

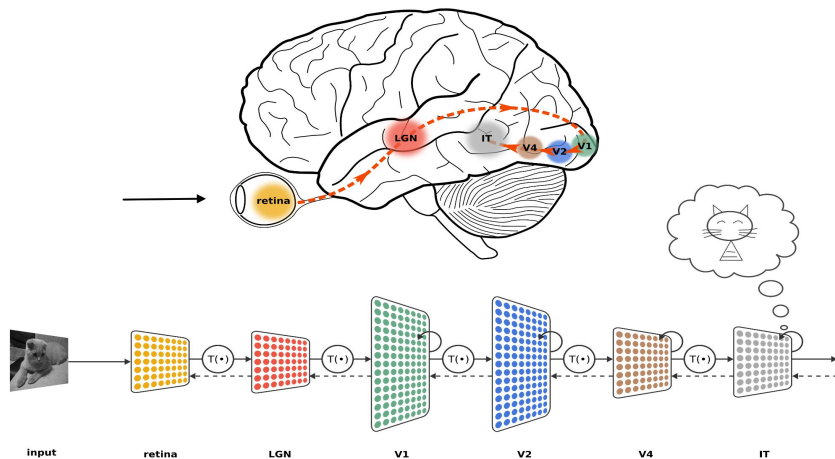
Does engagement matter: Do mice see the world differently when they don't care?



By: Ryan, Franck, Ankit, Jan, Helene, Manusmriti

Team: **AwareWolfs** (formerly the *Careful Wolves*)

Does engagement matter?



Kubilius, Jonas (2017): Ventral visual stream. figshare. Figure.
<https://doi.org/10.6084/m9.figshare.106794.v3>

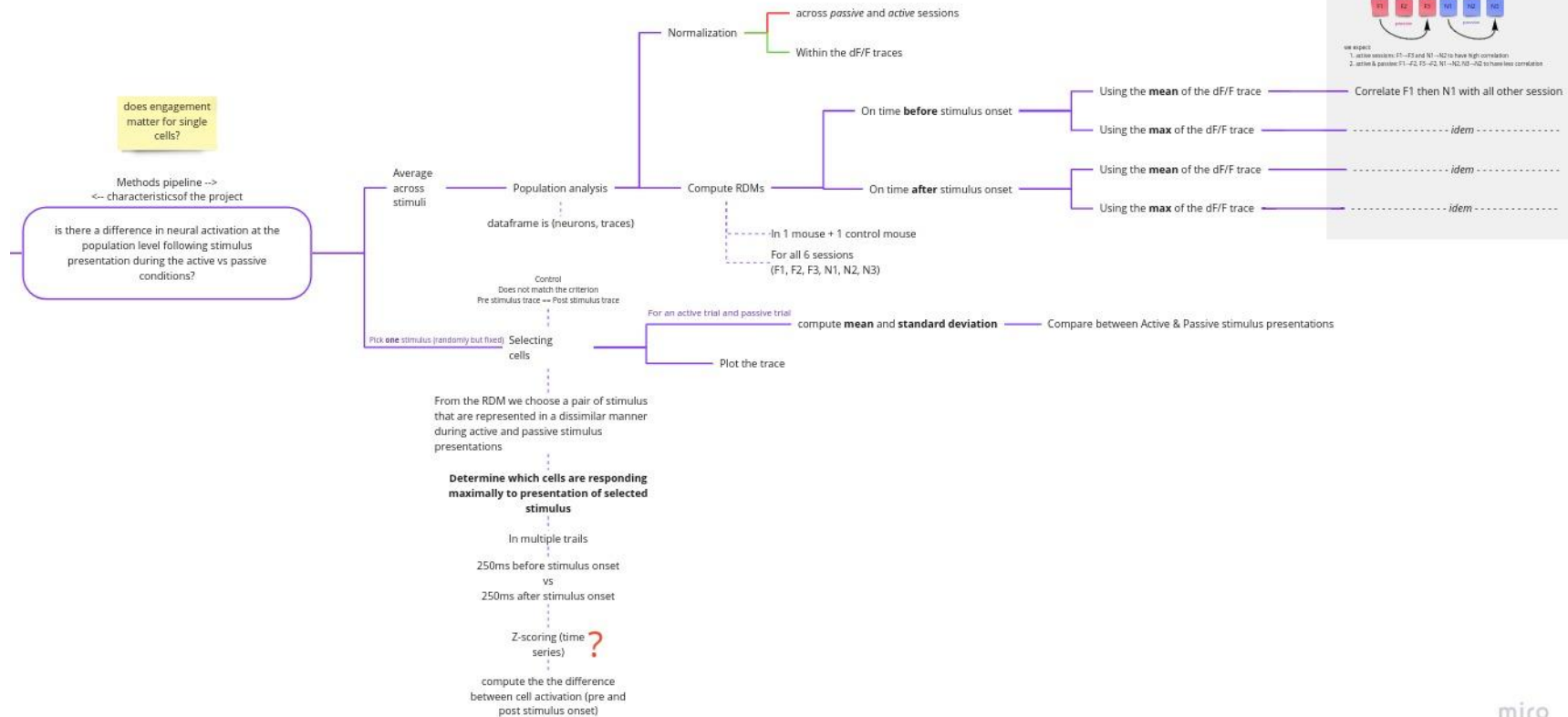
Traditionally V1/VISp is considered a simple feature detector

