Miniscope Analysis

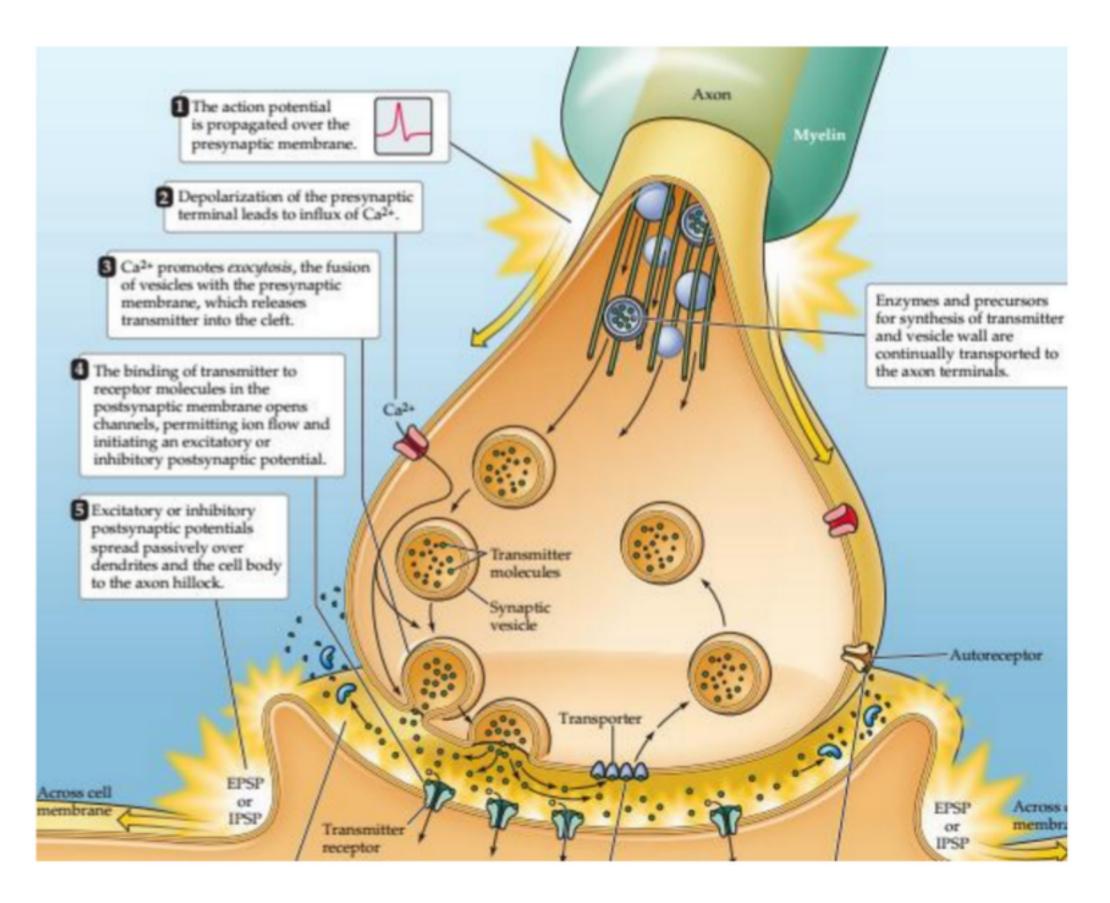
Wilke Lab Mtg.

Outline

- Motivation/Background
- Setup/Procedure
- Analysis
- Next steps

Motivation/Background

- ACC necessary for Effort-Based Decision-Making (EBD)
- Decision -> ACC activity
 -> Action Potentials ->
 Ca2+ ion influx



When the action potential arrives at the end of the axon terminal, voltage-gated calcium channels open. Entry of calcium triggers a biochemical cascade leading to fusion of neurotransmitter-containing vesicles.

Release of neurotransmitter from the pre-synaptic cell completes the goal of the action potential!

Background

- GCaMP7f binds to calcium and fluoresces green
- brightness of neurons ~ analog neural activity signal

