

# An adaptable analysis pipeline makes cortical wave phenomena comparable across heterogeneous datasets

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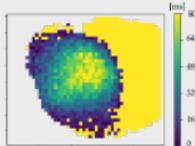


# Motivation: Slow Cortical Waves

## Imaging

a) Widefield VSD  
*(Figure proprietary)*

b) Widefield Calcium Imaging



c) Calcium Imaging  
*(Figure proprietary)*

## richness of data

Slow waves are observable

- across species,
- across scales,
- across methods.

## Lack of comparability

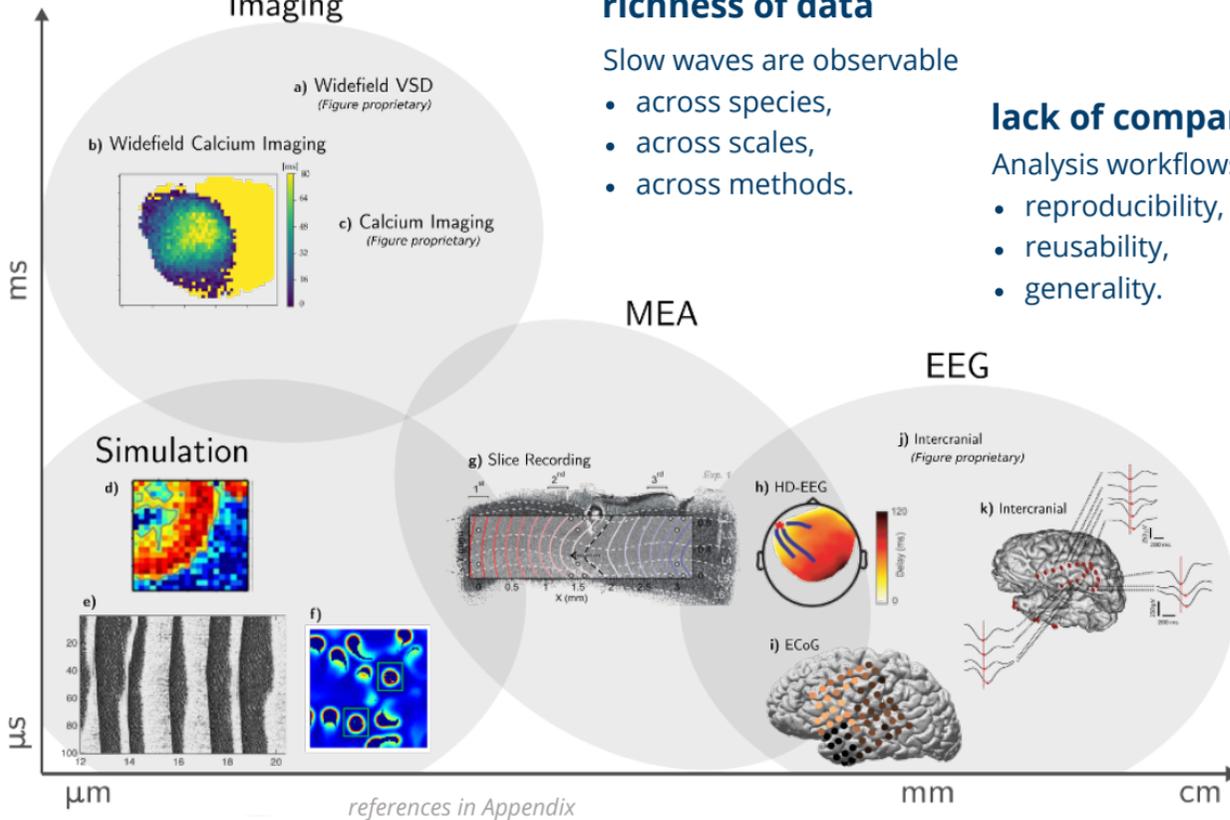
Analysis workflows often lack

- reproducibility,
- reusability,
- generality.

