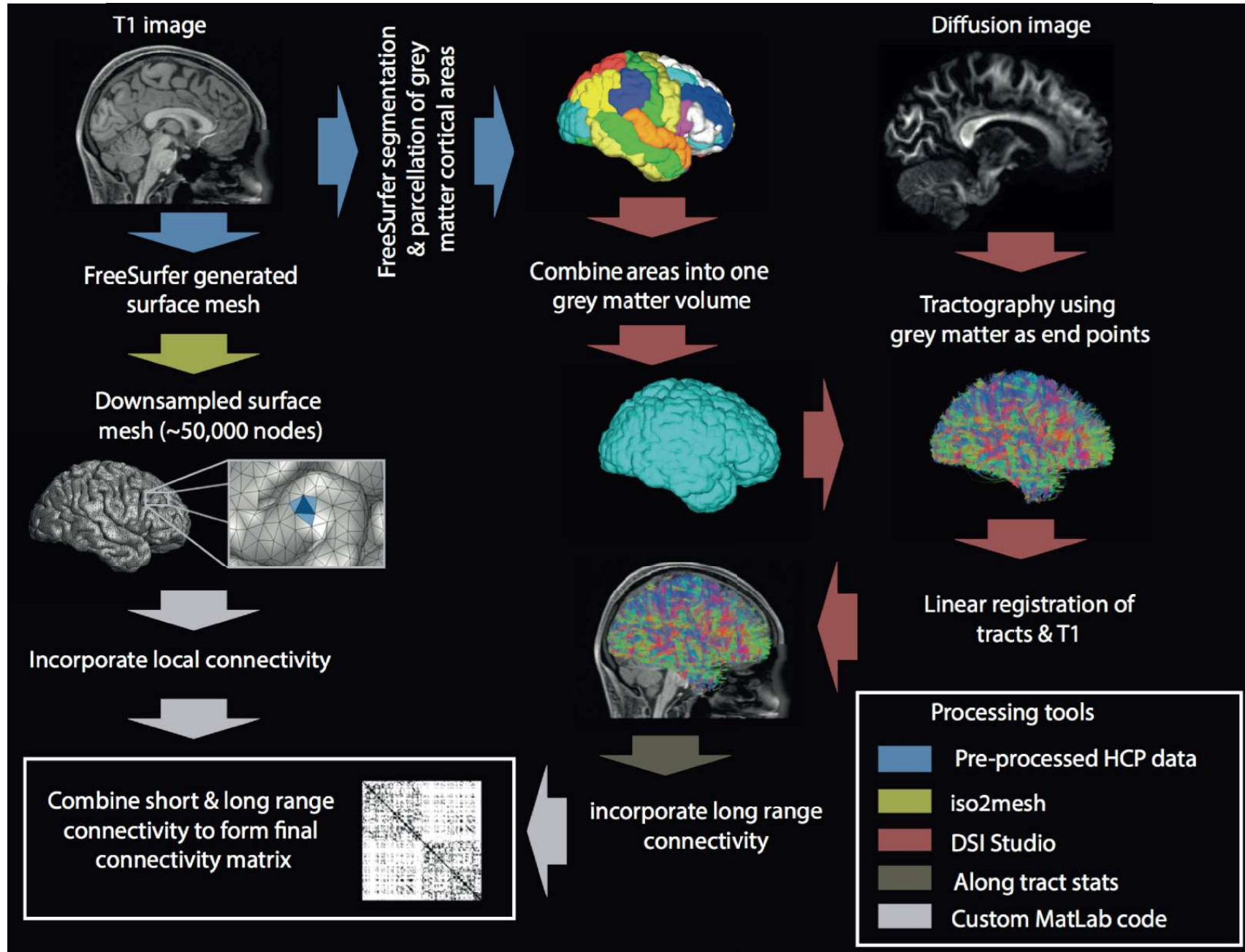
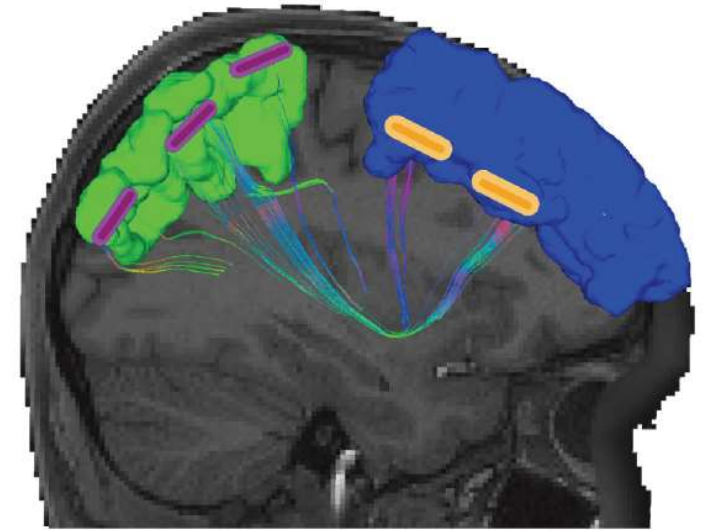
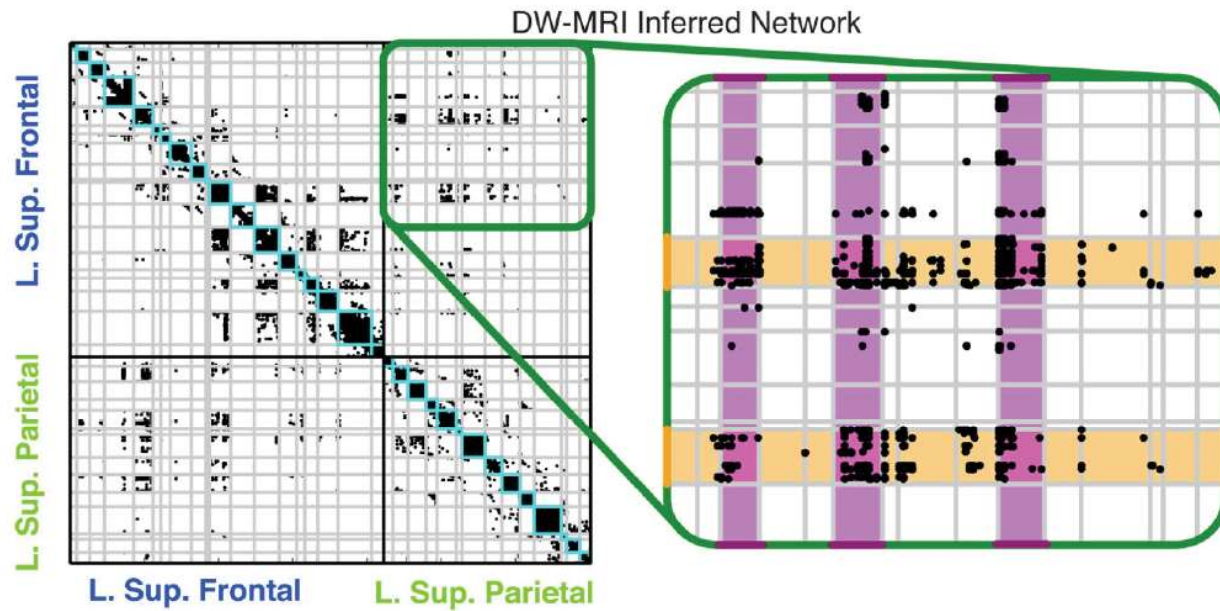