Stimulus representation in A1 adapts to engagement in go/no-go task
(Bagur et al 2018)

Does stimulus representation in V1 also differ depending on **engagement** (active) and **disengagement** (passive)?

Investigation on cell & population level

Our Plan



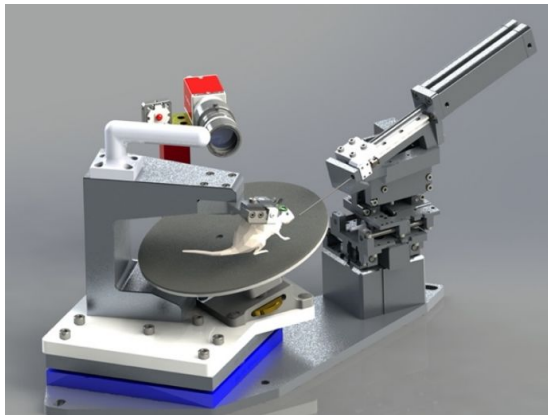
miro

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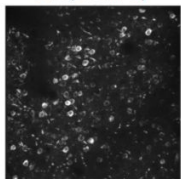


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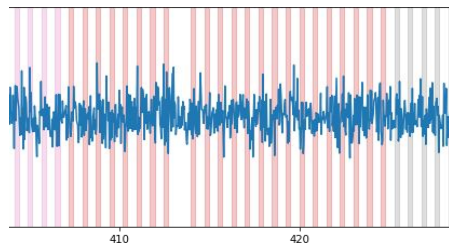
The Allen Institute 2P Visual Behavior Dataset



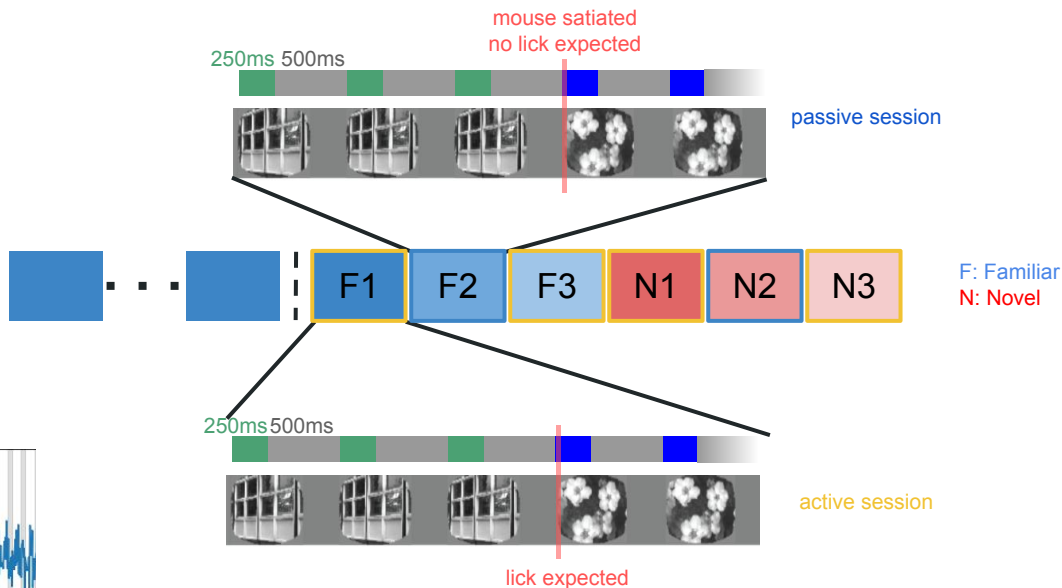
Excitatory
Slc17a7-IRES2-Cre;CaMk2-1TA;
Ai93(GCaMP6f)