## cross-domain comparison

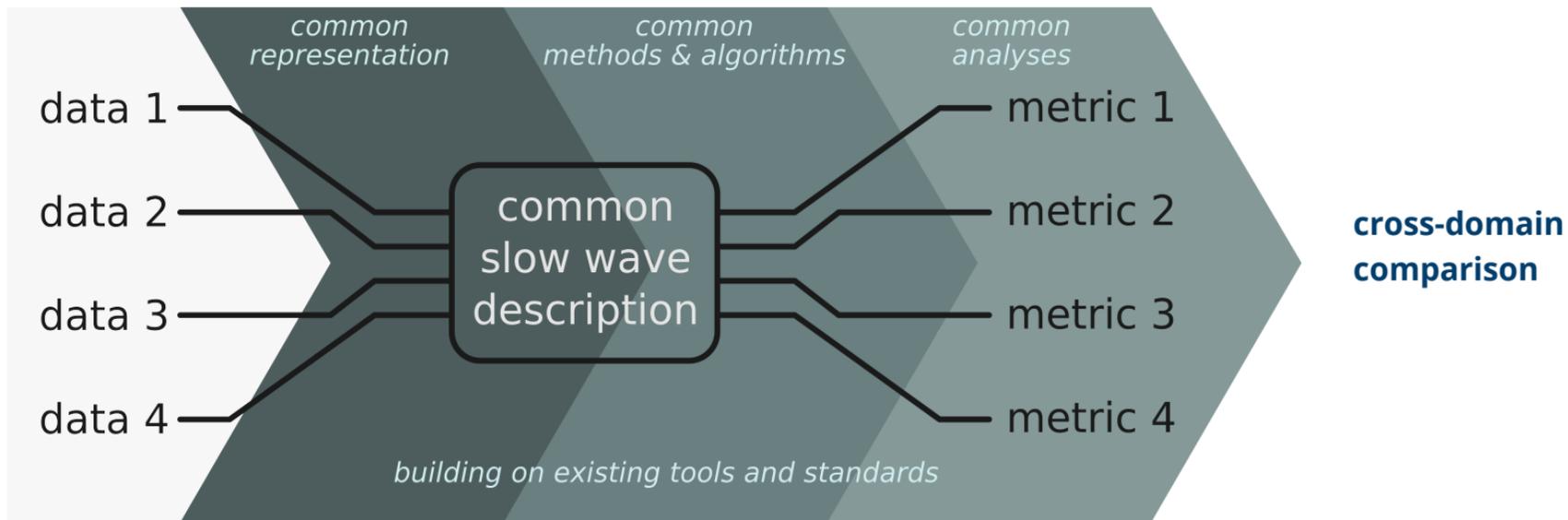
Comparability is needed for

- integration of data sources,
- model calibration & validation,
- quantifying experimental variability.



references in Appendix

# Modular Analysis Pipeline Approach



neo



CONDA



SPHINX  
PYTHON DOCUMENTATION GENERATOR



YAML

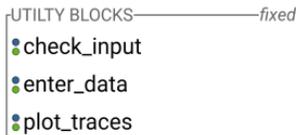


# The Slow Wave Analysis Pipeline

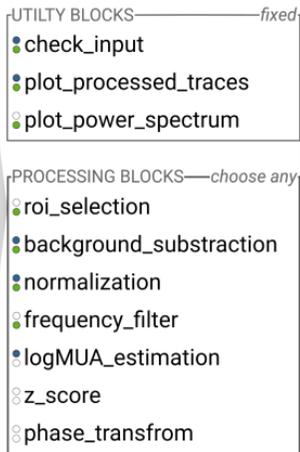
organizes the analysis steps in sequential stages  
of combineable blocks.

ECoG, Calcium Imaging,  
EEG, Spikes, LFP, Simulation, ...