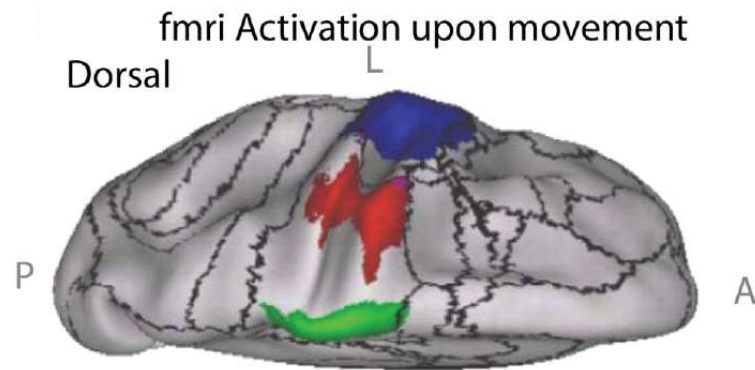
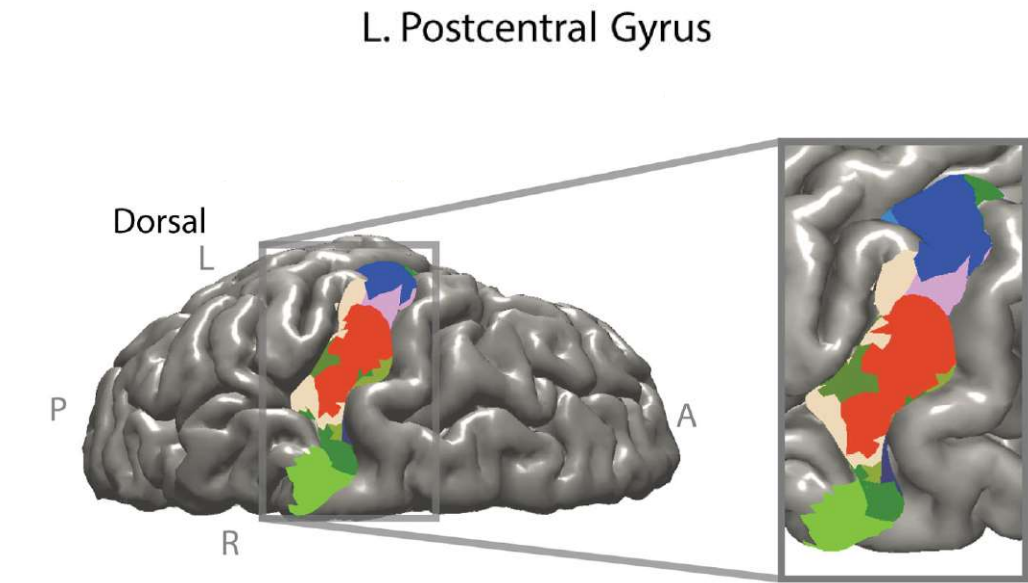
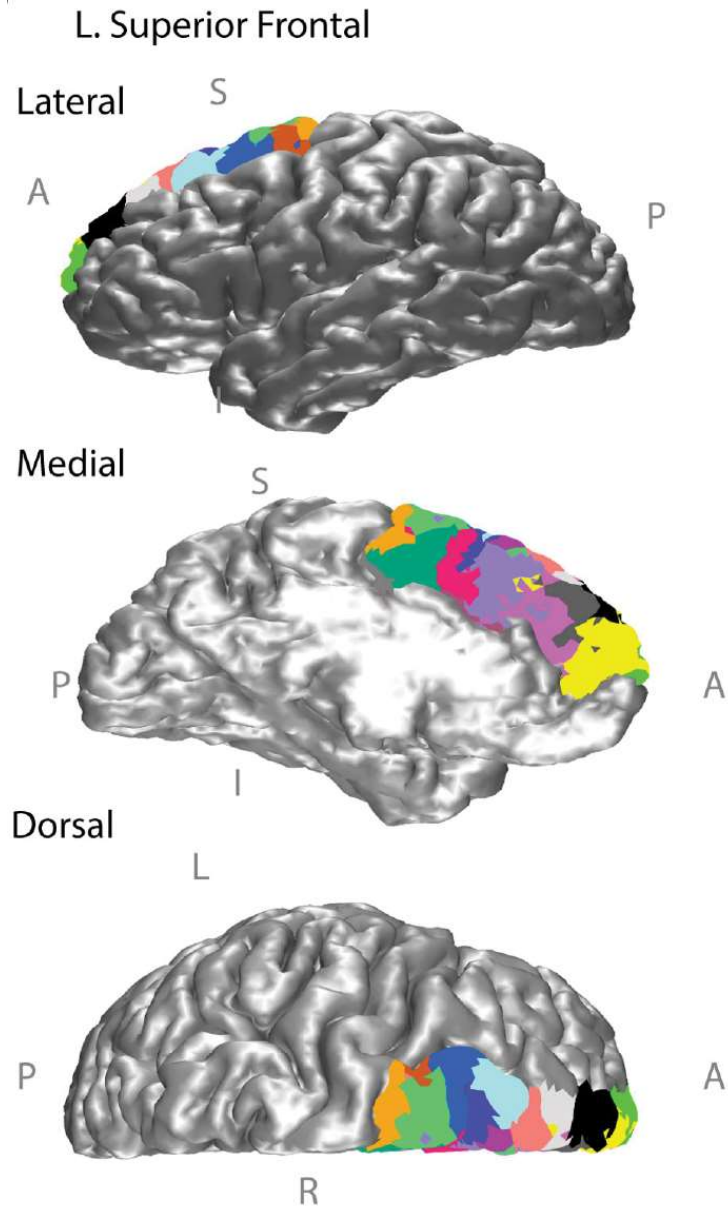
Super-resolution DTI: 50,000 nodes