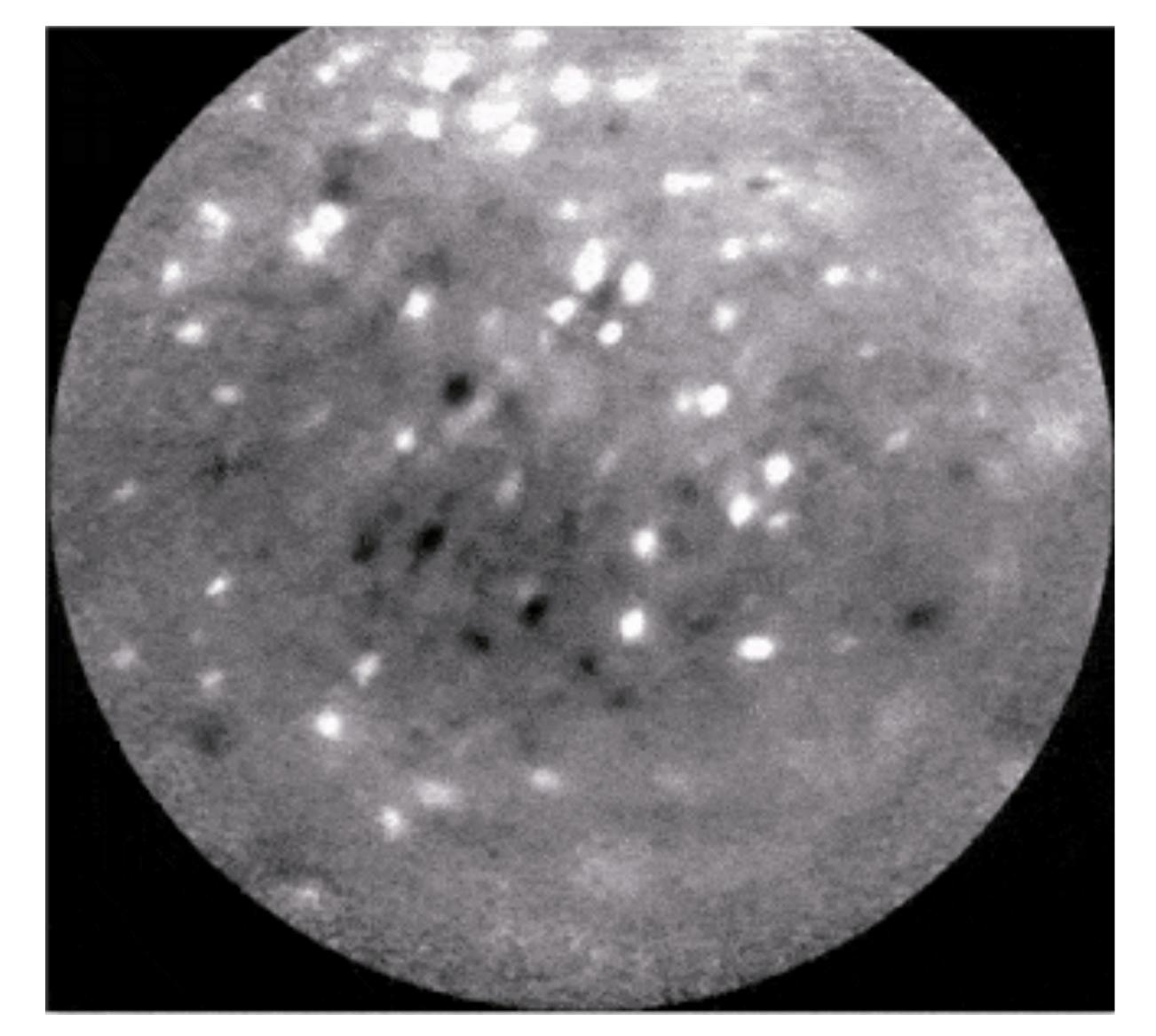
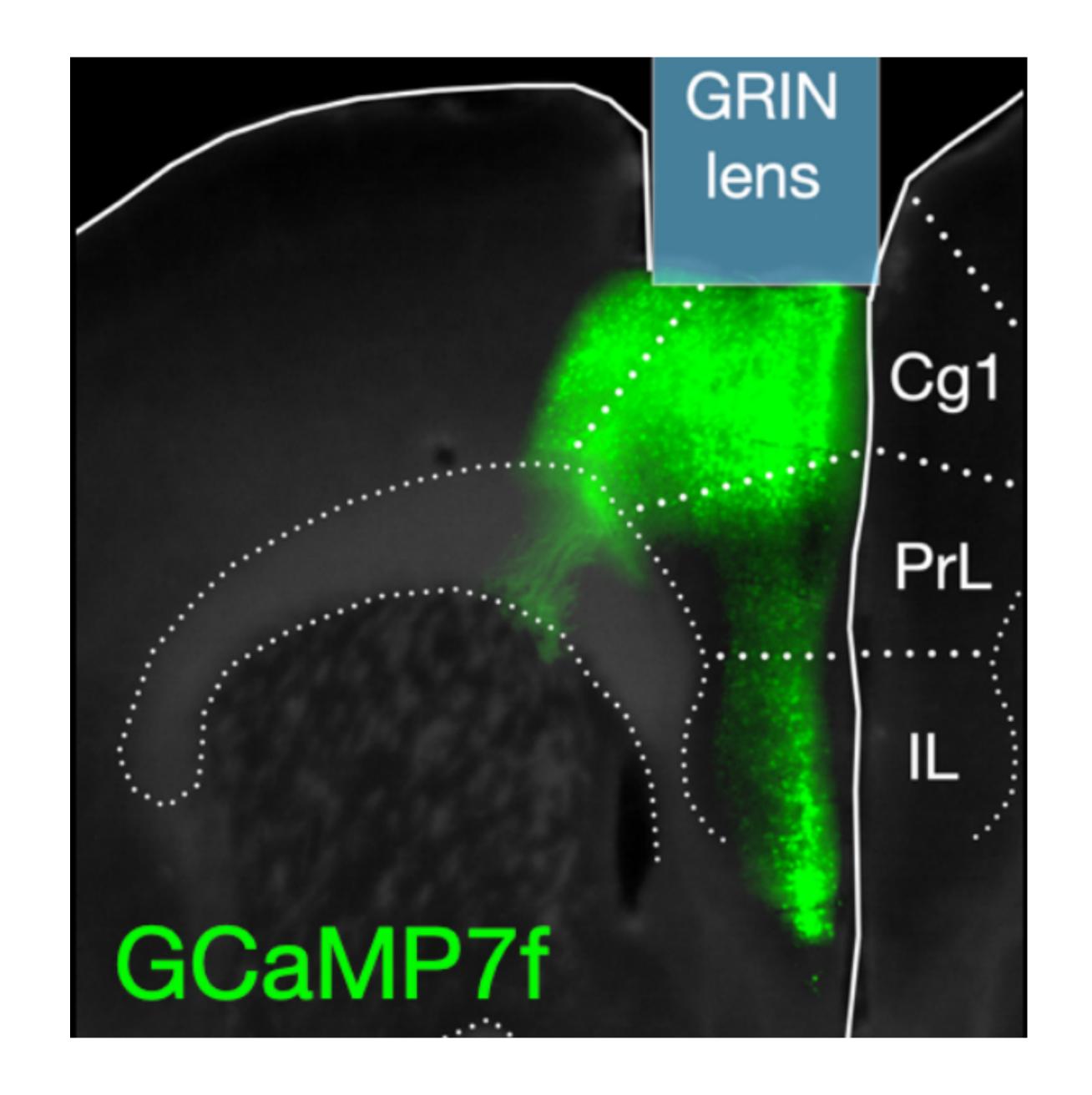


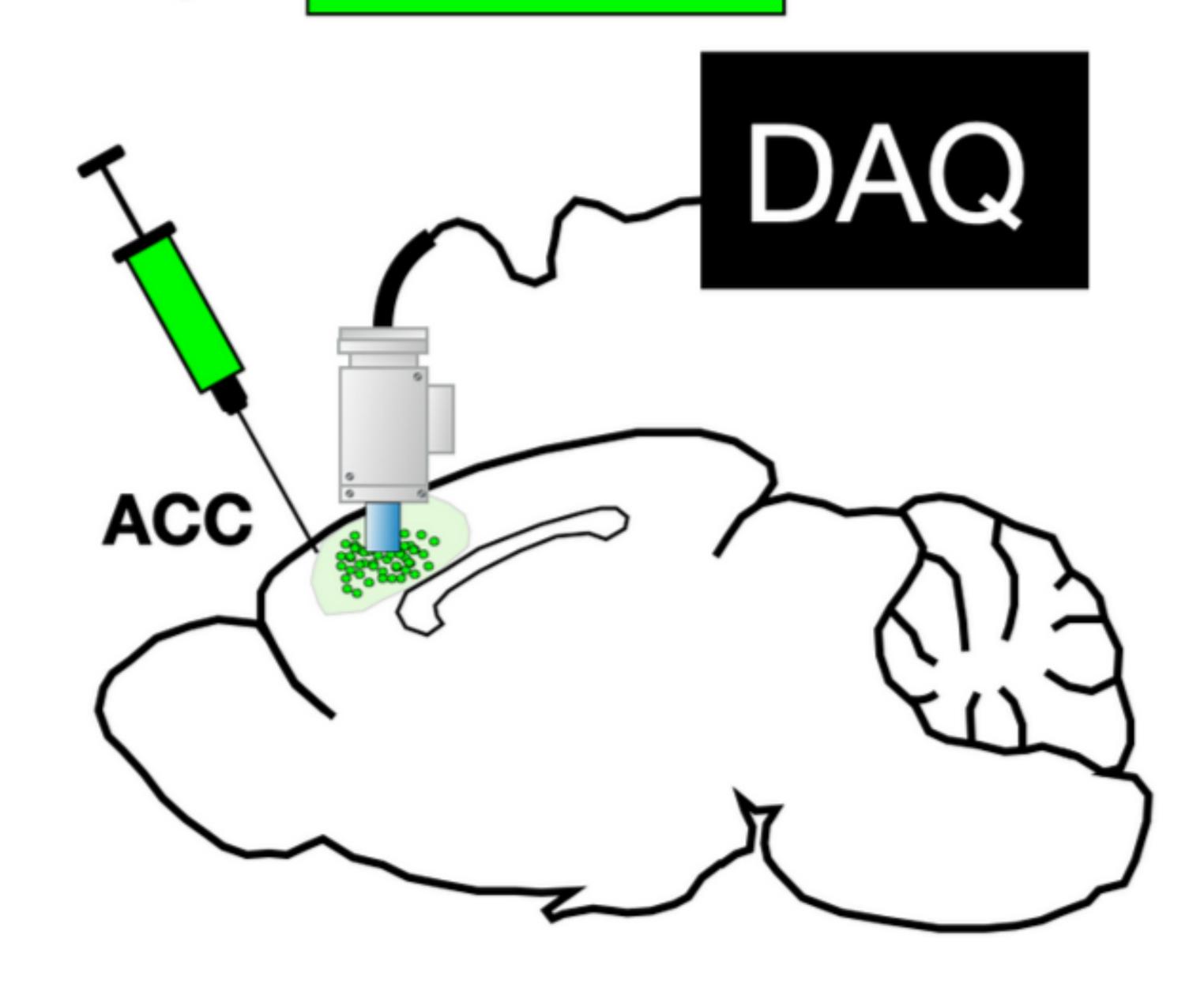
Photo Credit: Daniel Aharoni, Miniscope-v4 wiki