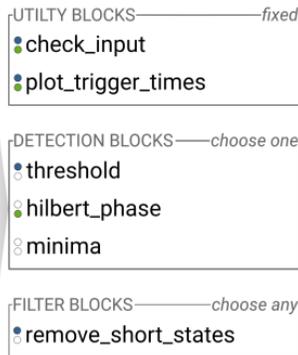
## Data Entry



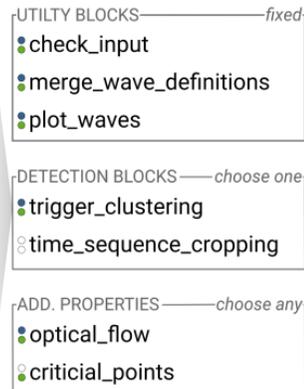
## Processing



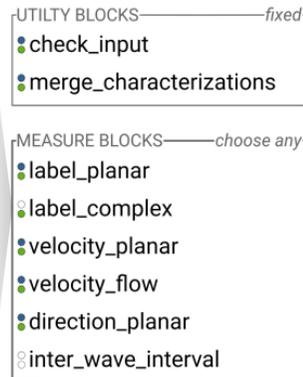
## Trigger Detection



## Wave Detection

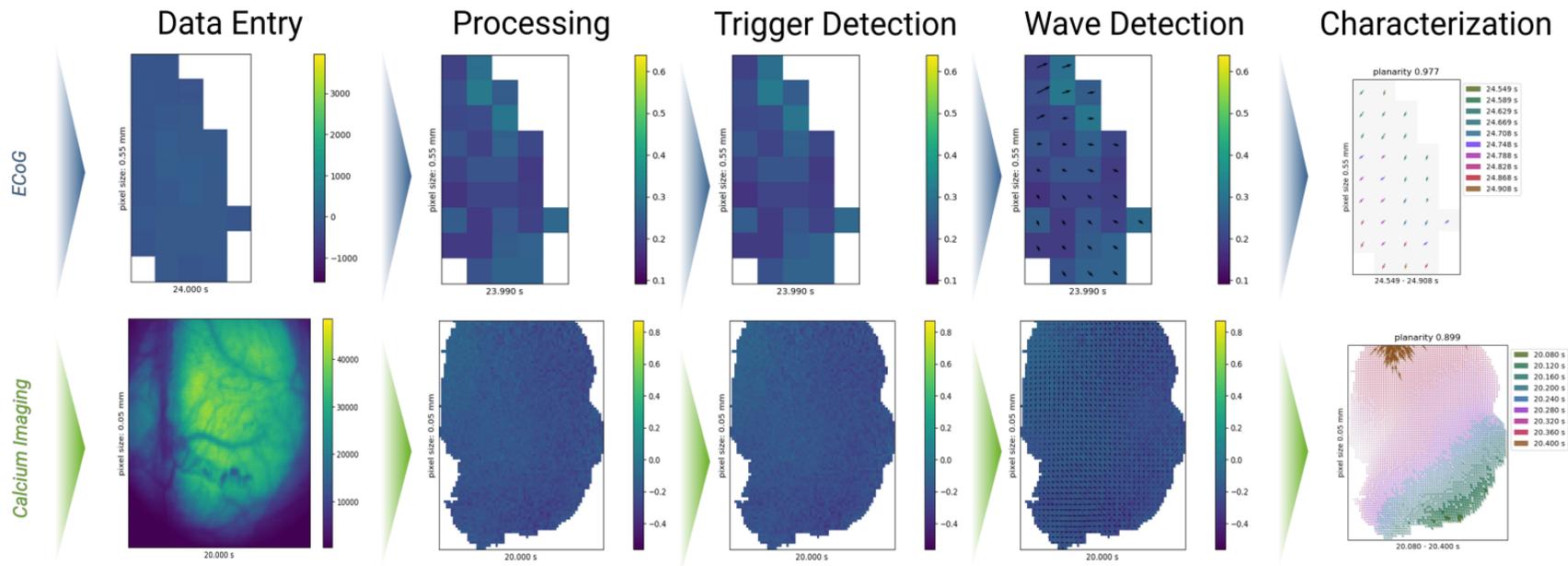