2-photon imaging



dff traces (fluorescence)



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Population Analysis

Let's look into the representation of each stimulus in **active** and **passive** sessions

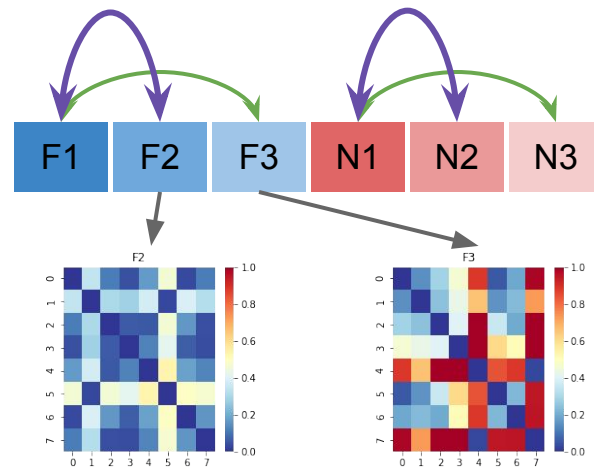
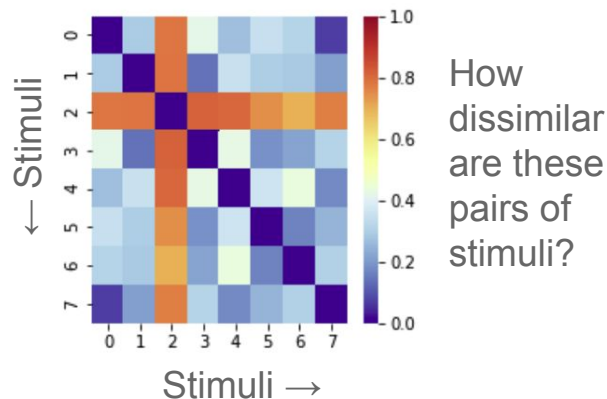
→ Representational dissimilarity matrices (RDM)

Time Series and sessions:

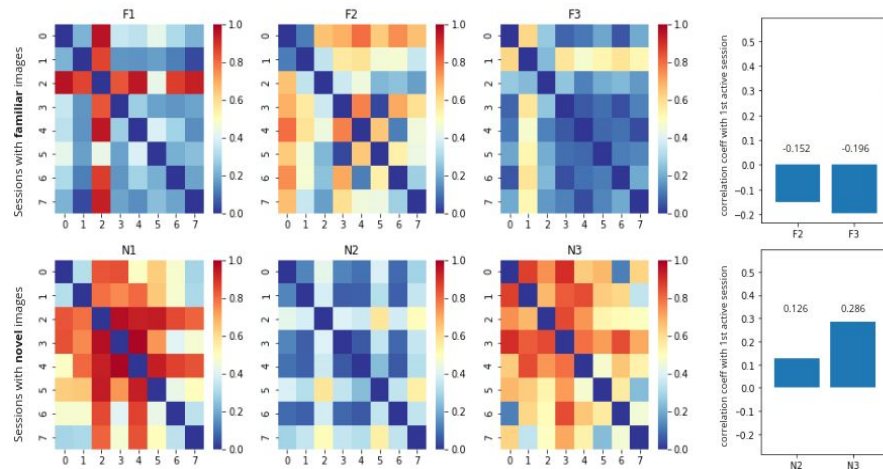
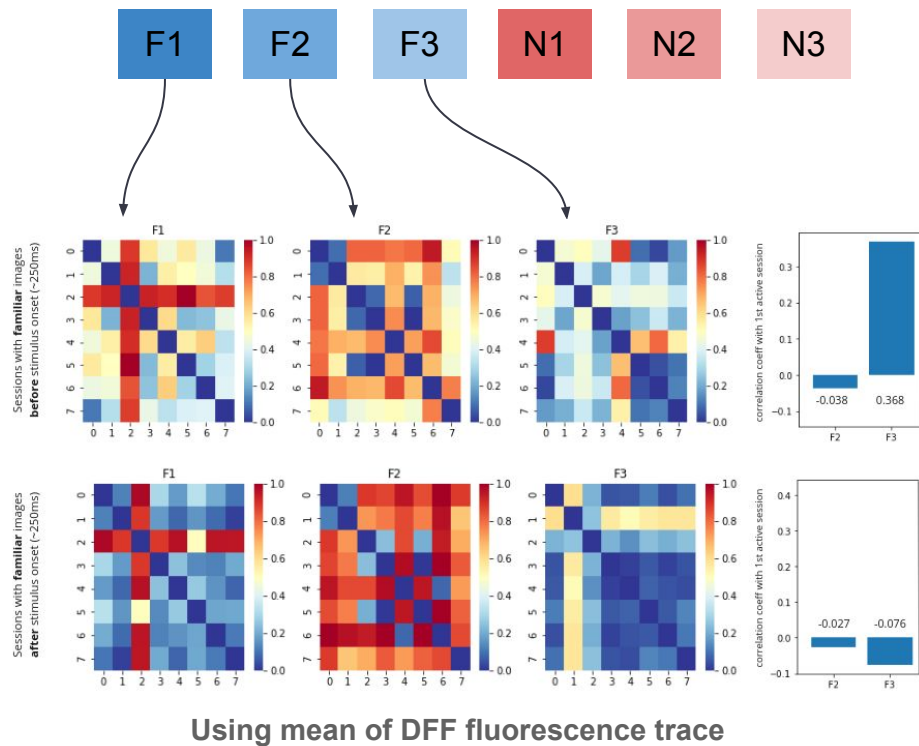
- 250ms **before** and **after** stimulus onset
- All 6 sessions: **familiars** (3) and **novels** (3)
- **Mean & max** of trace for all presentations of one stimulus

Correlate RDMs & expectations

- F1→F2 and N1→N2 | **Low correlation**
- F1→F3 and N1→N3 | **High correlation**



A few results



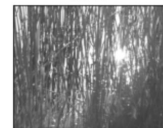
➔ Generally, the correlations remain consistent

Future directions :

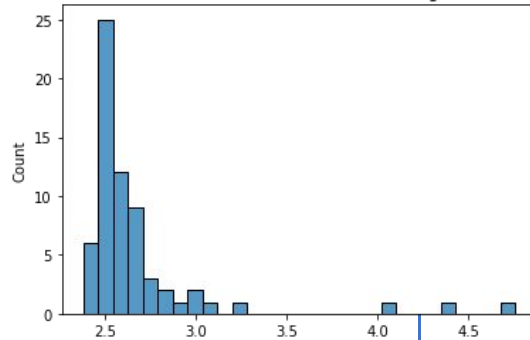
- Filter active cells
- Keep only cells common to sessions
- Take into account cell activity before versus after stimulus onset

