# Retinotopic Map Exploration

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Injera/ RetinaZ









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# Who Are We?



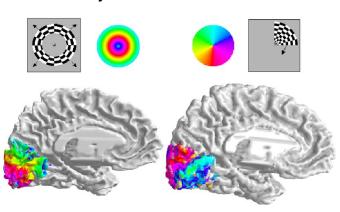
#### Contents:

- Intro
- Objective
- Methodology
- Results
- Conclusion

#### Retinotopy:

- Mapping of visual input from the retina to neurons
- The acquisition, representation, and Analysis of human retinotopic maps have required in the use and development of various methods to investigate the neural activity of the visual cortex

- Applications:
- Computes the population receptive field from responses to a wide range of stimuli
- Discovering the sizes and surface areas of different visual areas (V1, V2, V3) and finding the correlation between them [3]



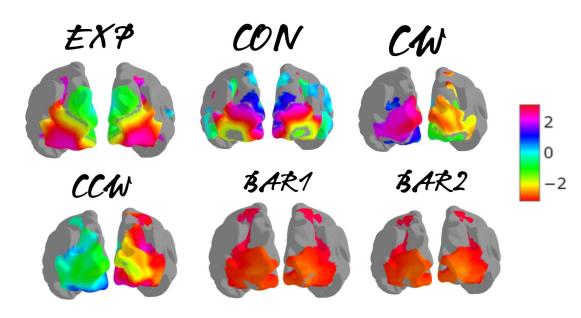
# Q: How do neural responses differ when exposed to different types of stimuli?

• Sub question 1: Does the whole brain activity (average responses) show difference for each

Stimulus?

Sub question 2:

How the neural responses differ among voxels?



## Hypotheses

**H1:** Same responses when comparing averaged voxels

The data is averaged over 183 human subjects and the distribution of 91282 voxels responses will hide the exact differences

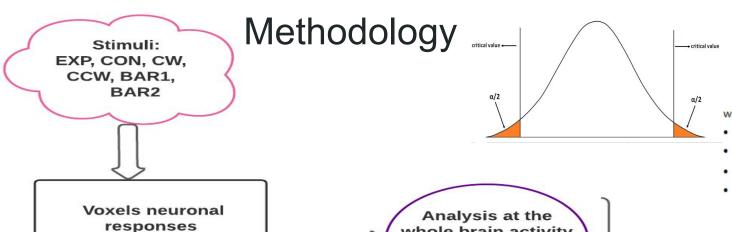
$$mu\_CON = mu\_EXP = mu\_CW = mu\_CCW = mu\_BAR1 = mu\_BAR2$$

H2: Each voxels respond differently to stimuli

The neuronal response of the voxel depends on the selectivity to visual stimulus features

### Dataset: HCP Retinotopic

183 Human subjects | B \* 22 s 22 s BLANK BLANK 6 stimuli: 1) RETCCW {CCW - CW - Exp - Con 22 s 22 s BLANK BLANK - Bar1 - Bar2} 2) RETCW 300\*91282 22 s 22 s BLANK **BLANK** 3) RETEXP 22 s 32 s 22 s BLANK BLANK BLANK 4) RETCON × 8 32 s 16 s 32 s 32 s 32 s 12 s 32 s 32 s 32 s 16 s 5) RETBAR1 **BLANK** BLANK **BLANK** 6) RETBAR2 UP RIGHT LEFT DOWN UPPER LOWER LOWER



T-test

$$t = \frac{\overline{x} - \mu_0}{s / \sqrt{n}}$$

where:

- x = the sample mean
- μ<sub>0</sub> = the hypothesized population mean
- s = the sample standard deviation
- n = the sample size

$$t = \frac{(\overline{x}_1 - \overline{x}_2) - (\mu_1 - \mu_2)}{\sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}}$$

Visualisation

Regions of Interest Analysis at the

Results for the

The Proposed process

91282 \*300 time steps

Analysis at the voxel-by-voxel level in all the brain with

t-test

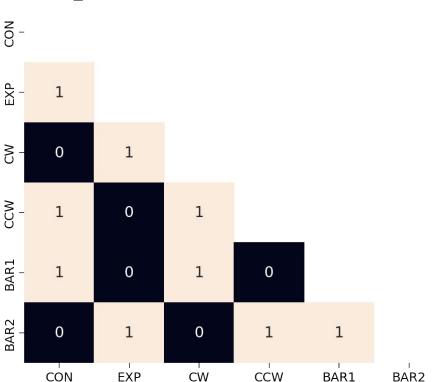
voxel-voxel level in the ROI with t-test