Super-resolution DTI: modules within brain regions



Super-resolution DTI: spatial (and functional?) modules

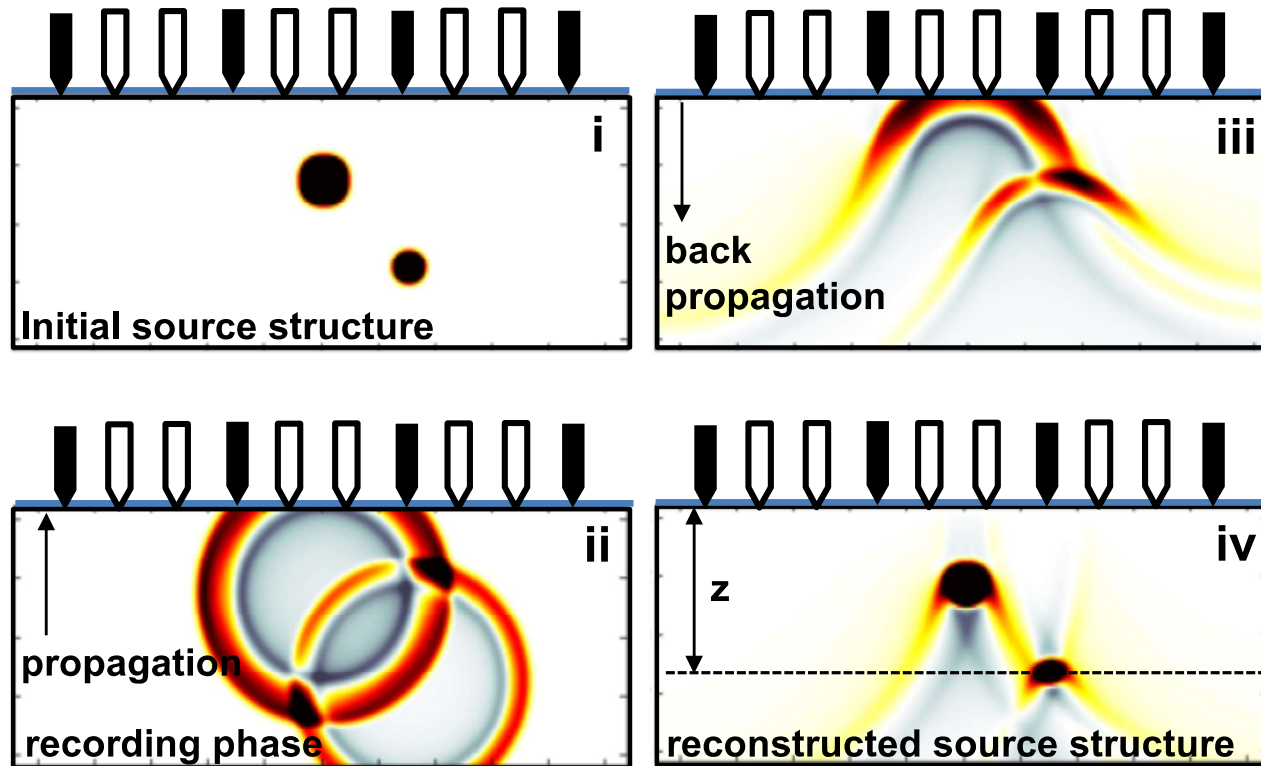


- Tongue
- Hand
- Foot

Yeo et al., *J. Neurophysiol.* 2011

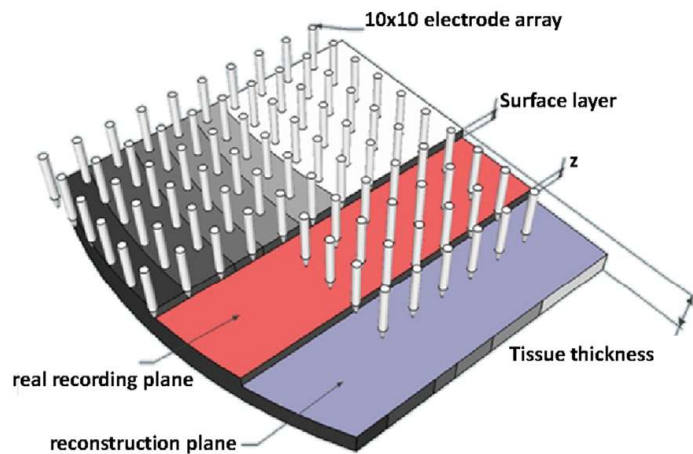
Taylor et al., *Scientific Reports*, 2017