Single-Cell Analysis

im077

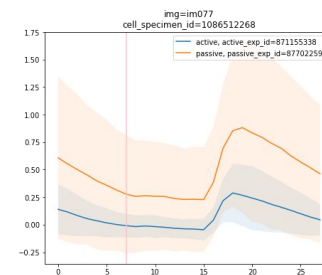
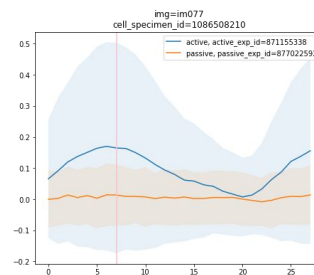
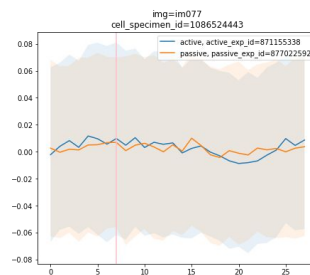
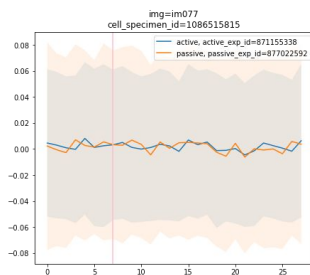
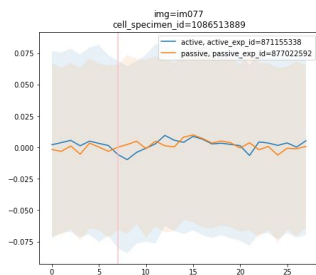
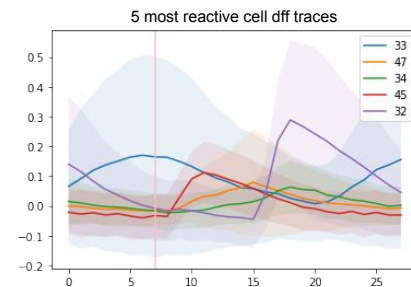
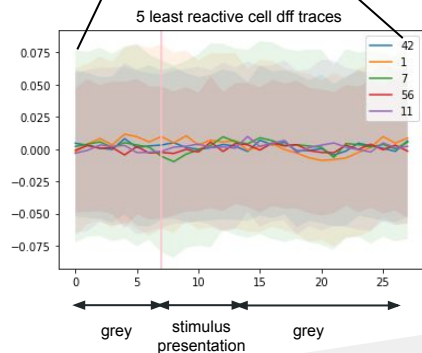


Cells count of maximum zscore (baseline before onset) averaged over stimulus presentation



Lowest z-score (least reactive)

Highest z-score (most reactive)

