This is the code for Ming’s paper: doi: https://doi.org/10.1101/2019.12.14.876201

The code uses CaImAn to analyze GCaMP dF/F imaging data.

**Ming’s software**

**Motion correction**

Ming uses TurboReg, a plugin for ImageJ for the motion correction.

<http://akiramuto.net/archives/208>

**CalmAn**

We then use CaImAn to find ROIs. Please set the MATLAB path to CaImAn because some routines are used by drgCaImAn\_script

<https://github.com/flatironinstitute/CaImAn-MATLAB>

**Olfactometer/readdropcspm\_hf**

This program is used to generate the percent correct vs trial plots. The input is from spm.mat files

**CaImAnDR/drgCalmAn\_script.** Finds components (ROIs) using CaImAn code. It ouputs a file called file\_name\_CalmAn.mat.

**CaImAnDR /drgCalmAn\_dropc.** This routine takes the output from drgCalmAn\_script (file\_name\_CalmAn.mat), the lick recordings from the INTAN board (.rhd format) and metadata from dropcspm\_hf.m to generate average deltaF/F and lick traces.

The output is saved in a file with the suffix \_pre\_per.mat

The output can be used for perceptron analysis, etc.

**CaImAnDR/drgCaImAn\_batch\_dropc**. This does a batch of warping and drgCalmAn\_dropc. If you want warping enter

do\_warp=1;

If you do not want to warp

do\_warp=0;

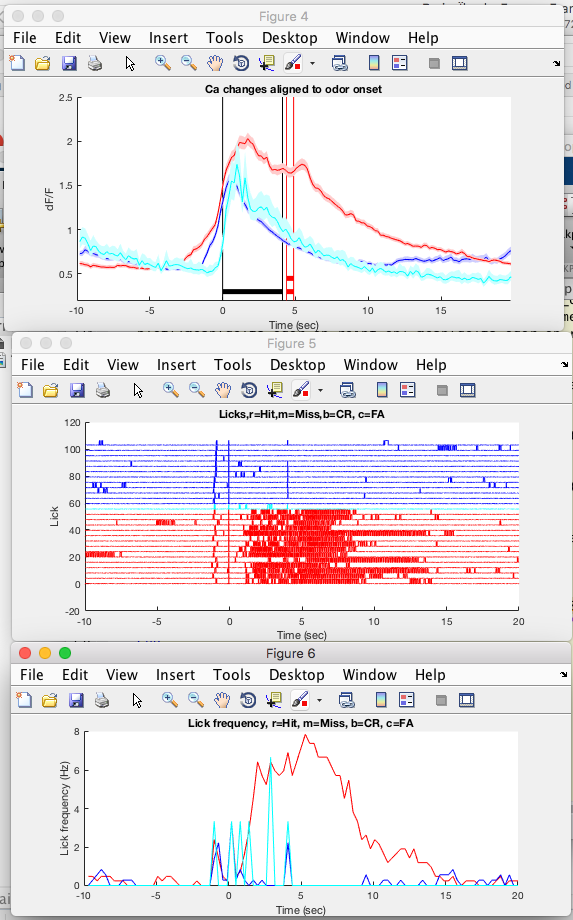
You need to have a file called:

drgCaImAn\_dropc\_choices\_some\_name.m

that has all the files to be processed.

**CaImAnDR /drgCaImAn\_dropc\_correlation.** This routine takes the output from drgCalmAn\_script (file\_name\_CalmAn.mat), the lick recordings from the INTAN board (.rhd format) and metadata from dropcspm\_hf.m to calculate correlation (and covariance) of Ca traces between components. The covariance is an input to for simulations.

**CaImAnDR /drgCaImAn\_average\_betweenSession\_dFF** averages the data between different sessions. It needs the output file generated by drgCaImAnBatchPerSession



**CaImAnDR /drgCaImAnBatchPerSession.** This routine takes input from a file named drgCaImAnChoices\_name of the file.m

It uses the output of drgCaImAn\_dropc to perform whole session analysis. Make sure to list the files in the Choices file in order of acquisition.