Super-resolution imaging: near-field holography

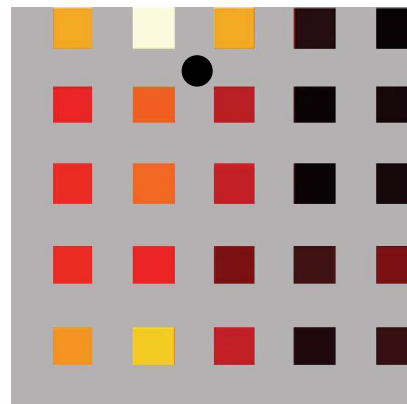


Kjeldsen, Kaiser, Whittington. *Journal of Neuroscience Methods*, 2015

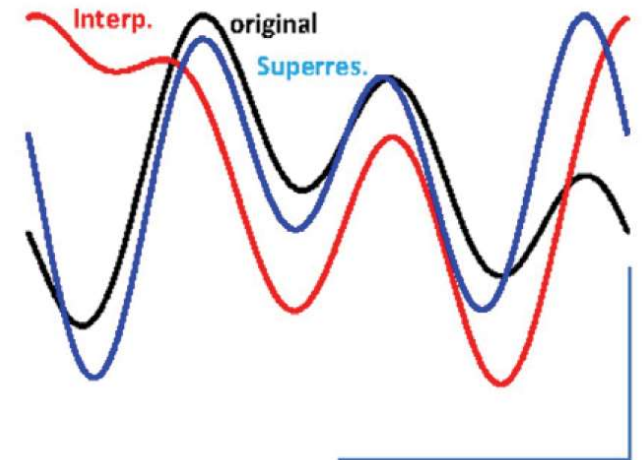
Super-resolution MEA: measuring delta rhythm activity



Signal reconstruction away from the recording plane



Compare real 10x10 data with super-resolution data based on down-sampled 5x5 data



Delta rhythm: semblance to original source signal

Kjeldsen, Kaiser, Whittington. *Journal of Neuroscience Methods*, 2015

Super-resolution EEG: Energy flow during processing



**Dynamic Energy Flow
Super-Resolution EEG
Change how you see the Brain**



Henrik Kjeldsen, Truust Neuroimaging (patent pending)

Computational models of effects and side effects

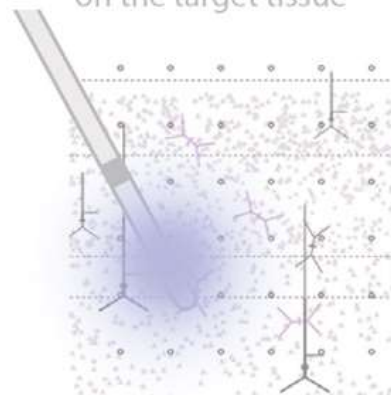
Not necessarily the ones that are most important for (altered) brain function

targeting involved nodes might lead to huge side effects

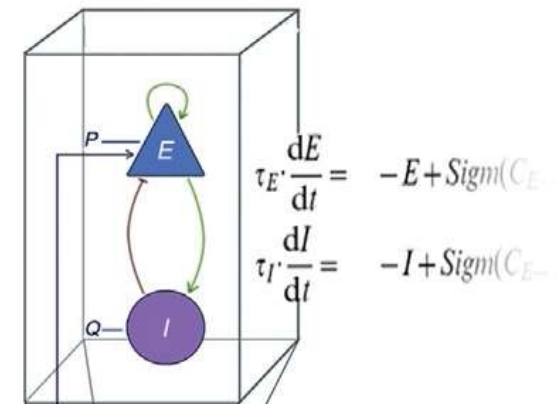
Magnetic resonance imaging (MRI)