Setup

- GCaMP7f injection into ROI (ACC)
- 2. Gradient Index (GRIN) lens implanted over injection site
- 3. Inscopix Miniscope mounted over lens



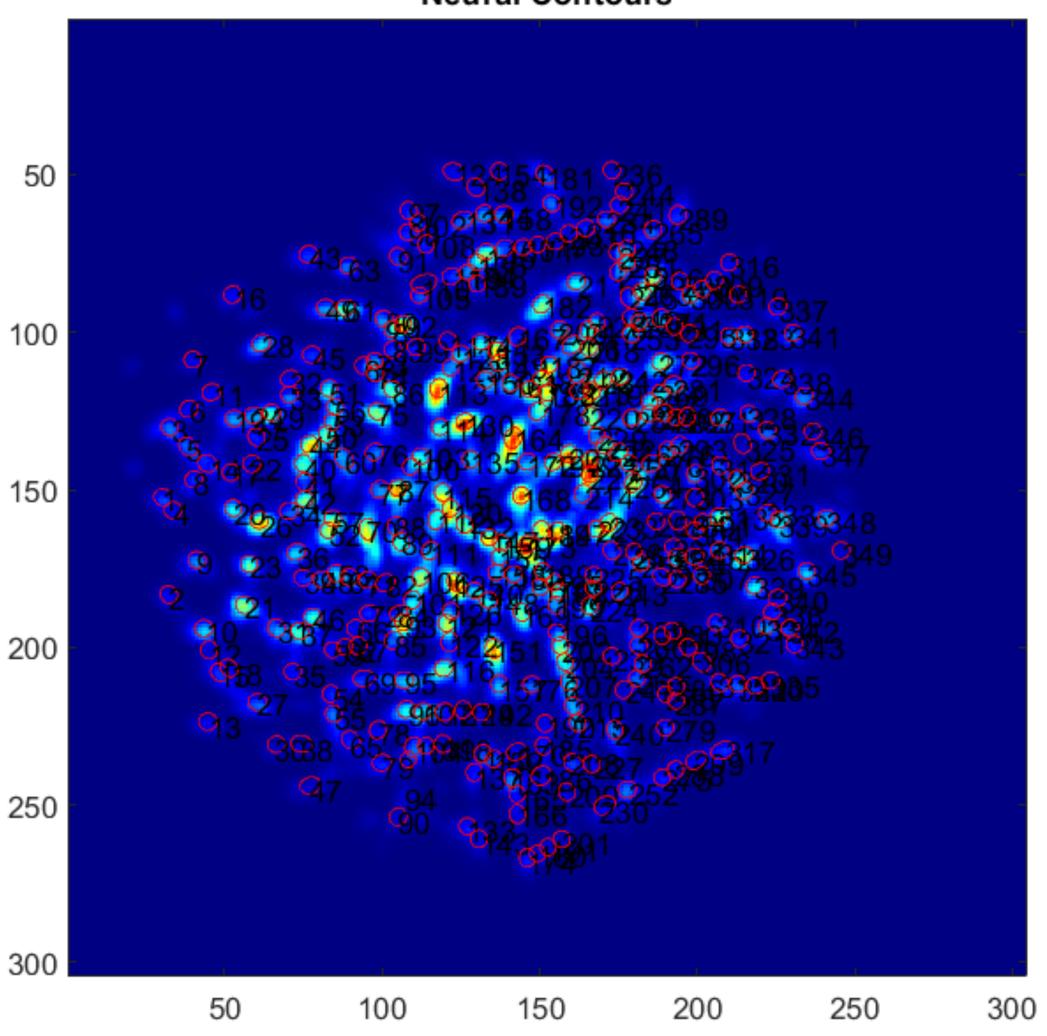
AAV-Syn-GCaMP7f

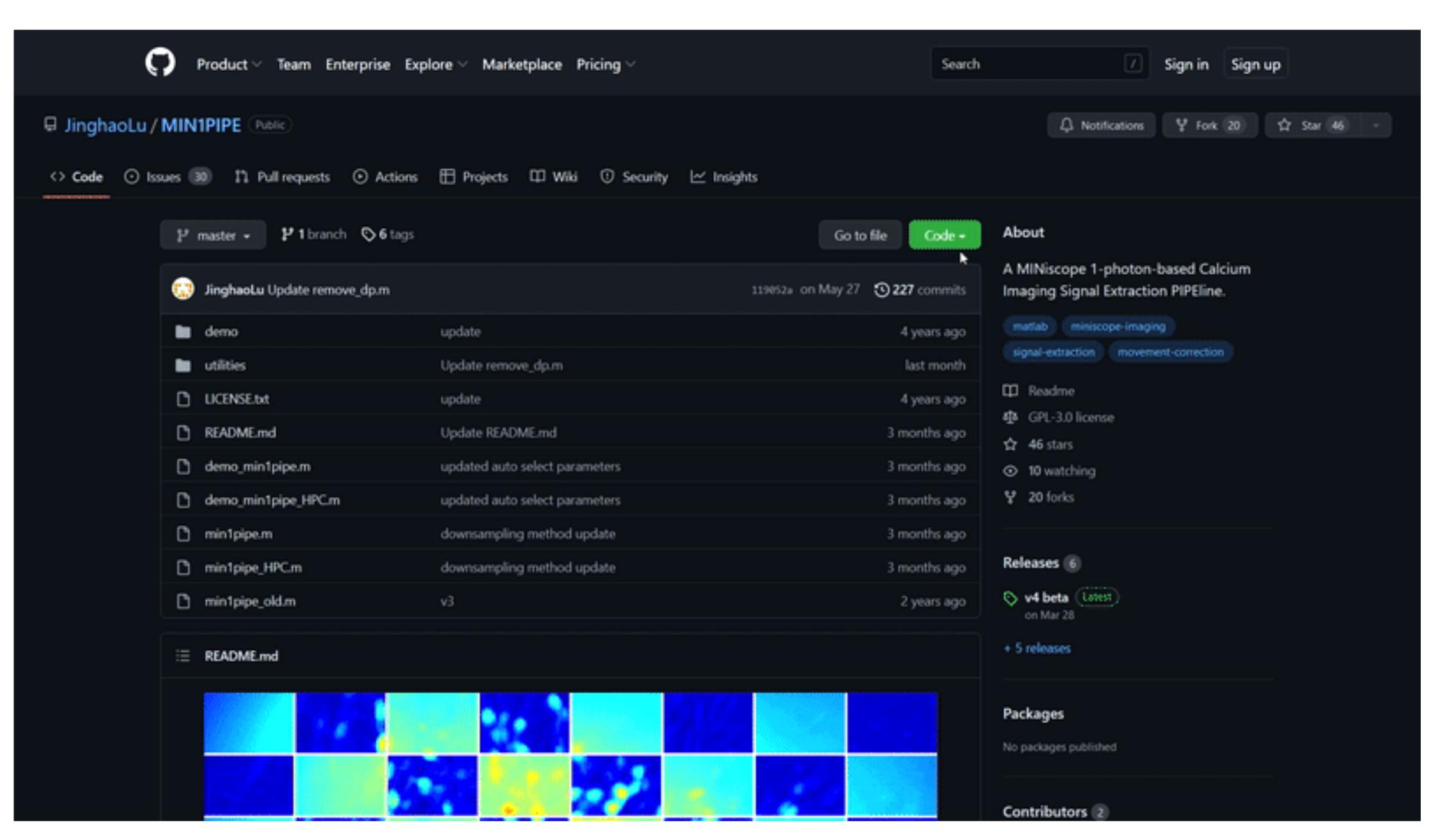


Procedure

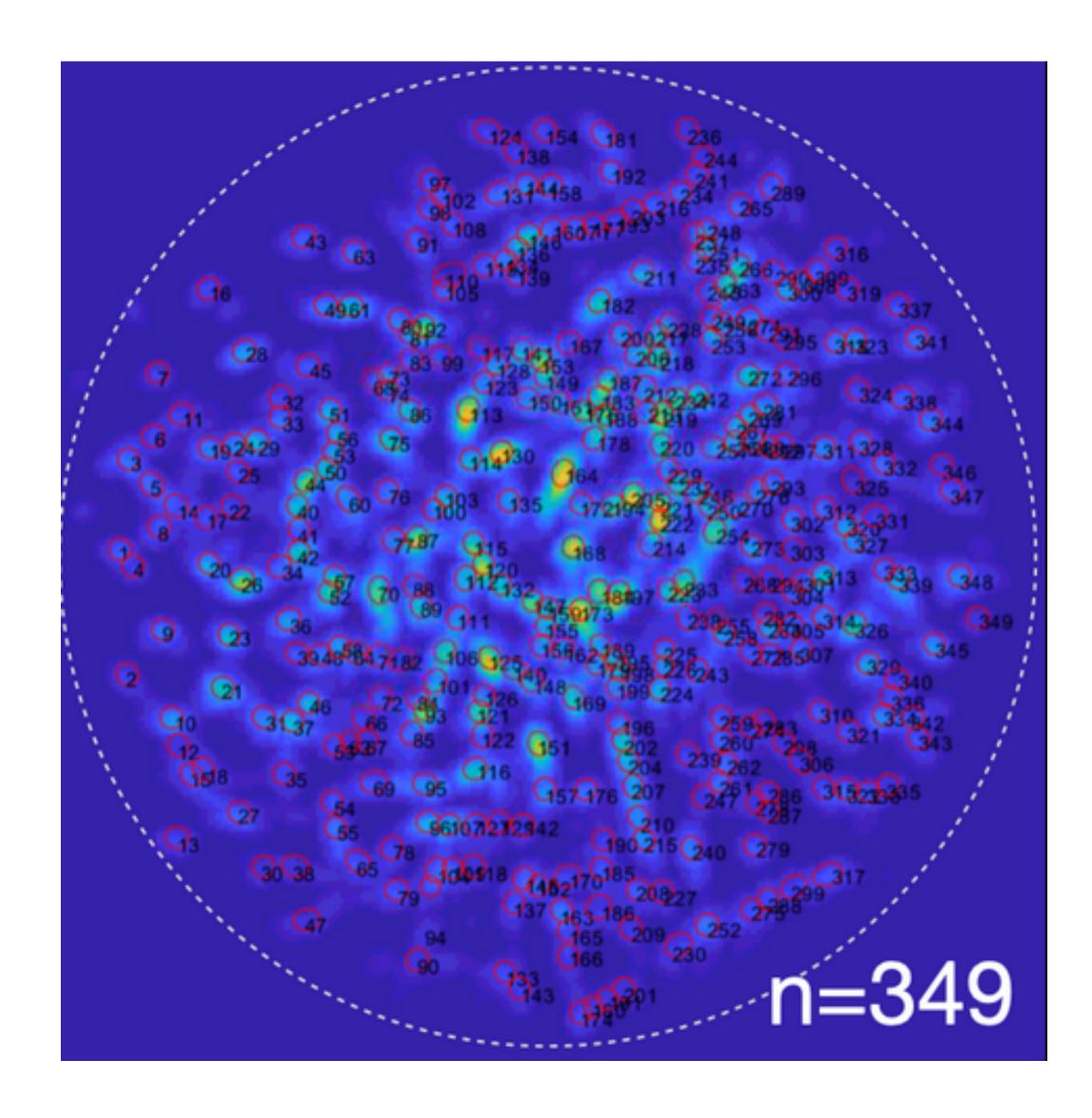
- Run Miniscope mice through maze
- Run video through MIN1PIPE