It generates:

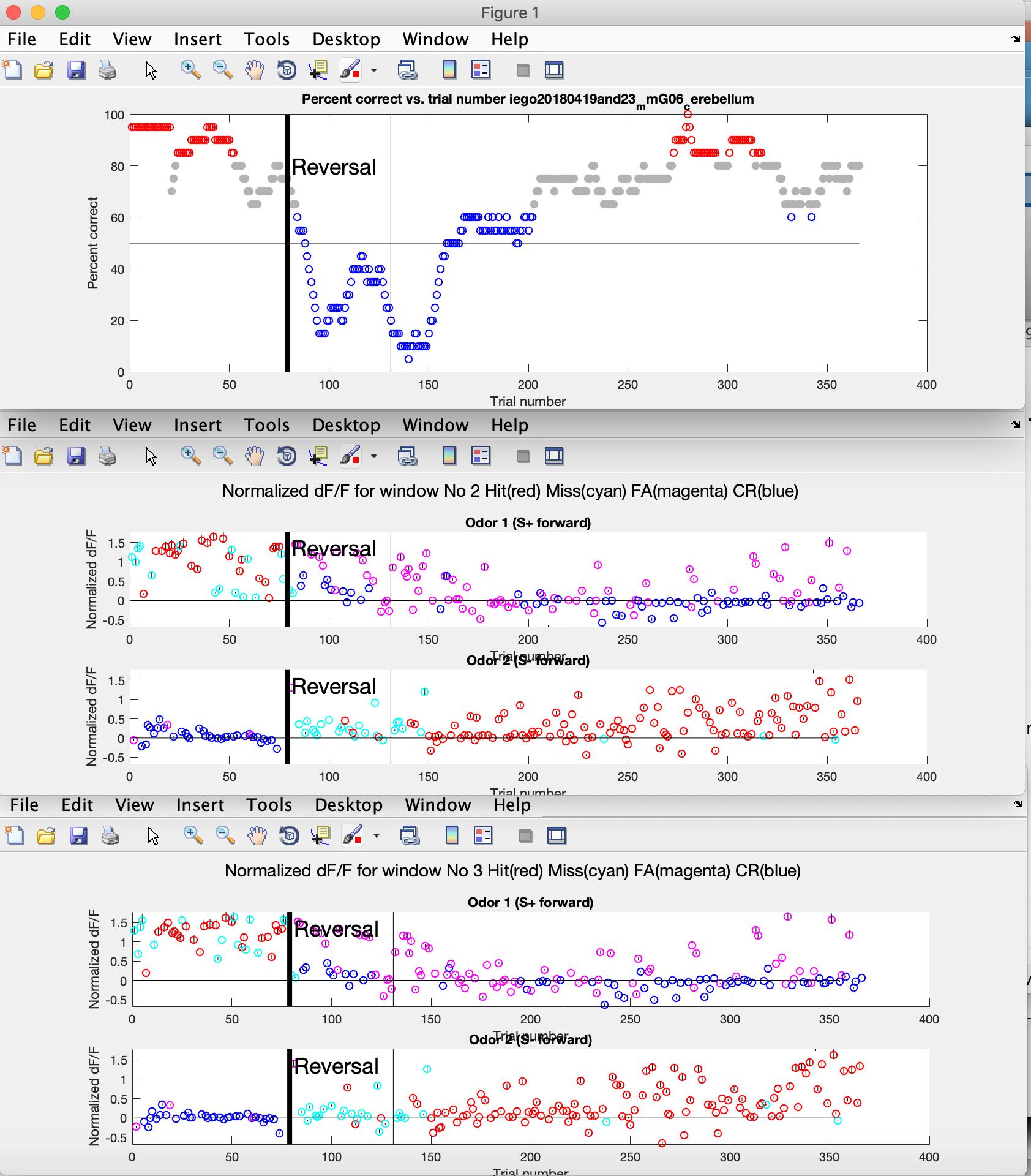
A screenshot of a video game

Description automatically generated

**CaImAnDR /drgCaImAnBatchPerSessionReversal.** This is used for the reversal figure. This routine takes input from a file named drgCaImAnChoices\_name of the file.m

It uses the output of drgCaImAn\_dropc to perform whole session analysis. Make sure to list the files in the Choices file in order of acquisition.