Simulating the effect of the stimulus on the target tissue



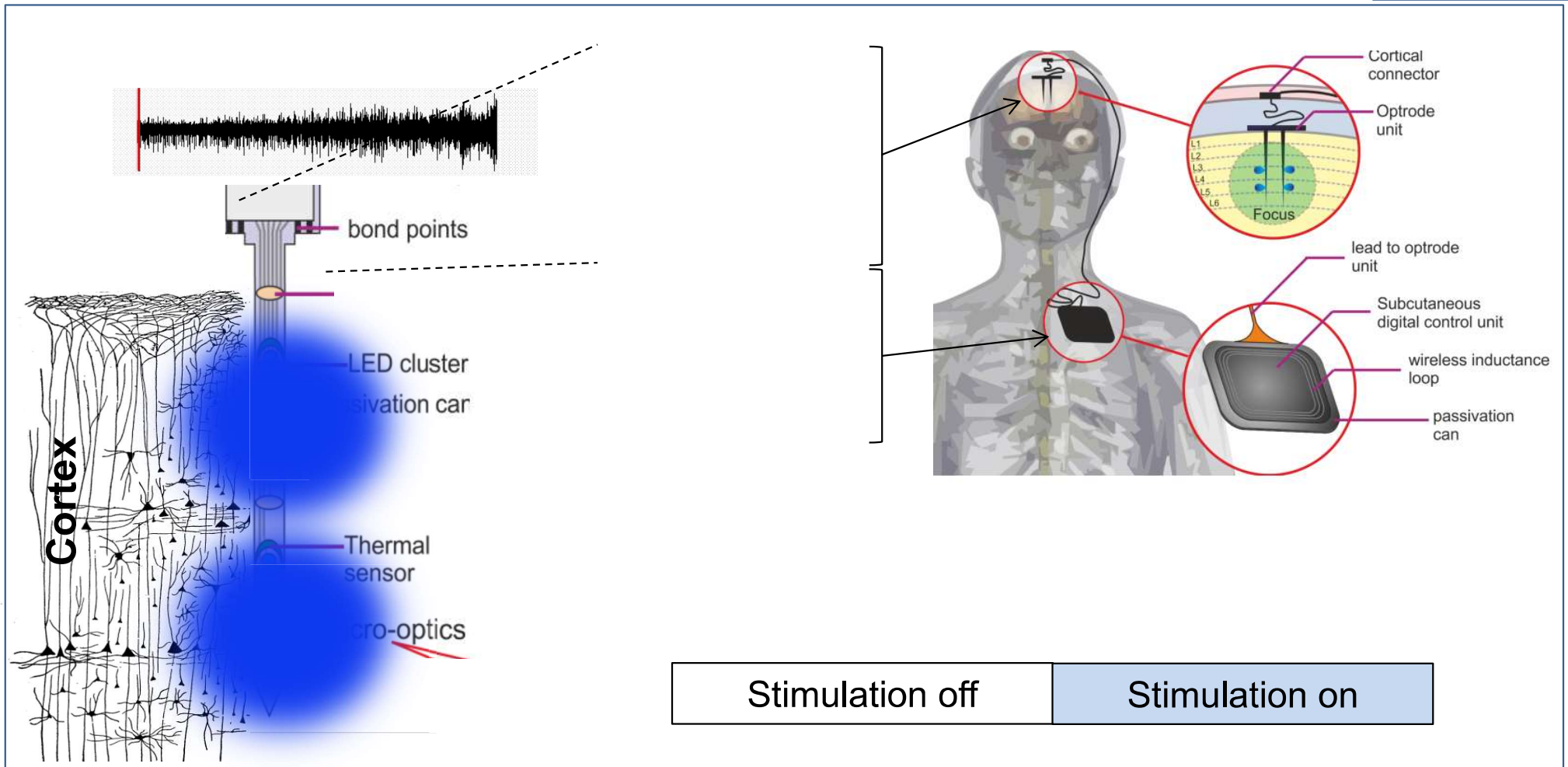
Simulating brain dynamics



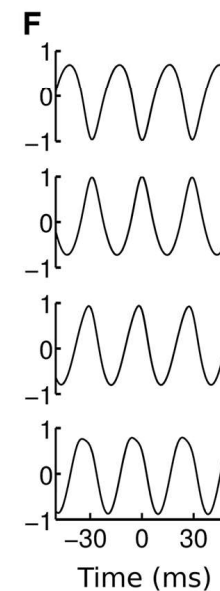
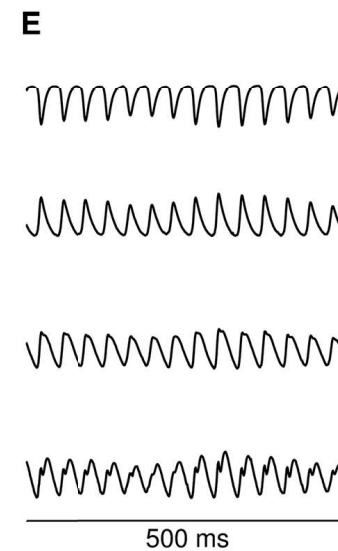
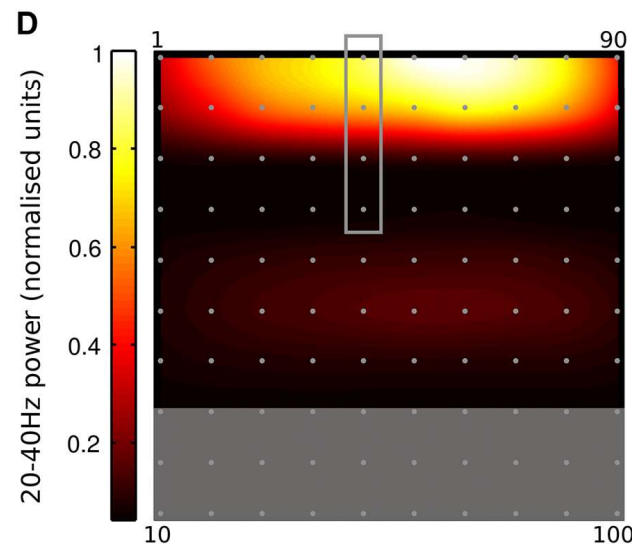