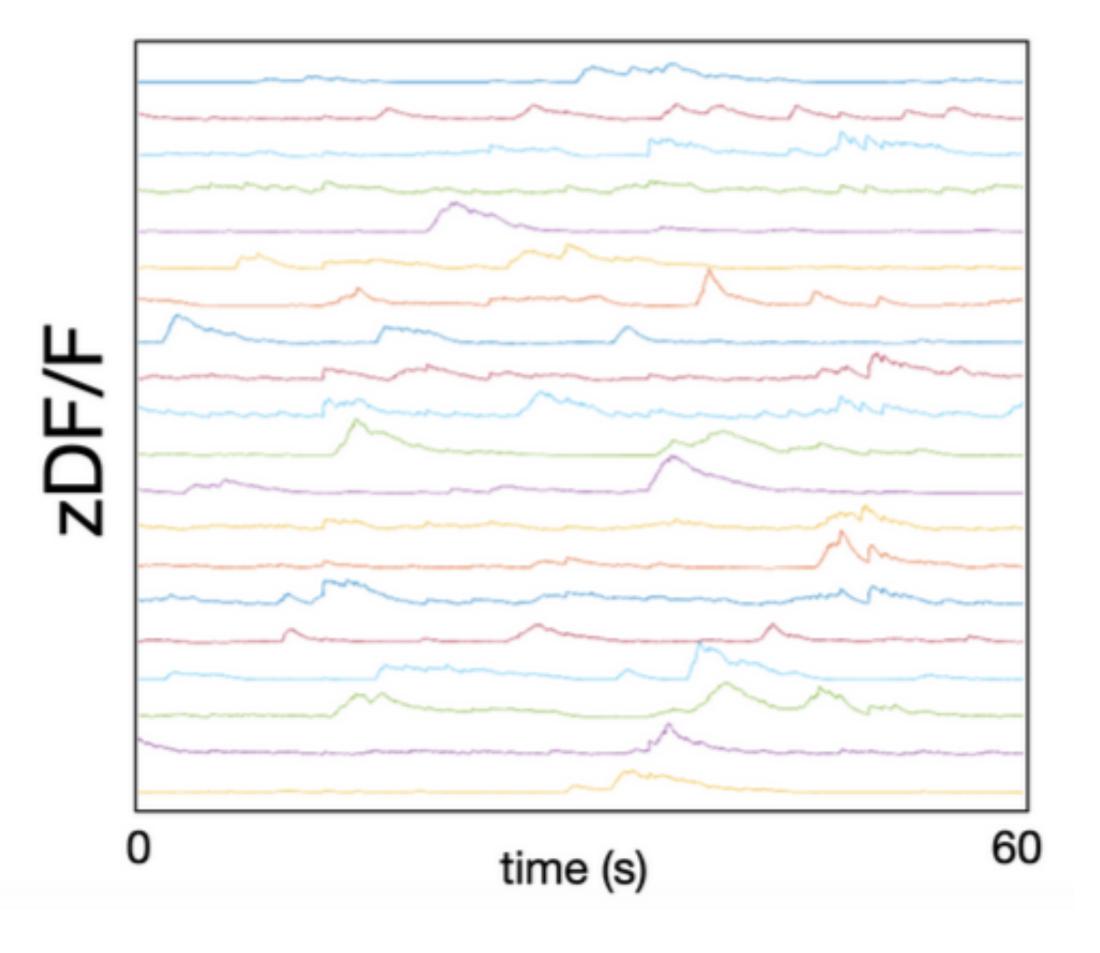
Neural Contours





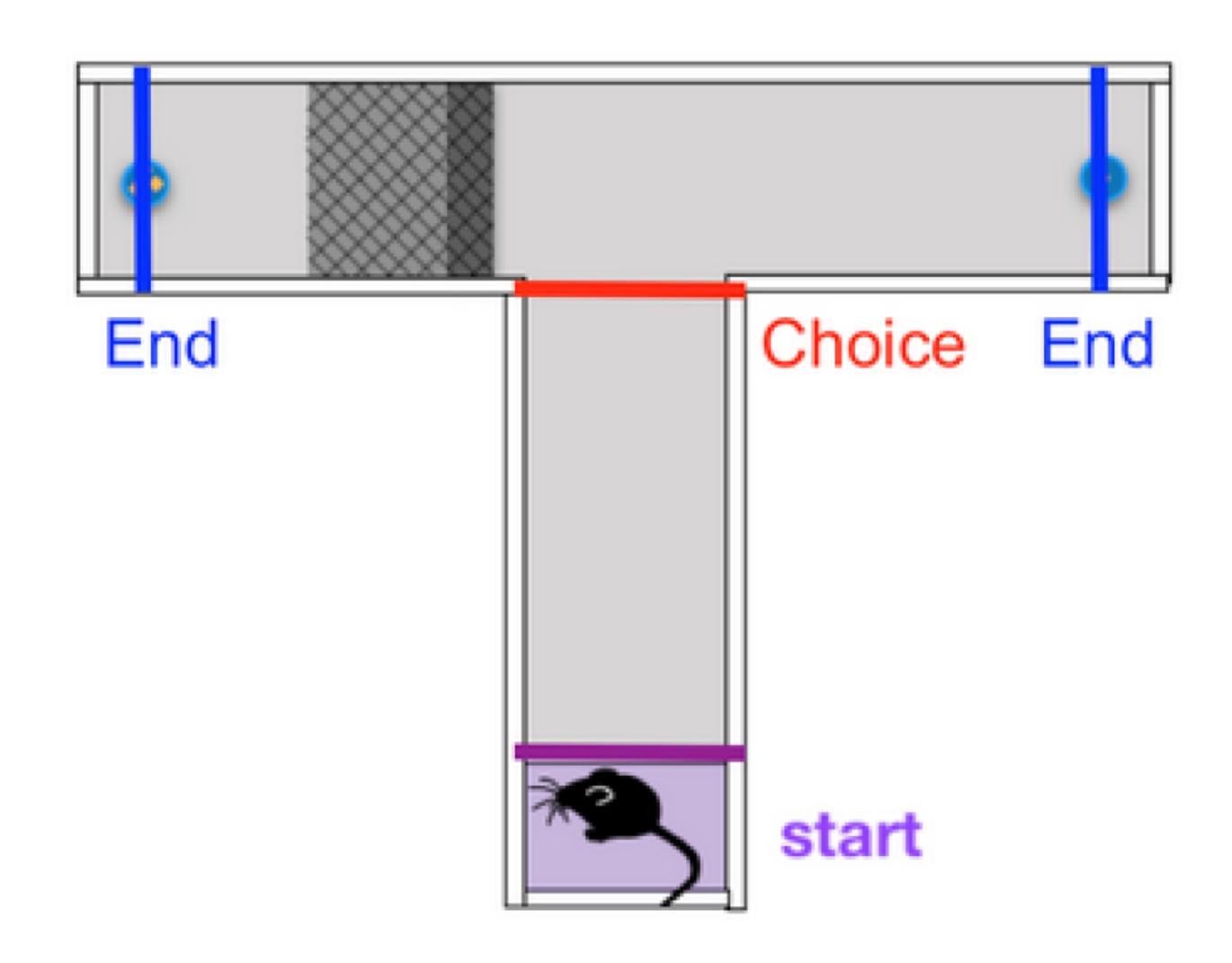
https://github.com/JinghaoLu/MIN1PIPE





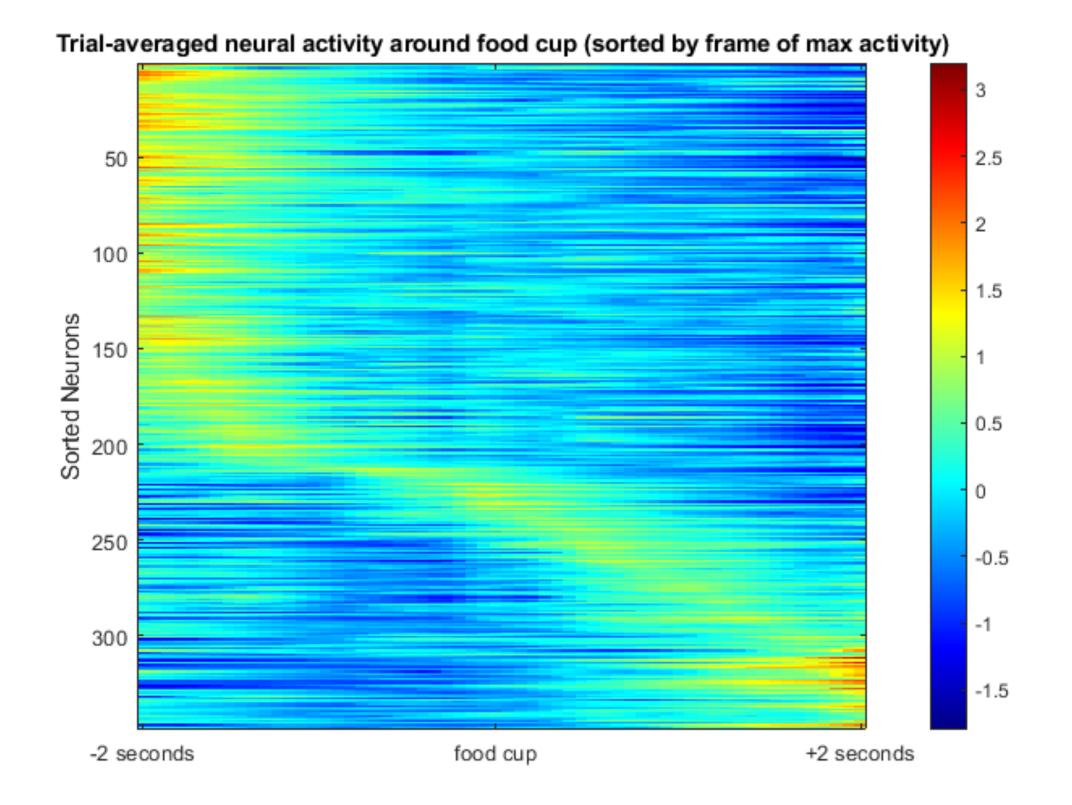
Analysis

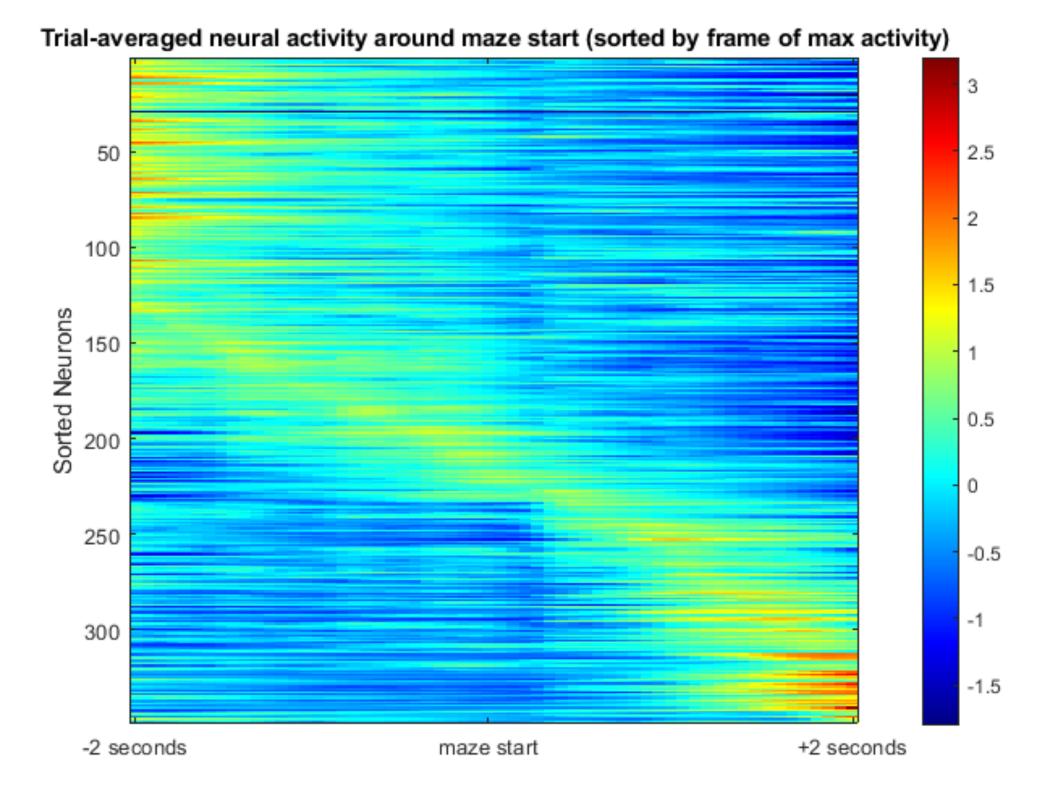
- 1. Synchronize overhead video with Miniscope video
- 2. Find frames for events of interest (i.e. start, choice, and end of each trial) and partition signals into individual trials
- 3. Center each trial around event of interest
- 4. Normalize signal, average across trials, and sort neurons by frame of maximum activity