## Characterization

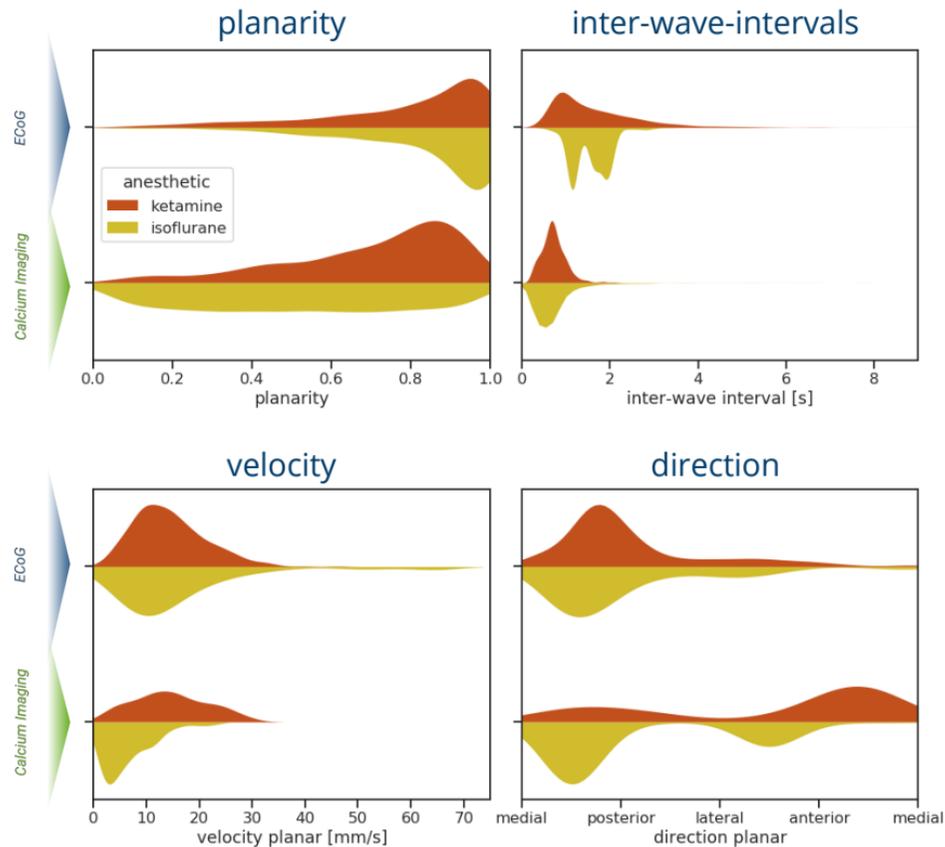


# The Slow Wave Analysis Pipeline

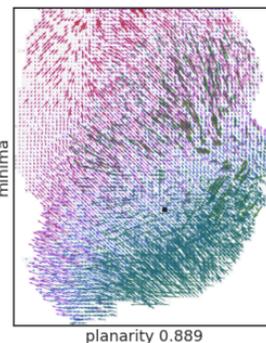
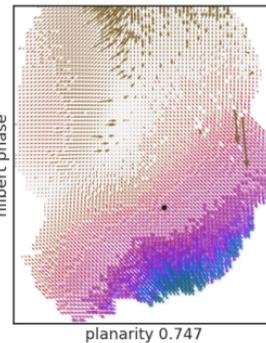
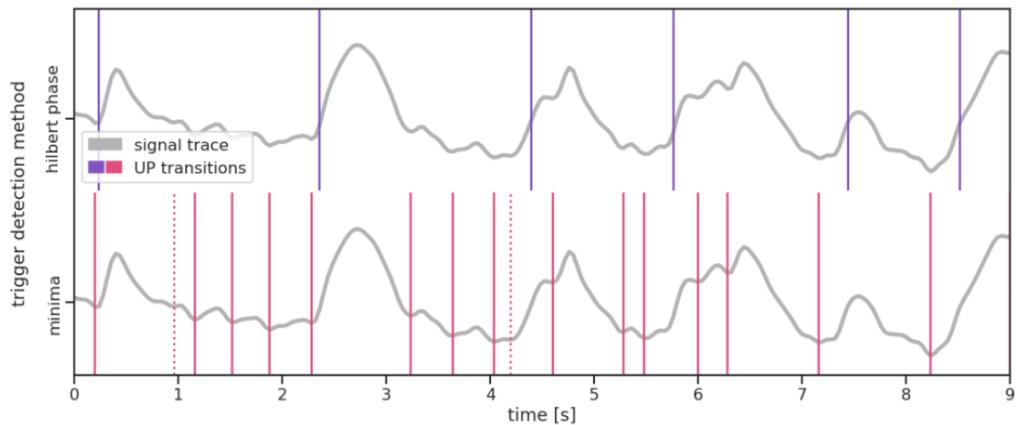
enables meta-studies,  
for which, we analysed 5 open-access datasets  