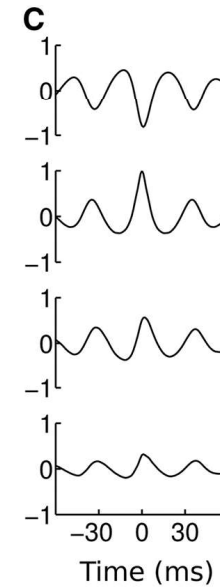
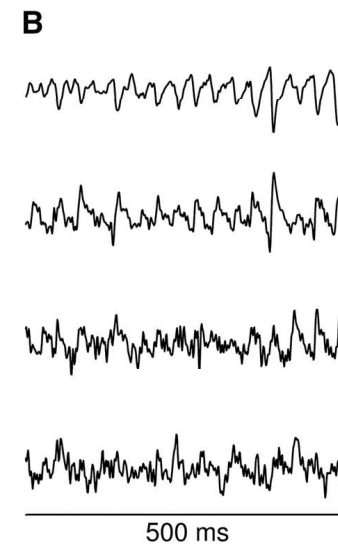
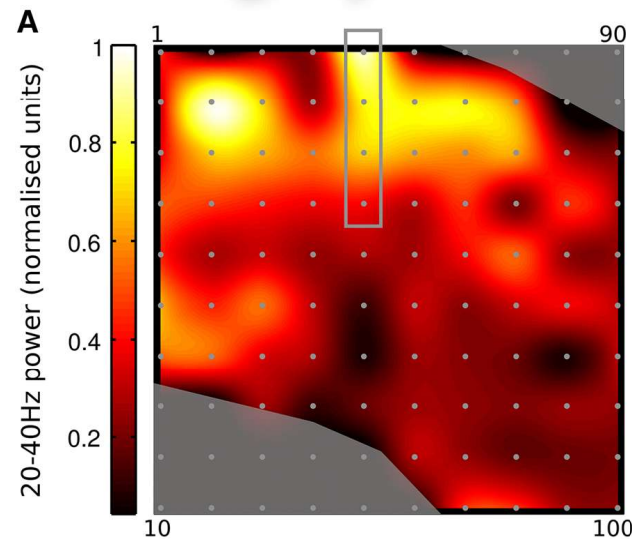
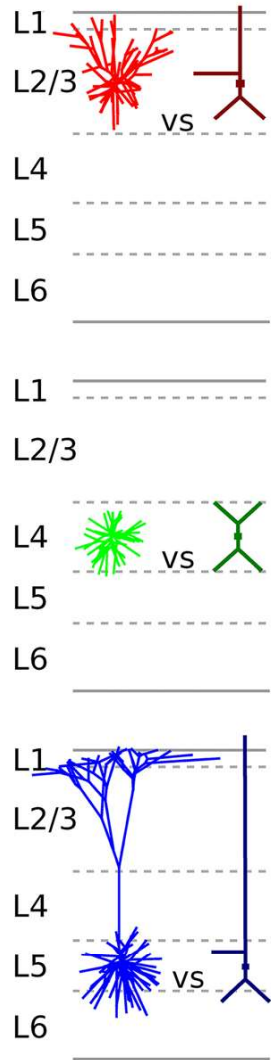
Computer simulations – predicting the effect of optogenetic stimulation

