Model-based analysis of neurobiological data to am study the representation of natural features

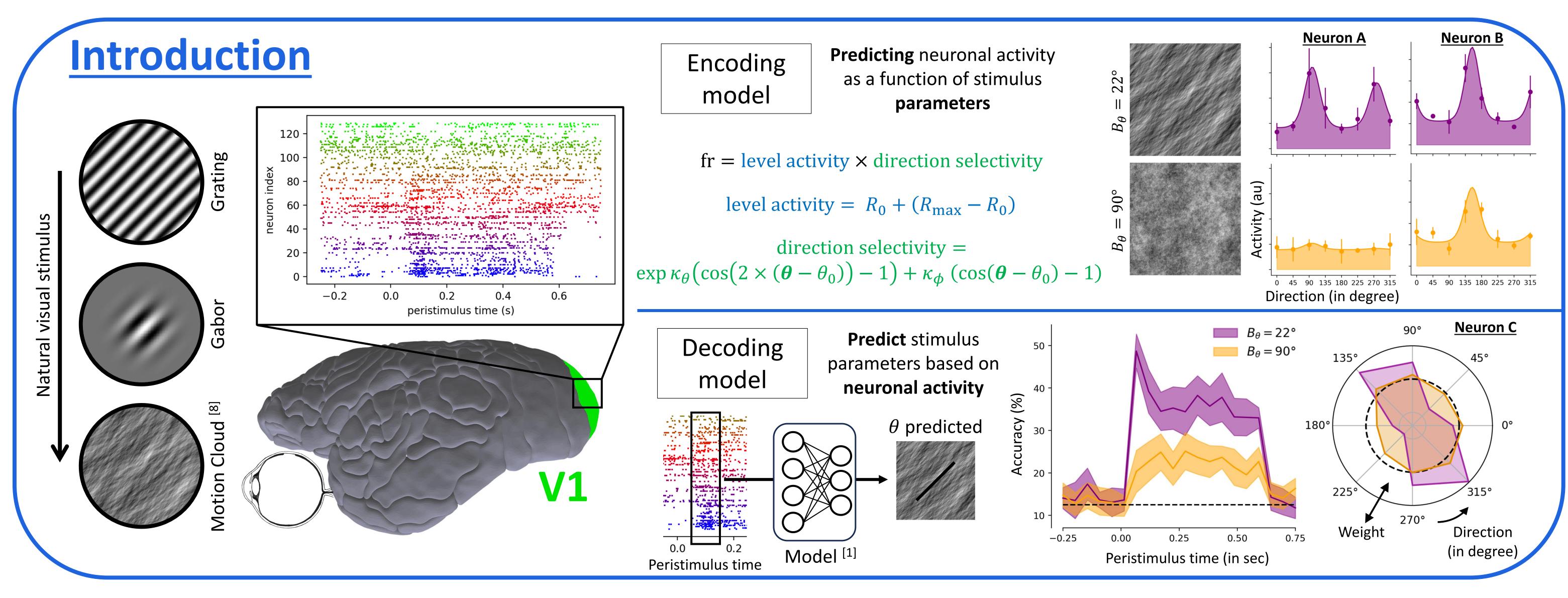
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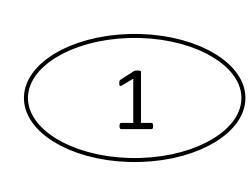




Project

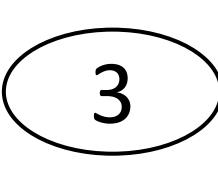
Decoding model **Encoding model** Rate code **Sparse connectivity** Stimulus F Synaptic diversity [3] Encoding window Reaction 111111 Slow 0.50 synapse 0.25 **Fast** synapse <u>Temporal code</u>: latency code <u>Temporal code</u>: interspike intervals -0.75**Spiking Neuronal** Network [3] Target neuron Input Stimulus Pattern detection [5] Predicted 10 stimulus Activity Plasticity mechanisms [2] **Neuron index** (%) E-I balance [4] slope Pre ____ Post ___ EPSP Normalized 48 96 144 -100 -150 Time (arb. unit) Pre/post spike interval (ms) Develop a population decoding model to test various Develop an encoding model that is biologically plausible code hypotheses, such as the existence of and whose result is close to a natural activity. spatio-temporal patterns.

Goals



Explore representations of information and hypothesise about the underlying mechanisms. [6, 7]

Developing new methods for analysing large-scale brain recordings.



Maintain the explicability of the analysis and avoid all the blackbox models.

References

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- [1]. Berens et al., 2012
- [2]. Bi & Poo, 1998
- [3]. Denève et al., 2017
- Denève & Machens, 2016
- Grimaldi et al., 2022
- [6]. Hénaff et al., 2020
- [7]. Ladret et al., 2023 [8]. Leon et al., 2012

[9]. Panzeri et al., 2010



GitHub

Bibliography

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from [9]

