# AUTOMATED ASSESSMENT AND COMPARISON OF CORTICAL NEURON MODELS

Justas Birgiolas, Russell Jarvis, Vergil R. Haynes, Richard C. Gerkin, Sharon M. Crook

Laboratory for Informatics and Computation in Open Neuroscience

Arizona State University



# WHAT CAN WE LEARN FROM THE LANDSCAPE OF DATA-DRIVEN MODELS OF CORTICAL NEURONS?

#### MOTIVATION

**NeuroML-DB** [2] catalogues over 1,500 published models obtained in NeuroML format from Open Source Brain [5]. Complementing OSB, NeuroML-DB provides systematic characterizations of model complexity, electrophysiology, and morphology, making it easy to find, evaluate, and reuse models and their components.



#### **WHICH FEATURES?**



Electrical behavior of neuron types according to the Petilla terminology [1]. (Image from [8].) Druckmann et al. features below [3].

#### c: continuous d: delayed b: bursting

AC: accommodating NAC: non-accommodating STUT: stuttering IR: irregular

## **CHANNEL MECHANISMS**

We investigate different channel types responsible for neuron model behaviors. Top: Projected clusters of cortical neuron models in feature space. Bottom: Normalized maximal conductance for slow inactivating K+ channel in soma for same models.



# How well do models cover the diversity of experimentally recorded neurons?

Which features maximally differentiate among neuron/model types and their dynamical behaviors?

Are there models of neurons and channels that are outliers when compared to each other and to cortical neuron electrophysiology data?

#### **MODELING METHODS**

We focus on models described in *NeuroML* – a simulator independent, modular, multiscale model description language [4]. This standardized format allows for automated model analysis.



Drop in AP amplitude (amp.) from first to second spike (mV)	Average accommodation at steady-state (%)	Percent change in AP width at half height, first to second spike (%)
AP amplitude change from first spike to steady-state (mV)	Average rate of accommodation during steady-state	Percent change in AP peak to trough rate of change, first to second spike (%)
AP 1 amplitude (mV)	Average inter-spike interval (ISI) coefficient of variation (CV) (unit less)	Percent change in AP fast AHP depth, first to second spike (%)
AP 1 width at half height (ms)	Median of the distribution of ISIs (ms)	Input resistance for steady-state current (Ohm)
AP 1 peak to trough time (ms)	Average change in ISIs during a burst (%)	Average delay to AP 1 (ms)
AP 1 peak to trough rate of change (mV/ms)		SD of delay to AP 1 (ms)
AP 1 Fast AHP depth (mV)	Average rate, suong sumutus (nz)	Average delay to AP 2 (ms)
AP 2 amplitude (mV)	Average delay to AP 1, strong stimulus (ms)	SD of delay to AP 2 (ms)
AP 2 width at half height (ms)	SD of delay to AP 1, strong stimulus (ms)	Average initial burst interval (ms)
AP 2 peak to trough time (ms)	Average delay to AP 2, strong stimulus (ms)	SD of average initial burst interval (ms)
AP 2 peak to trough rate of change (mV/ms)	SD of delay to AP 2, strong stimulus (ms)	Average initial accommodation (%)
AP 2 Fast AHP depth (mV)	Average initial burst ISI, strong stimulus (ms)	Average steady-state accommodation (%)
Percent change in AP amplitude first to second spike (%)	SD of average initial burst ISI, strong stimulus (ms)	Rate of accommodation to steady-state (1/ms)

### WHAT CAN WE LEARN?

Below: Comparisons of open data and cortical neuron model outputs for pairs of features (both from Druckmann et al. [3] and the Allen SDK [7]. There is good agreement between models and data but outliers could reveal specific experimental behaviors that are not captured well by models or models that do not fit data well.



Agglomerative clustering of model channel dynamics for multiple simulation protocols reveals channels that do not behave like other channels of that type.







#### Call T Type Tra Call T Type Tra Call T Type Tra Call H Call H

#### **DIY MODEL ANALYSIS!**

NeuroML-DB data are available via an *API*, which returns a JSON object in response to a URL. See documentation at https://neuroml-db.org/api and links to browser add-ons and examples for working with JSON URLs in programming languages like Python, R, and MATLAB.

#### Code Available at:

https://github.com/vrhaynes/LargeScaleModelAnalysis\_2019 Interactive Clustering Visualization Available at: https://iconlab.asu.edu/barcelona

## **MODEL/DATA COMPARISONS**



http://dash.scidash.org

**NeuronUnit** [9] provides a way to compare specific models to data using "unit tests". Results are shared at the **SciDash** dashboard. Feature extraction routines used here are providing the basis of novel unit tests for further work.

Allen Institute for Brain Science, Allen Cell Types Database [7] can be accessed using the SDK. http://celltypes.brain-map.org

Large-scale Model	Blue Brain Project (1035)	Allen Institute (170)
Publication	Markram et al. (2015) [8]	Gouwens et al. (2018) [6]
Exp. Protocol	Druckmann et al. (2012) [3]	Allen Cell Types Database Protocol
Features	38 Druckmann Features	12 Features



#### Hierarchical clustering of models with a density-based clustering method (HDBSCAN) on PCA-reduced features. Factor loading reveals features associated with top three components as shown directly above. Clusters correspond well to assigned electrical types, but outliers exist at different levels and will be the subject of further investigation.

#### REFERENCES

- 1. Ascoli GA, et al. (2008) Nature Reviews Neuroscience. 9(7):557.
- Birgiolas J, et al. (2015) In Amarnath Gupta and Susan Rathbun, editors. Proceedings of the 27<sup>th</sup>International Conference on Scientific and Statistical Database Management. New York, NY: ACM. 37.
- 3. Druckmann S, et al. (2012) Cereb. Cortex 23:2994–3006.
- 4. Gleeson P, et al. (2010) PLoS Comput Biol. 6, e1000815.
- 5. Gleeson P, et al. (2019) Neuron. doi:10.1016/j.neuron.2019.05.019.
- 6. Gouwens N, et al. (2018) Nature Comm. doi:10.1038/s41467-017-02718-3.
- 7. Hawrylycz M, et al. (2016) PNAS. 113(27):7337-44.
- 8. Markram H, et al. (2015) Cell 163:456-492.
- 9. Omar C, et al. (2014) In Companion Proceedings of the 36th International Conference on Software Engineering. ACM. 524–527.

### ACKNOWLEDGEMENTS

This work was supported in part by grant R01EB021711 from the National Institute of Biomedical Imaging and Bioengineering and by grant R01MH106674 from the National Institutes of Mental Health.