Controlling Abnormal Network Dynamics with Optogenetics (CANDO)

7yrs (till 2021), £10m www.cando.ac.uk



VERTEX: simulating dynamics within cortical columns

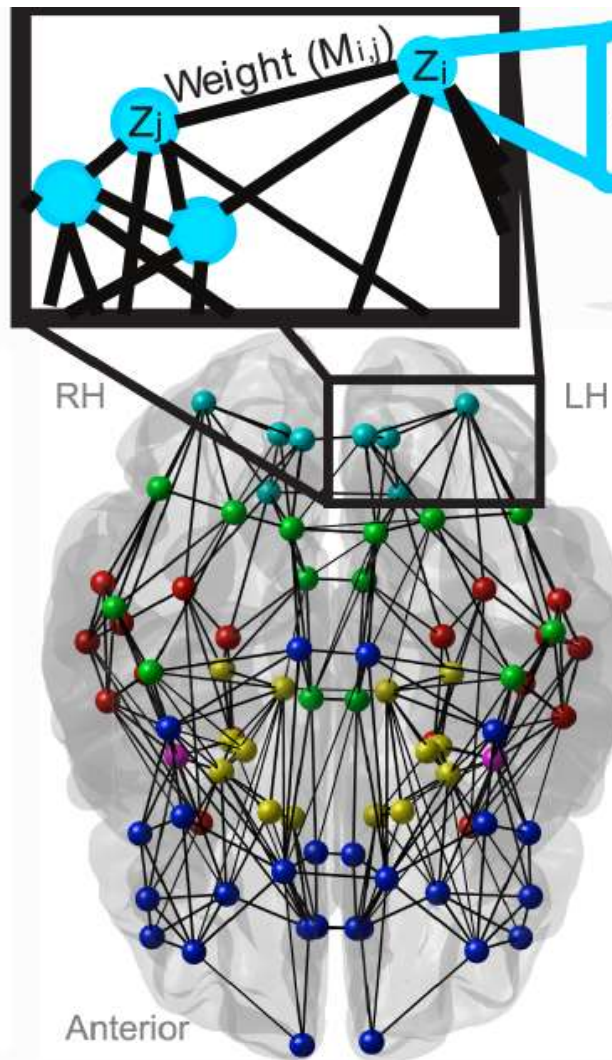


<http://www.vertexsimulator.org/>

Tomsett et al. *Brain Structure and Function*, 2015

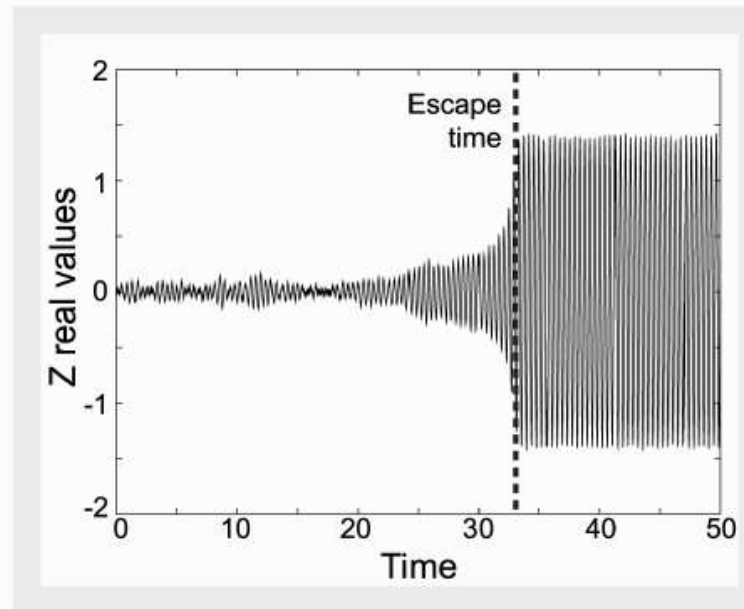
Roopun et al. *PNAS*, 2010

Computer simulations – predicting the location of epileptic tissue



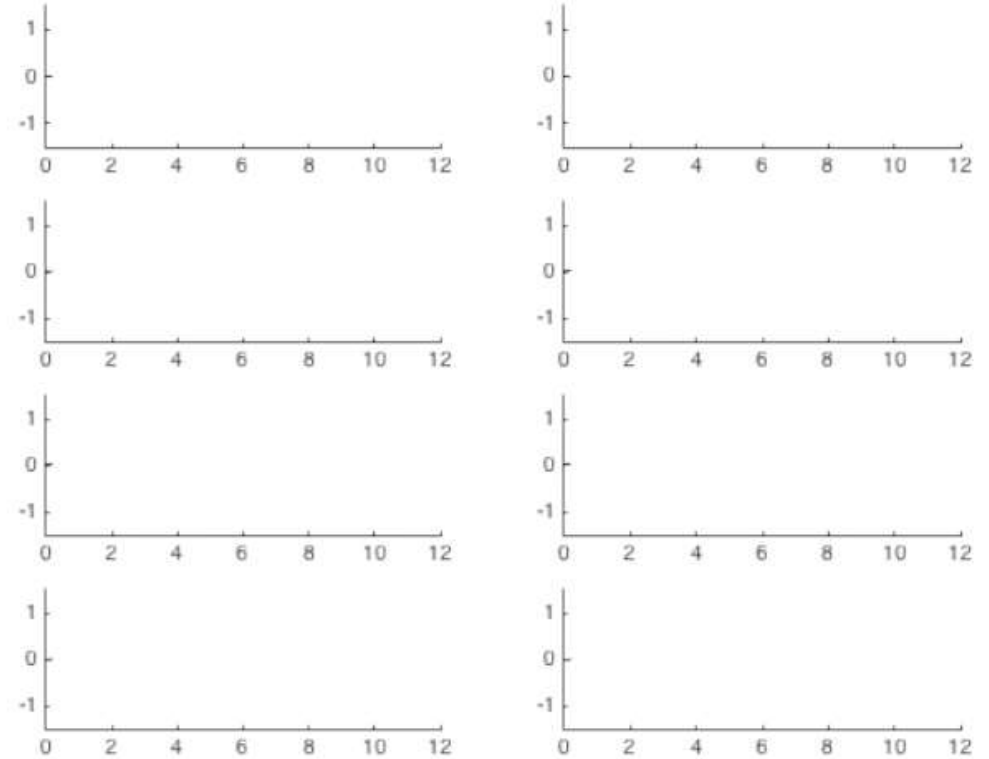
$$dz_i(t) = \left(f(z_i) + \beta \sum_{j \neq i} M_{ji} (z_j(t - \tau_{ij}) - z_i(t)) \right) dt + \alpha dw_i(t)$$