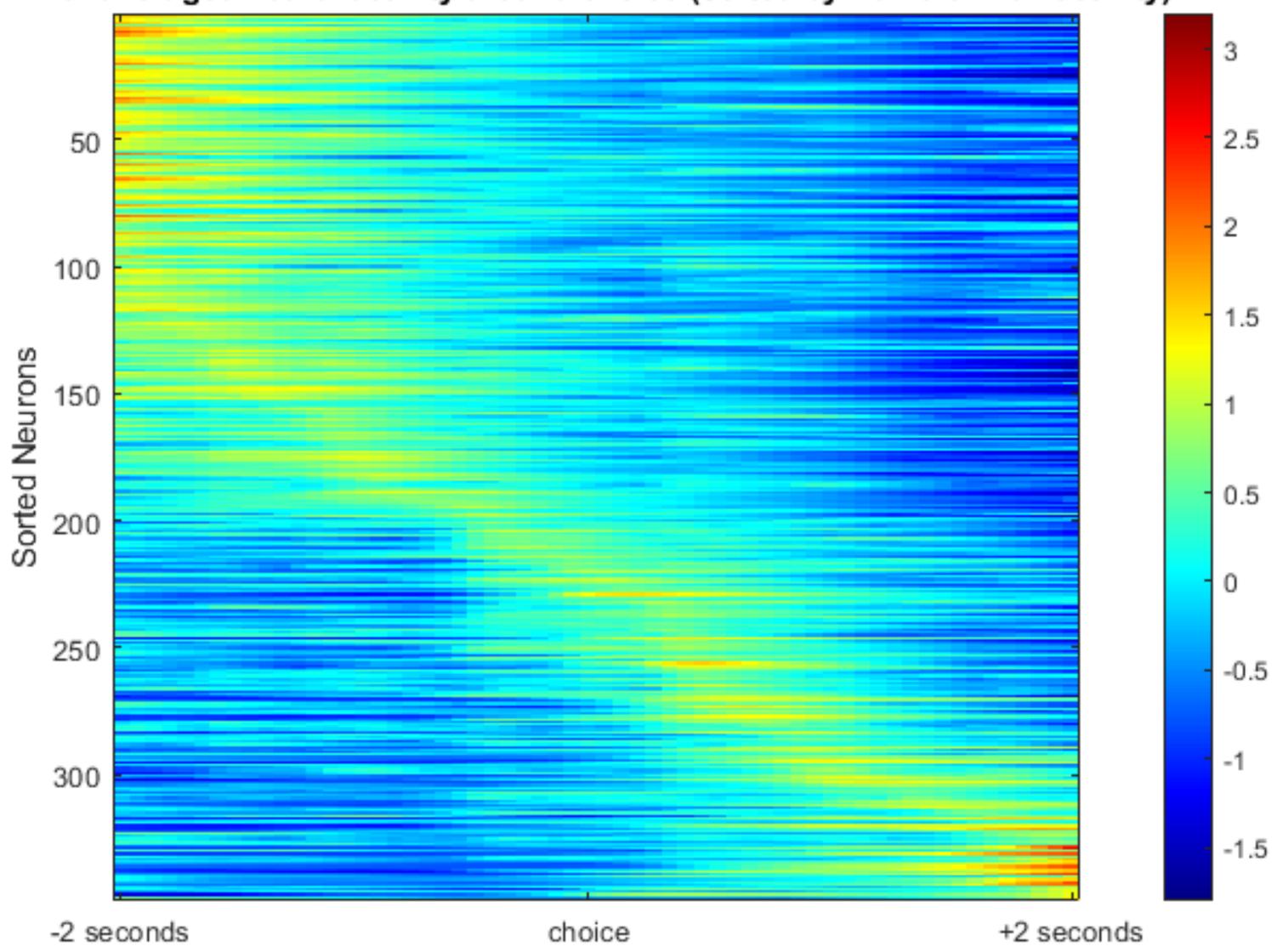
```
%loop through each neuron
\exists for neuron = 1:size(dff,1)
     % for one neuron
     % number trials x event length
     one_neuron = zeros([length(start_frame), seconds*frames per sec*2]);
     % loop through each trial, normalize neuron signal
     for trial = 1:length(start_frame)
         one neuron(trial,:) = normalize(dff(neuron, beginning(trial):endi
     end
     % take average of single neuron across all trials
     neuron activity(neuron,:) = mean(one neuron);
 end
 % sort neurons by peak activity
 [~, max pos] = max(neuron activity, [], 2);
 [~, max_order] = sort(max_pos);
```

nas2 = neuron activity(max order, :);

