



# Data-driven model of auditory thalamocortical system rhythms

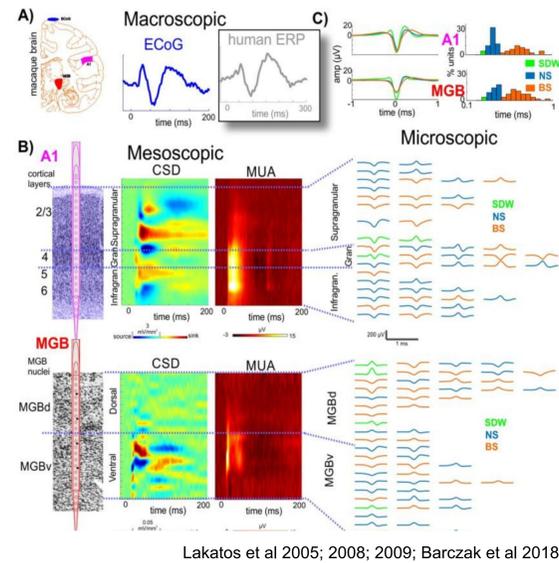
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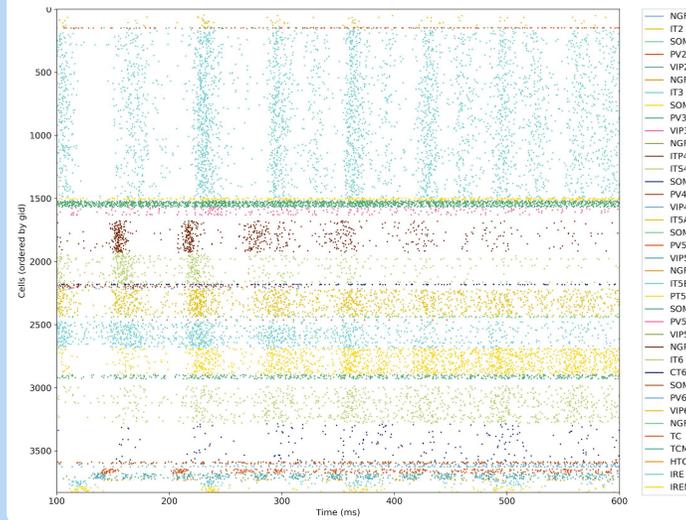
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## Summary

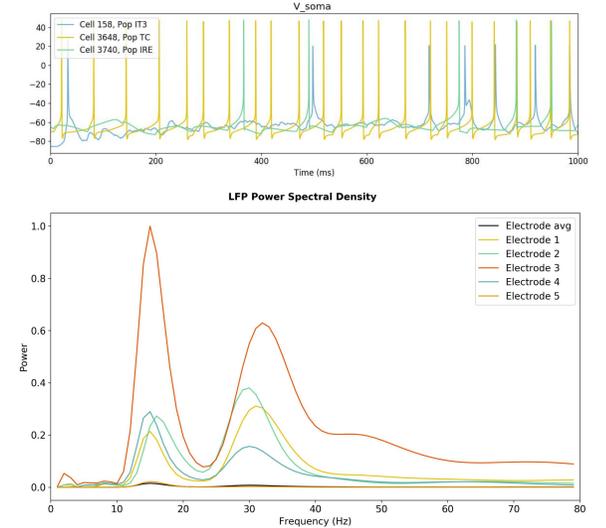
- **Detailed biophysical computer model** of the auditory thalamocortical system, using the NEURON simulator and the NetPyNE tool, with 39 neural populations consisting of multiple cell types (7 excitatory types, 5 interneuron types) distributed across the 6 A1 cortical layers and 2 thalamic regions (MGB and RTN).
- **Layer boundaries**, neuronal **densities** and cell type **distributions** per layer were based on macaque and rodent thalamocortical data.
- **Model neurons** are multicompartment, conductance-based and include multiple ion channels, with parameters optimized to reproduce physiological responses.
- **Connectivity** depended on pre- and post-synaptic cell type and layer and was derived from cortical data from macaque and rodents.
- Model will be refined using data analyses from multisite recordings, and used to **explore** the mechanistic origins of the spatiotemporal neuronal **oscillatory patterns** observed *in vivo*.
- Model will make **predictions** on the proposed roles of neuronal oscillations in auditory information processing, which will be **tested** via targeted deep brain electrical microstimulation and pharmacological manipulations.