This code accepts multiple sessions



**CaImAnDR/drgOpticalFlowEstimate.** Optical Flow Estimation Using the Farneback Algorithm

The input is the .mp4 movie. Saves \_optflow.mat

**CaImAnDR/drgOpticalFlowEstimateBatch**

Batch optical flow estimation for multiple files using the Farneback algorithm. This code needs a choices file containing the names of the .mp4 input files (like drgOptFlowChoices20180419\_mmG06\_cerebellum.mat). The output is one \_optflow.mat file for each input file.

**CaImAnDR/drgOpticalFlowAligned.** Alignment of Optical Flow Estimation with the deltaF/F

The inputs are the pre\_per.mat generated by drgCalmAn\_dropc\_plot and optflow.mat generated by drgOpticalFlowEstimate.

**CaImAnDR/drgAlignOpticalFlow.** Allows the user to find whether the alignment is correct.

**CaImAnDR/drgCaImAnBatchPerSessionEventsPerTriallickvsdFFOptFlow**. This code does a movement and lick derivative analysis per trial. The program needs a choices file such as drgCaImAnChoicesDiego20180910\_mmPVG04\_CerebellumLDAOpFlow

**CaImAnDR/drgCaImAnBatchPerSessionEventsPerTriallickvsdFFOptFlowDimensionality.** This code does a dimensionality analysis.

**CaImAnDR/drgCaImAn\_glm.** This code does the glm analysis of a file generated by drgCaImAnBatchPerSessionEventsPerTriallickvsdFFOptFlow.m

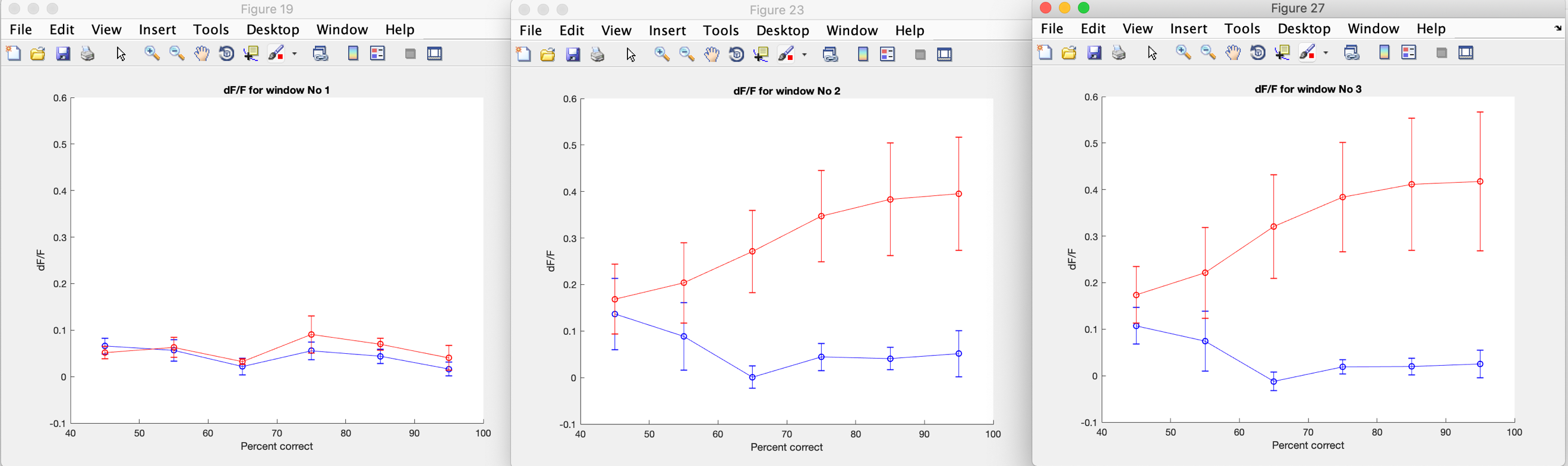
**CaImAnDR/drgCaImAn\_dropc\_perceptron.** Performs perceptron analysis.