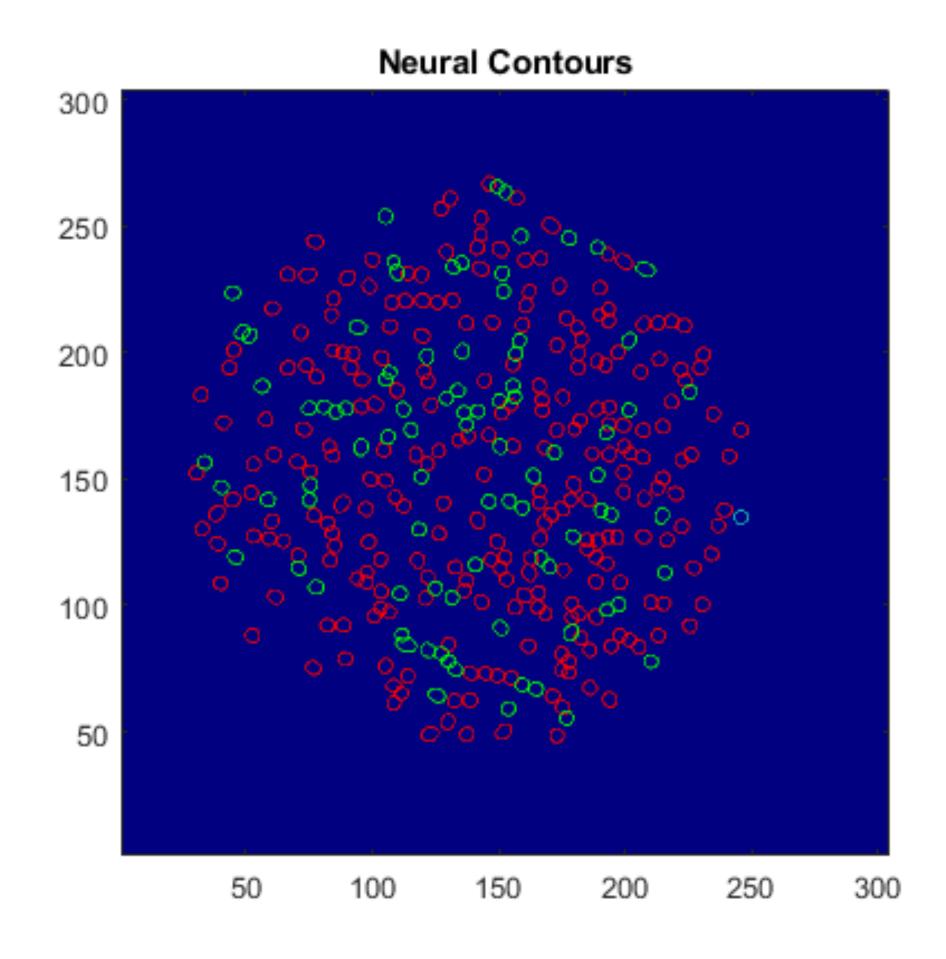


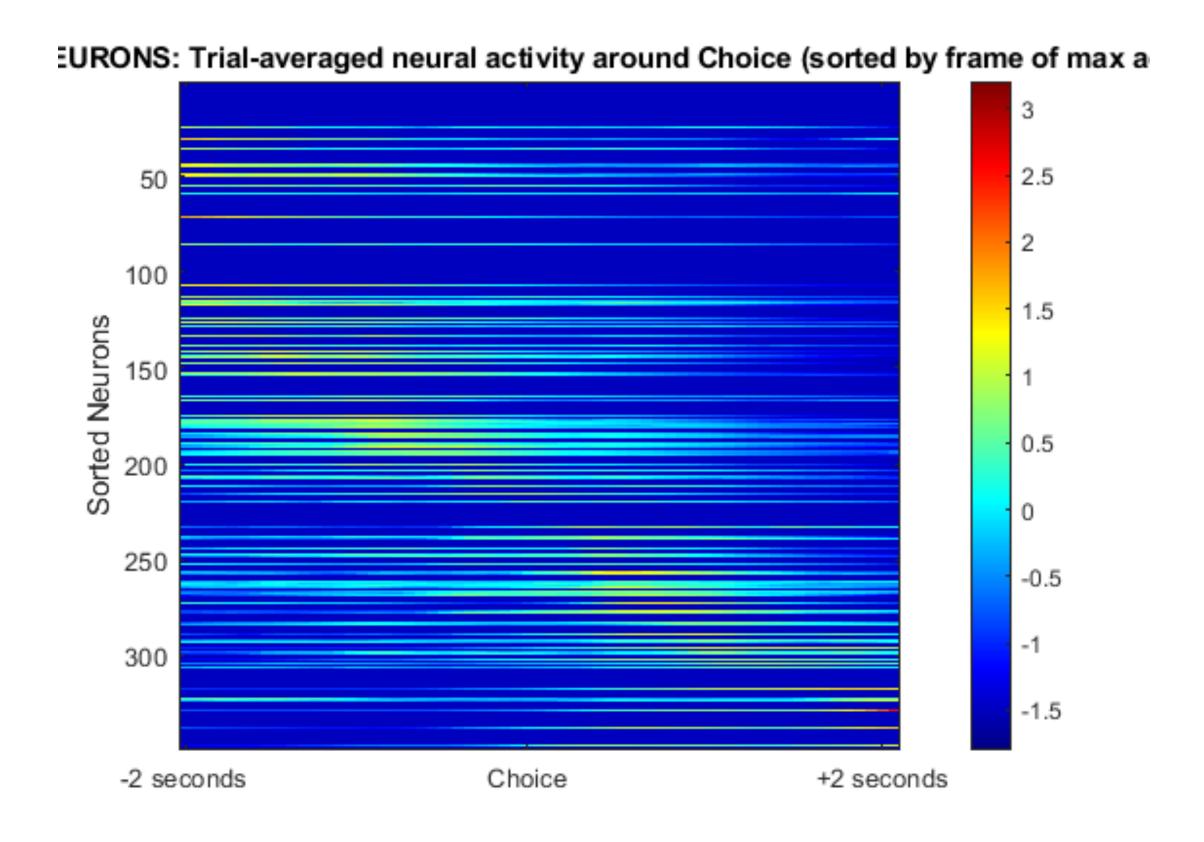






Permutation Test





PCA

This view is capturing what the *neural* population activity is doing! Each trajectory combines the activity from all neurons into a lower dimensional subspace, and each neuron contributes to this overall trajectory. We call this trajectory the "neural state."