## Preliminary simulation results



Model scaled down to 7.5% cell density: 4k neurons; 1M synapses



## Data-driven cell models

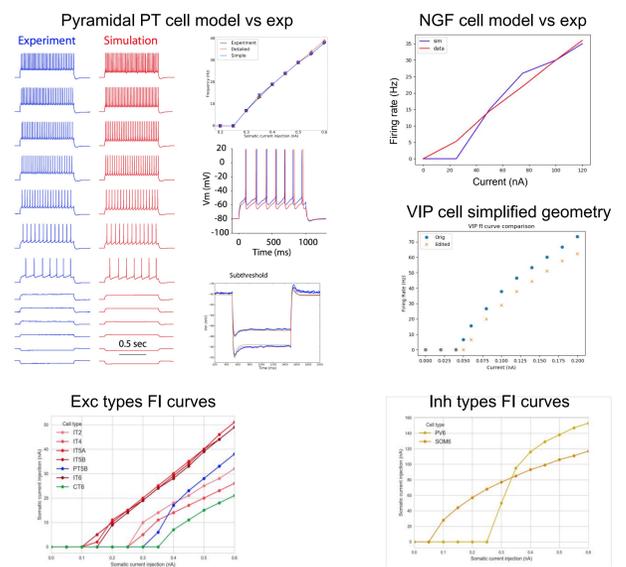
Cell type models developed were multi-compartment and conductance-based.

**Excitatory** cell types: intratelencephalic (IT), pyramidal tract (PT), spiny stellate (ITS), corticothalamic (CT) and MGB thalamocortical (TC).

**Inhibitory** cell types: somatostatin (SOM), parvalbumin (PV), neurogliaform (NGF), vasoactive intestinal peptide (VIP), and reticular nucleus (RT).

**Geometry:** Simplified morphologies (1 - 6 compartments). Dendritic lengths were sized to match the macaque cortex dimensions.

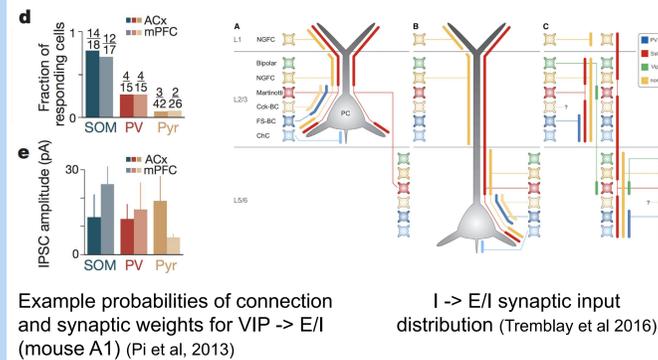
**Parameter Fitting:** Cell types in each layer were fitted to macaque (if available) or rodent electrophysiology data. This was done via either multi-objective evolutionary optimization algorithm or via hand-tuning. Passive parameters (e.g. leak channel conductance) were tuned to fit RMP and other features of subthreshold traces, including steady state voltage and sag. Active parameters (e.g. fast Na<sup>+</sup> channel density) were then tuned to fit features like firing rate curves and adaptation.



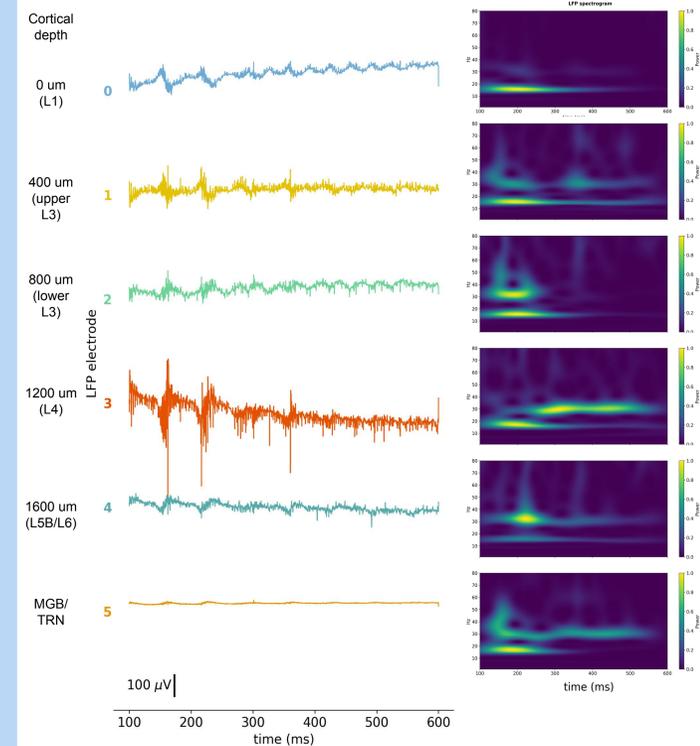
## Data-driven local microcircuit connectivity

• E -> E/I from mouse V1,S1,A1,M1 (Levy & Reyes 2012; Yoshi et al 2015; Billeh et al 2019; Lefort et al 2009; Allen Brain Institute)