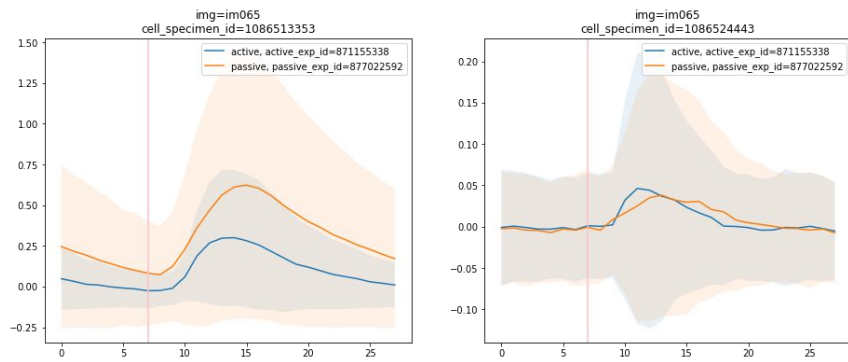


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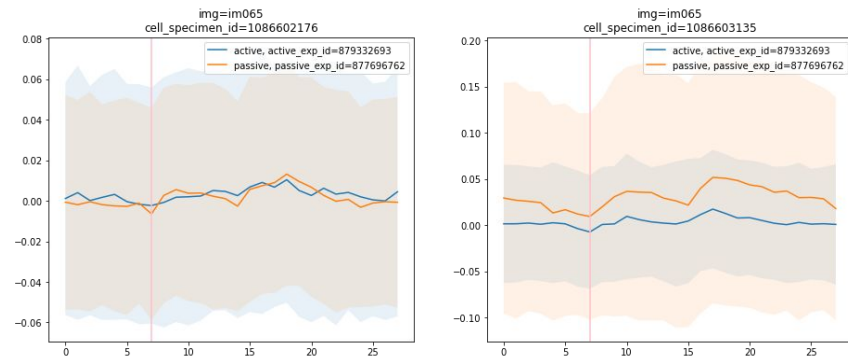
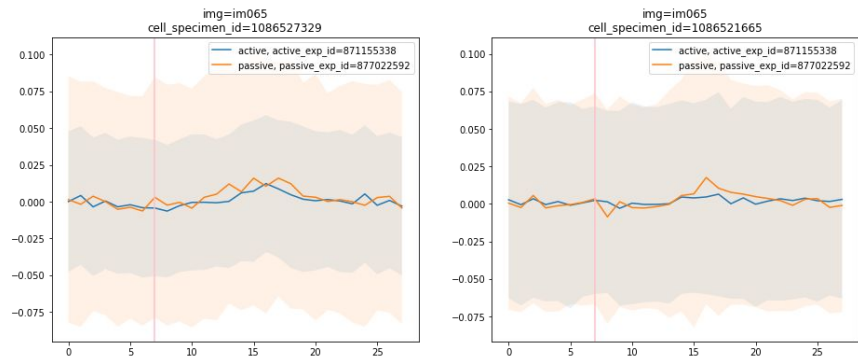
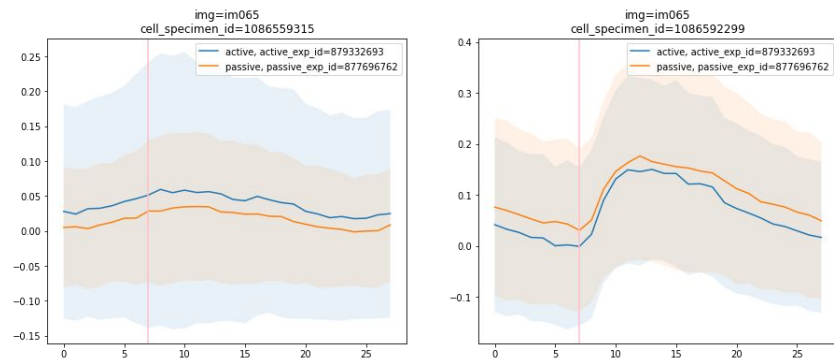


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Experiment (fast degrading reporter line)



Experiment (slow degrading reporter line)



Control (fast degrading reporter line)

Control (slow degrading reporter line)

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Conclusions

- On **single cell** level our methods yield no clear results.
- **Population level** dissimilarity (RDM) hints at the difference between active and passive session, but with control (pre-stimuli) this difference appears *independent of the stimuli*.

Discussion

- Despite the above findings there is some variability from general observation. This can be potentially mitigated by using more **behavioral features** (e.g., pupil dilation, running speed).
- **Single cell analysis** — in the passive condition cell responses *seem to increase* in many cases (though not consistently observed).

The Awarewolfs thank their stars!



Dr. Juliane Jaepel

Our kind mentor



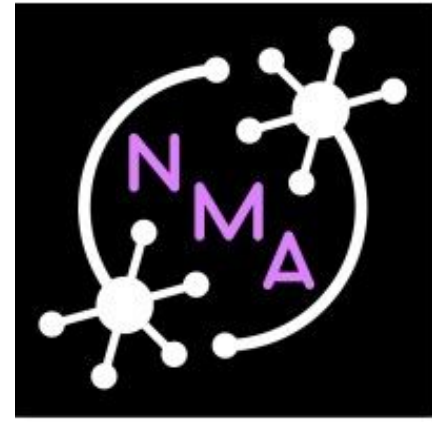
Nishant Jana

Leading the pack



Dr. Agustín Sanchez

Our dataset magician!



Neuromatch Academy

for helping us learn together

References

Bagur, S., Averseng, M., Elgueda, D., David, S., Fritz, J., Yin, P., Shamma, S., Boubenec, Y., & Ostojic, S. (2018). Go/No-Go task engagement enhances population representation of target stimuli in primary auditory cortex. *Nature Communications*, 9(1). <https://doi.org/10.1038/s41467-018-04839-9>

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Montijn, J. S., Goltstein, P. M., & Pennartz, C. M. (2015). Mouse V1 population correlates of visual detection rely on heterogeneity within neuronal response patterns. *ELife*, 4, e10163. <https://doi.org/10.7554/eLife.10163>

Our fabulous dataset can be found at https://allensdk.readthedocs.io/en/latest/visual_behavior_optical_physiology.html