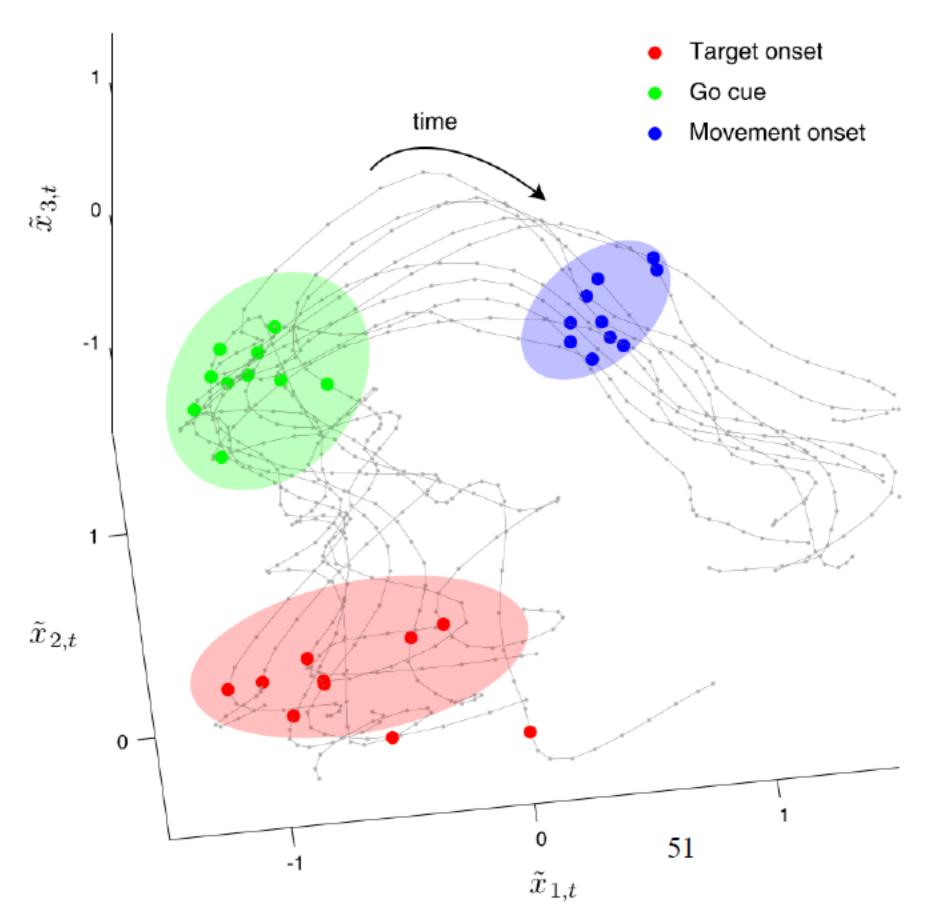
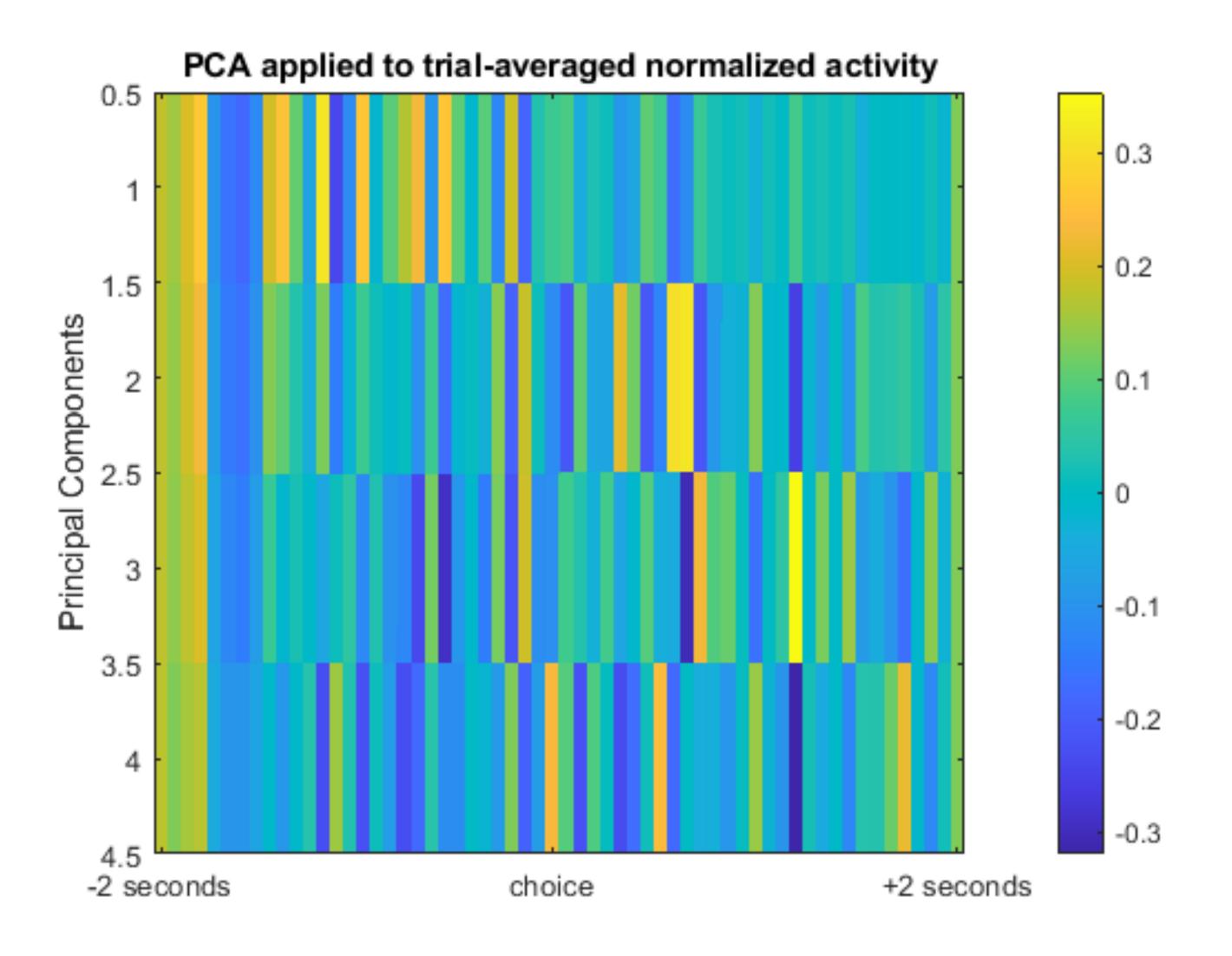


Photo Credit: Jonathan Kao, ECE C143A Lecture 18 Slides



- apply PCA on 349 x 61 matrix
- first principal component: high activity before choice
- second principal component: high activity after choice



Next Steps

- We need a more robust way of ID'ing "significant" neurons
- Use neural data in a neural network to predict binary outcome
- Read in more data with more animals
- Create computational model with various parameters such as perceived cost, velocity, etc.
- Optogenetic activation/silencing of ACC neuron subsets
- Project average population for different trial outcomes into lower-dimensional space

Neuron Projection

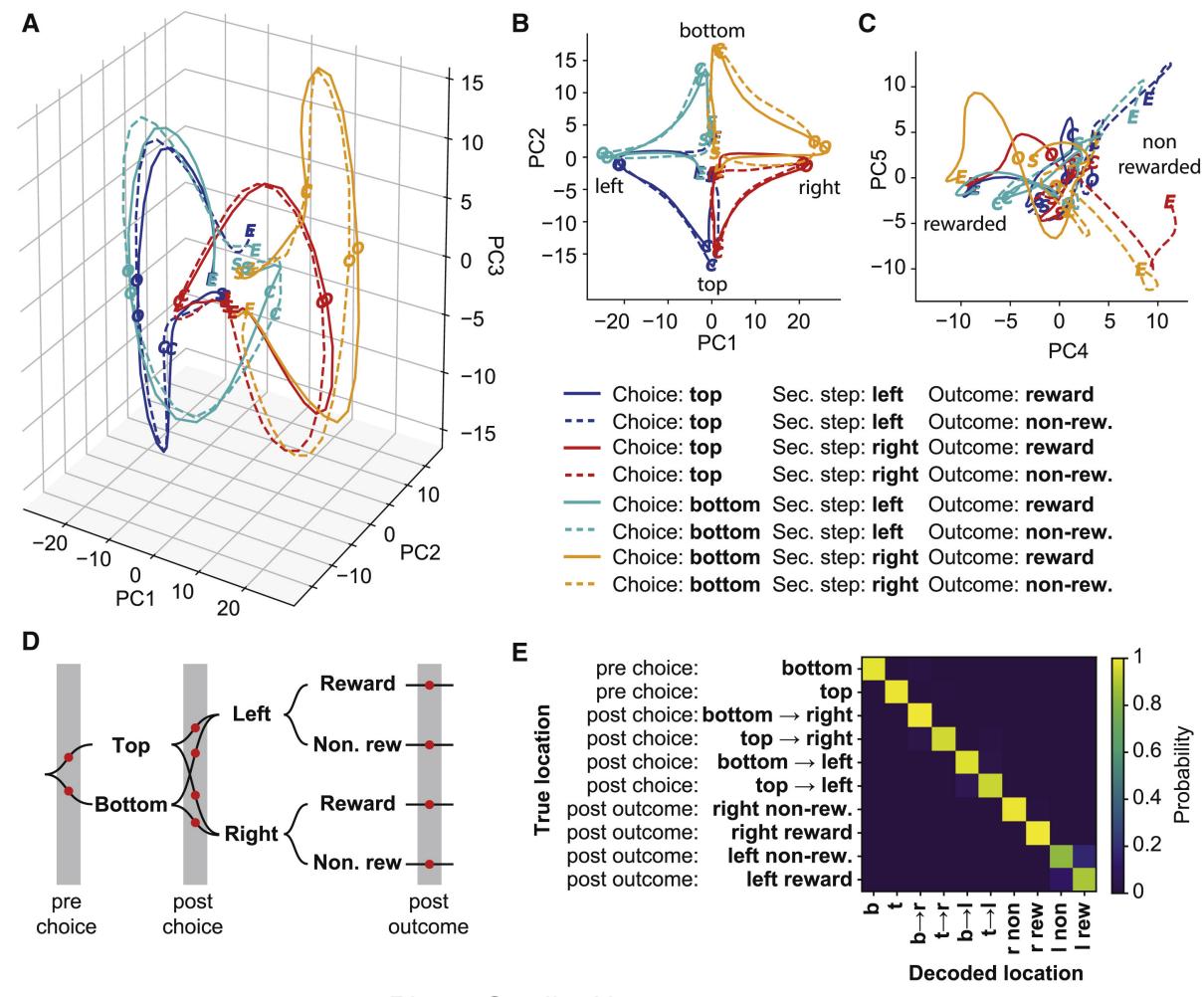


Photo Credit: Akam et al. 2021