Model Simulation



Hutchings et al. *PLOS Computational Biology*, 2015

Computer simulations – predicting the location of epileptic tissue



Hutchings et al. *PLoS Computational Biology*, 2015

Computer simulations – predicting epilepsy surgery success

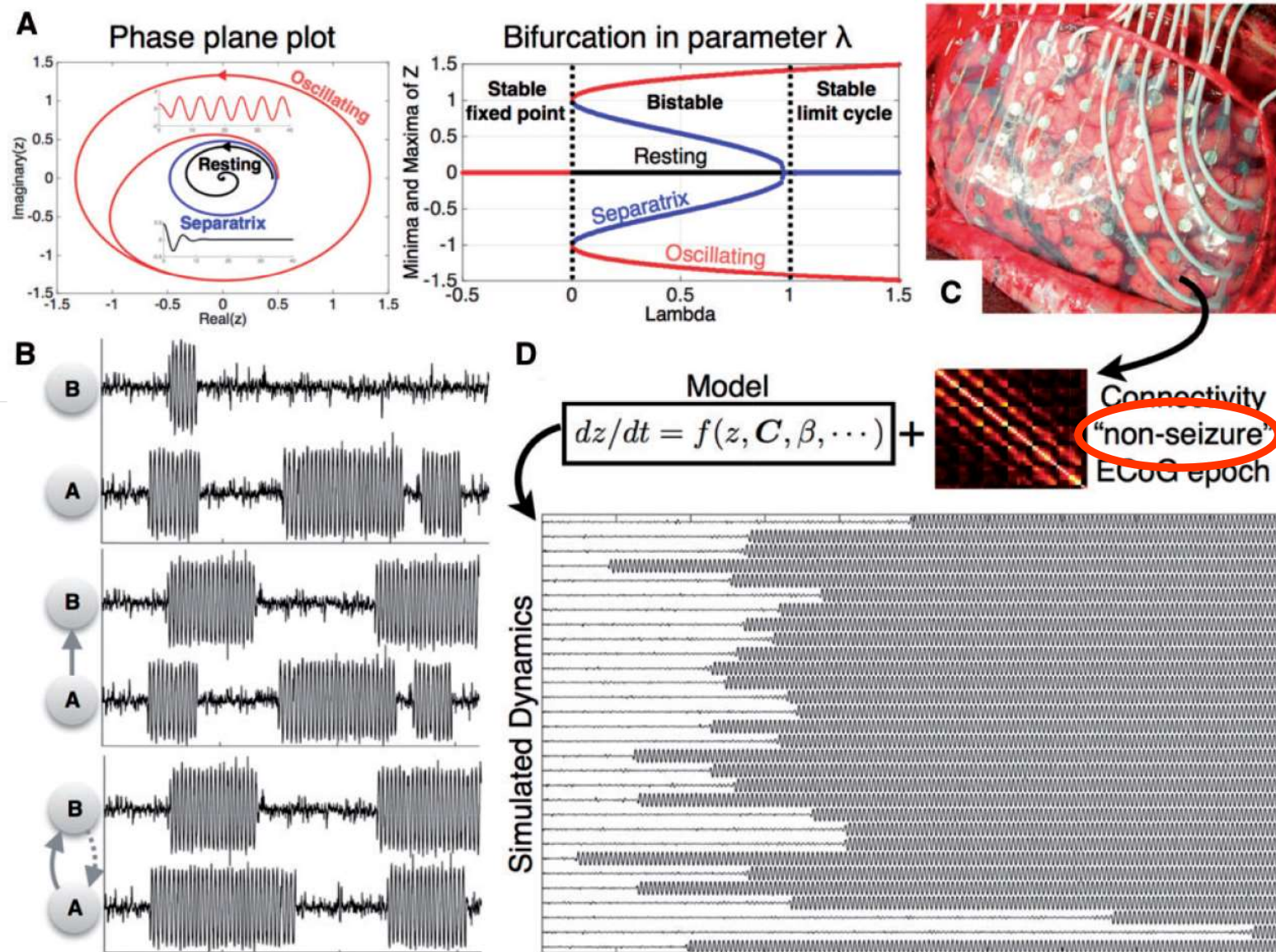
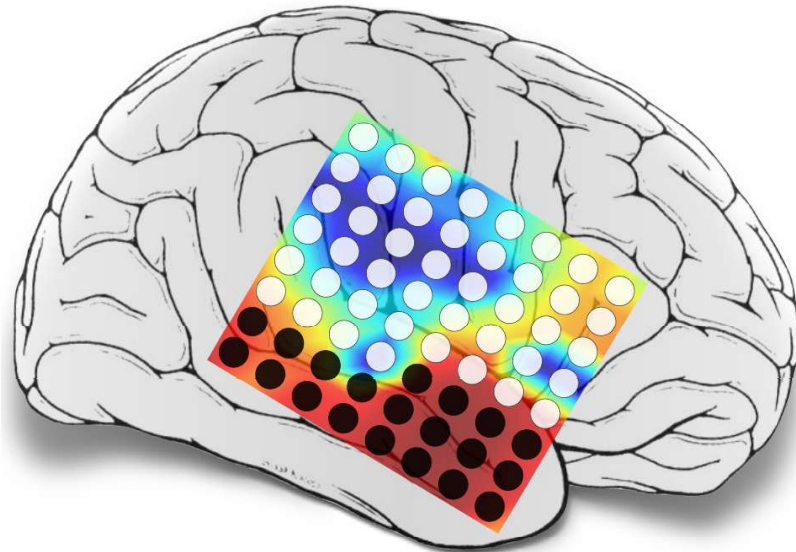


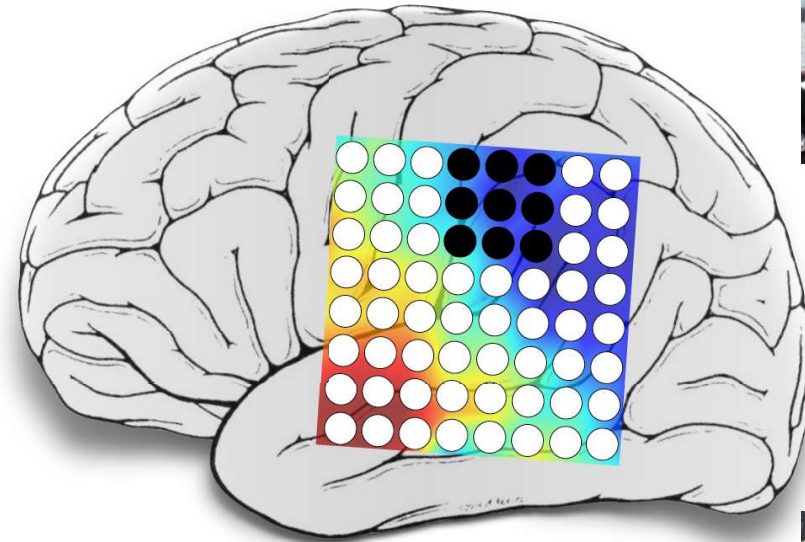
Figure 1 Illustration of model dynamics. (A) Deterministic dynamics of a single node representing the bi-stability of the model. (B) Stochastic dynamics in a two node network. The two nodes are initially disconnected having independent dynamics. Depending on the strength and direction of connections, the dynamics of each node is influenced by the other. (C and D) Patient-specific connectivity matrix is obtained from intracranial, interictal ECoG recording, which is incorporated as a model parameter to simulate the model dynamics.

Computer simulations – predicting epilepsy surgery success

P1: Seizure free outcome



P2: Not Seizure free outcome



● Location of surgical resection

Low Simulated Seizure likelihood High

| Prediction | | Real outcome |
|------------------|---|-----------------------|
| seizure-free | → | 70% seizure-free |
| | → | 30% |
| not seizure-free | → | 100% not seizure-free |

Sinha et al. *Brain*, 2016

Summary

- Diagnosis for individual patients including aetiology (developmental origin) and disease subtype
- Identification of potential treatment targets
- Model for effects *and side effects* of treatment

All three components are needed to lead to new alternative treatments for brain network disorders

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PostDoc



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Now **faculty**



Yujiang Wang
Now **faculty**



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PhD student



Frances Hutchings
PhD student



Chris Hayward
PhD student



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Chris Thornton
PhD student



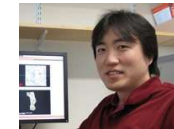
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<http://www.dynamic-connectome.org>



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