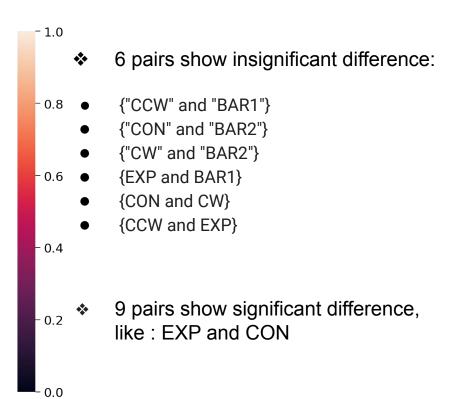
whole brain activity

with T-test

### The Analysis at the distribution level

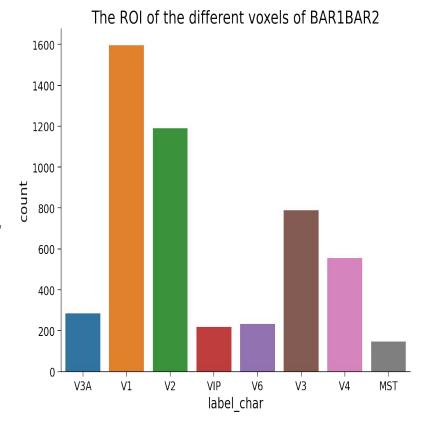
T\_Test Results at the distribution level



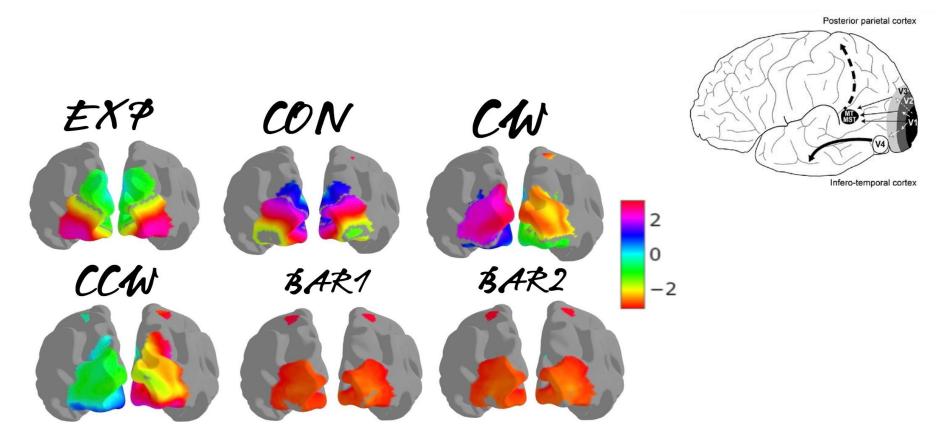


## The analysis at the voxels level

- Voxels Level for whole brain:
  - The t-test: more than 93% of the voxels are statistically different.
- Voxels Level for Regions of interest of:
  - ROIs: Visual cortex areas tuned to moving objects, color, orientation, speed, direction. [5][6][7]
  - There is slight difference for count of different voxels in each bair.

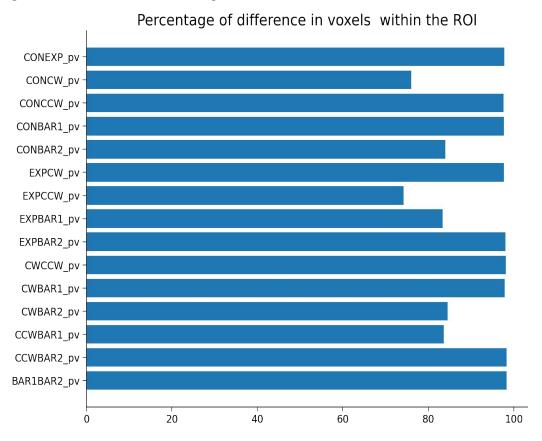


### Visualization of the Regions of interest activity



## Results of voxel-by-voxel analysis of ROIs

The t-test: Ranges in (74~98) % of the ROIs voxels are statistically different.



#### Conclusion

- **At the whole brain level**: neural response showed difference between stimuli of opposite patterns (ex. CW & CCW), which means we can still use the overall distribution of responses to differentiate between opposite patterns.
- At the voxel-by-voxel level for whole brain: >93% of voxels respond differently even with pairs that showed insignificant difference in the whole brain analysis level.
- At the voxel-by-voxel level for ROIs: 74% ~ 98% of voxels are selective to pairs of stimuli.

#### **Future work:**

Analyze difference between stimuli at regions level (average voxels response over region).

Individual fMRI dataset can be used in Biometrics widely as the dataset is individual selective.

#### References

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# Many Thanks to 💛 💛



Aryan



Dilip