of 60 ECoG and calcium imaging recordings.



# Comparing Heterogeneous Data

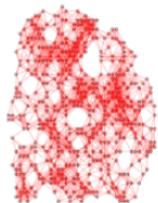


# Comparing Methods on Same Data

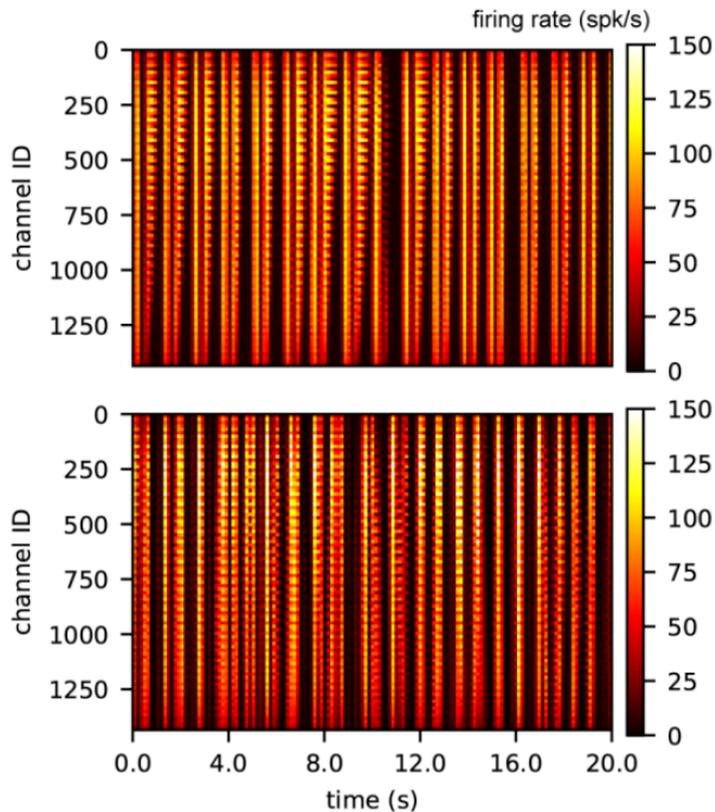
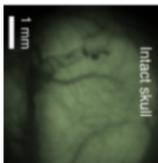