**CaImAnDR/drgCaImAn\_rainbow**. Generates pseudocolor plots of the deltaF/F traces per ROI

**CaImAnDR/drgCaImAn\_get\_warped\_components**. This function warps the ROIs of the reference tif to the other tif. It saves data appropriate for drgCalmAn\_dropc

**CaImAnDR/drgCaImAn\_average\_betweenSession\_dFF** averages the data between different

sessions. It needs the output file generated by drgCaImAnBatchPerSession

****

**drgCaImAnBatchPerSessionReversalPerTrial**

Needs a choices file such as drgCaImAnChoicesDiego20180910\_mmPVG04\_Cerebellum. Needs the output files from drgCaImAn\_batch\_dropc.m

**drgCaImAnBatchPerSessionReversalPerTrialLDA.** Generated the example of LDA timecourse in Ming’s Fig. 3. This example is generated with drgCaImAnChoicesDiego20180910\_mmPVG04\_CerebellumLDA

**drgCaImAnBatchPerSessionEventsPerTrialPCA**

Sorts the predicted responses for Hit, Miss, CR and FA and generates a PCA analysis plot. It needs as input the output files of drgCaImAn\_batch\_dropc and a choices file such as drgCaImAnChoicesDiego20180910\_mmPVG04\_CerebellumLDA defining the windows (3-4 sec, etc) and making sure the sessions do not include reversals.

**drgCaImAnBatchPerSessionEventsPerTrialLDA**

Sorts the predicted responses for Hit, Miss, CR and FA and generates the output file for lda\_per\_event\_all\_files.m. It needs as input the output files of drgCaImAn\_batch\_dropc and a choices file such as drgCaImAnChoicesDiego20180910\_mmPVG04\_CerebellumLDA defining the windows (3-4 sec, etc) and making sure the sessions do not include reversals.

**lda\_per\_event\_all\_files.m**

Generates LDAperevent.fig showing that the Ca changes during mistakes can predict the odor, with some very interesting exceptions. It needs input generated by drgCaImAnBatchPerSessionEventsPerTrialLDA

**drgCaImAnBatchPerSessionEventsPerTriallickvsdFF**

Does the analysis of the slope of lick frequency vs. slope of dFF (as in Gaffield 2017). It needs as input the output files of drgCaImAn\_batch\_dropc and a choices file such as drgCaImAnChoicesDiego20180910\_mmPVG04\_

**drgCaImAnPerTrialLDASubsampleSimulation**

This code simulates activity of multiple components with different covariant activity to analyze the dependence of LDA prediction on the number of components

**drgCaImAn\_batch\_dropc\_no\_microscope.m**

This is used to process files acquired in a session to study the changes

in behavior and licks when the MLIs are activated with optogenetics

Needs as an input the ouput files from the dropcspm\_hf.m (.mat) and from

INTAN (.rhd)

Needs a dropc choices file such as

drgCaImAn\_dropc\_choices\_PVhM4Di\_04292019.m

or if you want to save time typing:

drgCaImAn\_dropc\_choices\_PVhM4Di\_all\_files\_04292019.m

**drgCaImAnBatchPerSessionLicks**

This code does a lick analysis for sessions where we record the behavior and licks

for optogenetic modulation of behavior/licks by turning MLIs with

optogenetics

Needs a choices file such as

drgCaImAnChoices\_20190312\_mmPVsoChR02\_Cerebellum\_licks.m

% Needs the output files from drgCaImAn\_batch\_dropc\_no\_microscope.m

**drgCaImAnBatchMultiSessionLicks**

drgCaImAn\_multichoices\_PVhM4Di\_all\_files\_05052019

**drgCaImAnBatchPerSessionPerTrialLDASubsample**

This code does a linear discriminant analysis for spm data using subsets of components

Needs a choices file such as drgCaImAnChoicesDiego20180910\_mmPVG04\_Cerebellum

Needs the output files from drgCaImAn\_batch\_dropc.m

**Zoe Donaldson’s LDA demo**

This is a demo of our use of LDA

**LDA\_demo** calls drgCaImAnLDAtimecourse.m to process data from Ming’s Figure 3C (MLI data.mat data available on request).

**drgCaImAnLDAtimecourse** performs LDA for MLI data.mat

**Connor McCoullough’s software**

**drgcmCaImAn\_batch\_dropc\_rhd.m** reads the images, olfactometer file and rhd for Connor’s analysis. You need a choices file such as drgCaImAn\_dropc\_rhd\_choices\_157565\_9\_27CA.m

**drgcmPostHocLickReward.m** re-evaluates the scoring of the hits, CRs, etc.

drgcmCrossPCArhddFF takes as an input the \*\_batch\_per\_file.mat and outputs the correlations between LFP and dFF.

drgCaImAn\_dropc\_nsampler