Before stimulus onset

After stimulus onset

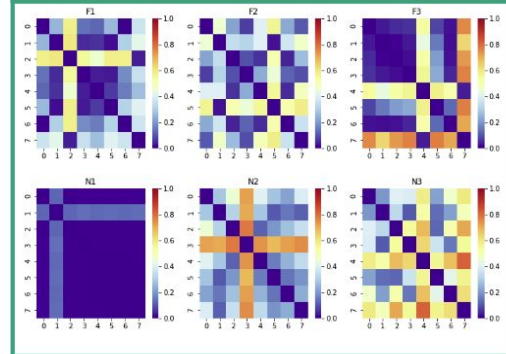
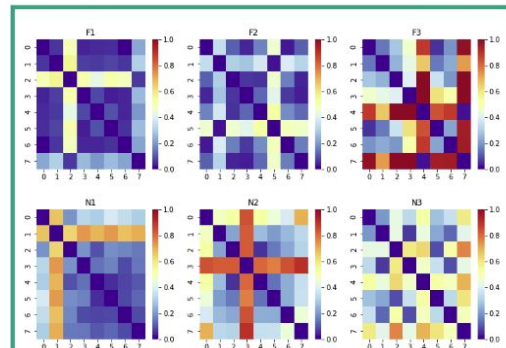
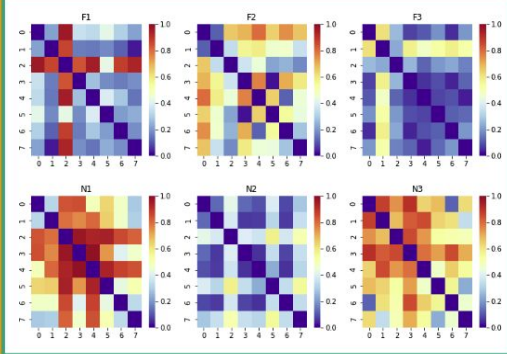
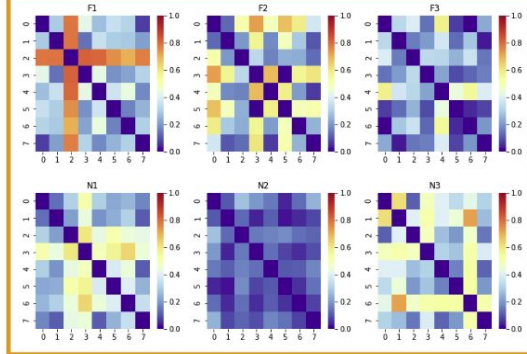
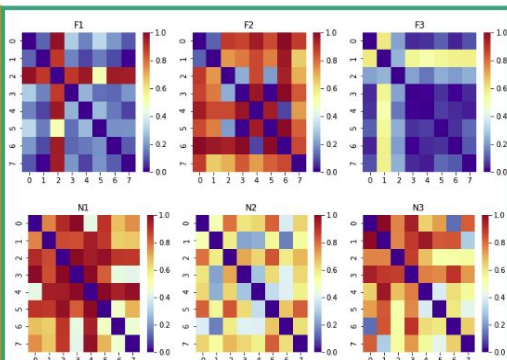
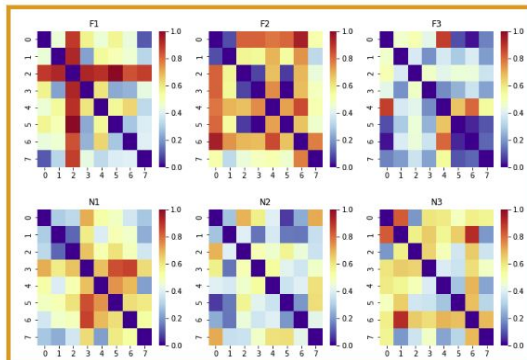
After stimulus onset

mean DF/F trace

mean DF/F trace

maximum DF/F trace

maximum DF/F trace



MOUSE 1

MOUSE 2

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The Journey

- many ideas, we choose one
- population or individual cells \Rightarrow why not both?
- population:
 - hey, t-SNE and PCA might be an option
 - use RDMS
- individual cells:
 - mean traces? but how?
 - use z-scoring
 - how to select them?
- later realized there are estimated spiking events
 - but we stayed with dff traces