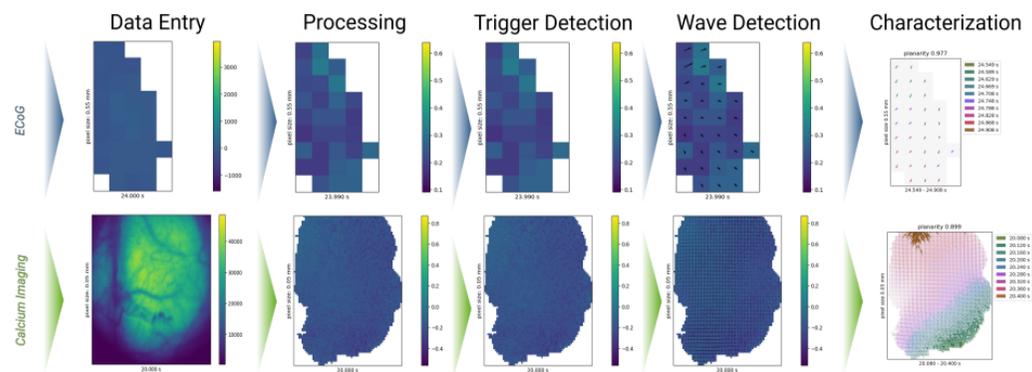
# Calibrating & Validating Models

Simulation



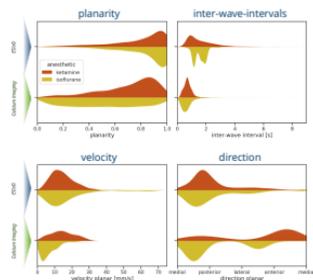
Data



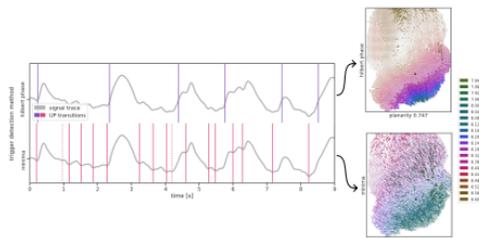


## Modular Wave Analysis Pipeline

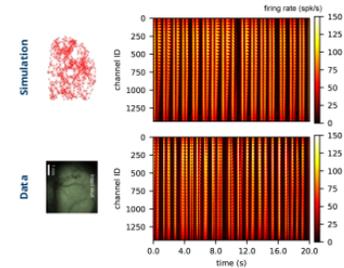
### Comparing Heterogeneous Data



### Comparing Methods on Same Data



### Calibrating & Validating Models



# Acknowledgments

for more information:



[http://go.fzj.de/wave\\_analysis\\_pipeline](http://go.fzj.de/wave_analysis_pipeline)

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# Appendix

## References for Figure on Slide 1

- a) Chan et al. (2015) doi:10.1038/ncomms8738
- b) Celotto et al. (2020) doi:10.3390/mps3010014
- c) Stroh et al. (2013) doi:10.1016/j.neuron.2013.01.031
- d) Pastorelli et al. (2019) doi:10.3389/fnsys.2019.00033
- e) Bazhenov et al. (2002) doi:10.1523/JNEUROSCI.22-19-08691.2002
- f) Keane & Gong (2015) doi:10.1523/JNEUROSCI.1669-14.2015
- g) Capone et al. (2017) doi:10.1093/cercor/bhx326
- h) Massimini et al (2004) doi:10.1523/JNEUROSCI.1318-04.2004
- i) Muller et al. (2016) e17267. doi:10.7554/eLife.17267
- j) Nir et al. (2011) doi:10.1016/j.neuron.2011.02.043
- k) Botella-Soler et al. (2012) doi:10.1371/journal.pone.0030757

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## Datasets

- Resta et al. (2020) doi:10.25493/3E6Y-E8G
- Resta et al. (2020) doi:10.25493/XJR8-QCA
- Sanchez-Vives (2020) doi:10.25493/WKA8-Q4T
- Sanchez-Vives (2019) doi:10.25493/ANF9-EG3
- Sanchez-Vives (2019) doi:10.25493/DZWT-1T8