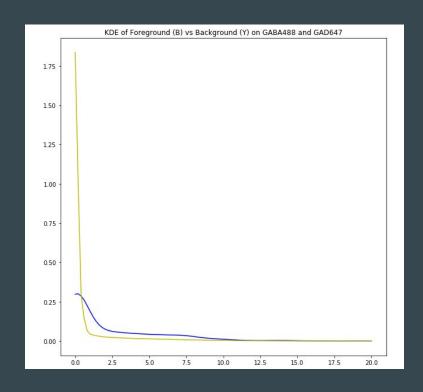
- This plot is confusing!
- LM per EM = prediction per synapse
- EM per LM = synapse per prediction

- There is ONE merged synapse in the 2 column of the EM per LM chart
- This merge accounts for ALL of the 1 LM per EM



#### GABAergic Synapses are Hard

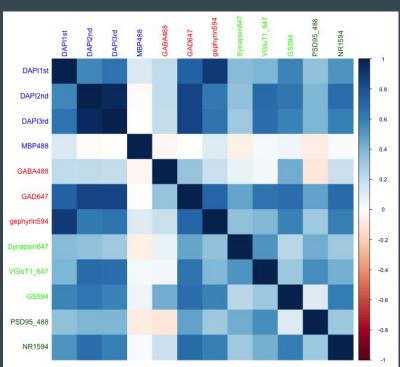




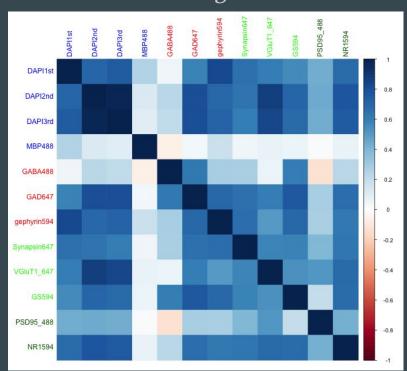
# Unsupervised Algo-Stuff around TP and FN look similar for Collman15

LINK

True Positives



False Negatives



#### Synaptome - Calculating Features

- Collman15v2
  - o 236 synapses + 12 channels
  - $\circ$  ~5 minutes to finish
- Collman14v2
  - 1451 synapses + 11 channels
  - ~20 minutes to finish
- Only integrated sum for now.
- <u>Link</u>

## **NEXT WEEK**

- Refactor Sprint Goals
- Talk With Jovo
- Implement MEDA and synaptograms
  - Port XYZ Spacial Cluster Plot to Python