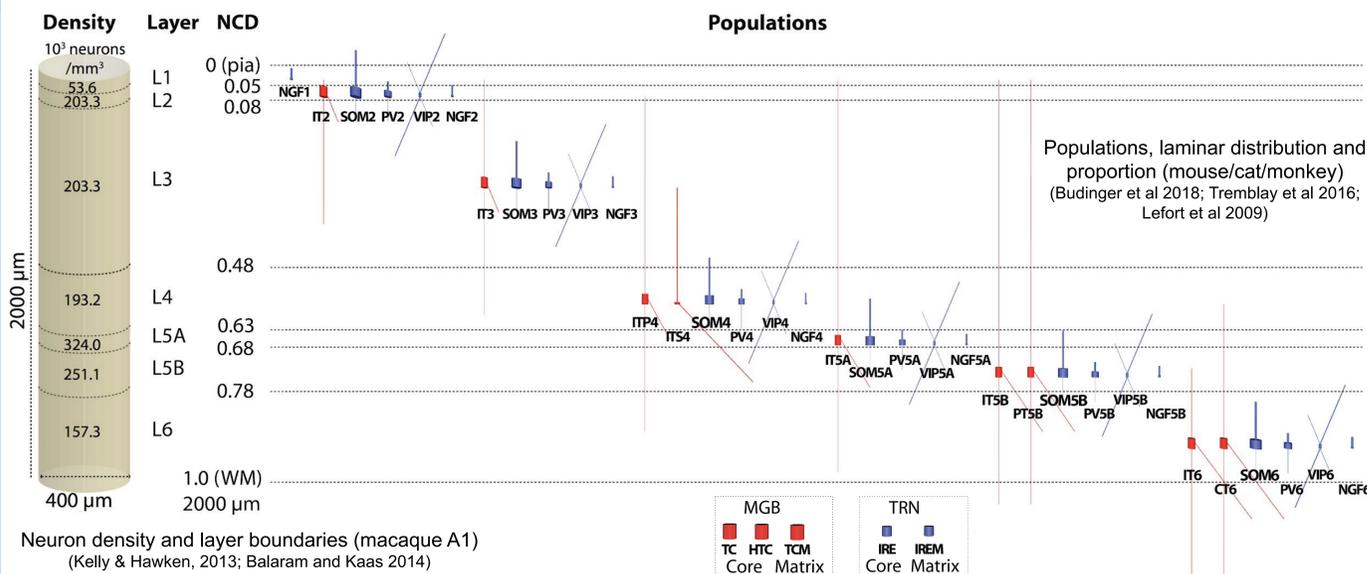
• I -> E/I from mouse V1,S1,A1,M1 (Sohn et al 2016; Naka et al 2016; Tremblay et al 2016; Pi et al 2013; Pi et al 2013)



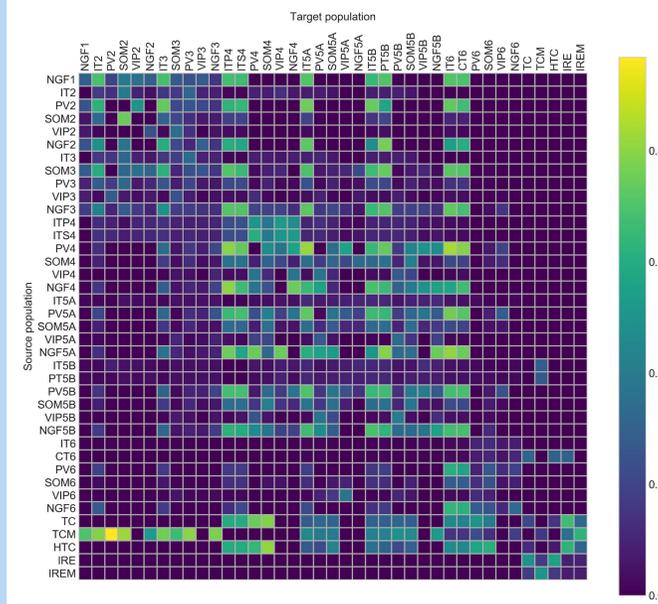
## Local Field Potential (LFP) recorded at different depths



## Data-driven network structure



## Connectivity Matrix



## Modeling software tool

Dura-Bernal S, Suter B, Gleeson P, Cantarelli M, ... , Neymotin SA, McDougal R, Hines M, Shepherd GMG, Lytton WW. (2019) **NetPyNE: a tool for data-driven multiscale modeling of brain circuits.** *eLife* 2019;8:e44494 ([www.netpyne.org](http://www.netpyne.org))



## Funding

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