



Translational Neuromodeling Unit



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Swiss Federal Institute of Technology Zurich

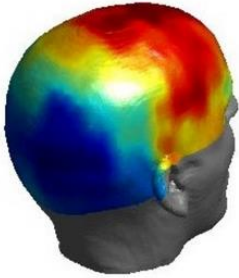
DYNAMIC CAUSAL MODELING FOR EEG

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TRANSLATIONAL NEUROMODELING UNIT (TNU)
UNIVERSITY OF ZURICH & ETH ZURICH

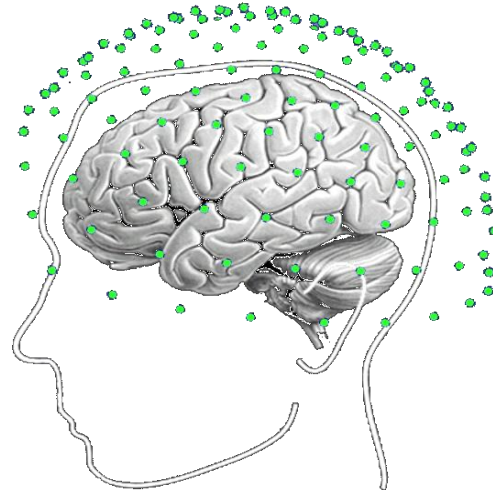
Computational Psychiatry Course 2018

EEG, MEG

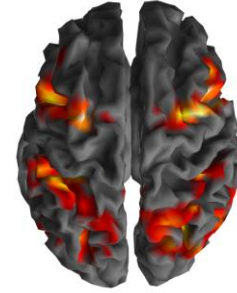


Forward model:
Predicting measured activity

$$y = g(x, \theta) + \varepsilon$$



fMRI



Model inversion:
Estimating neuronal
mechanisms

$$\frac{dx}{dt} = f(x, u, \theta)$$

Friston et al., 2003, *NeuroImage*; David et al., 2006, *NeuroImage*



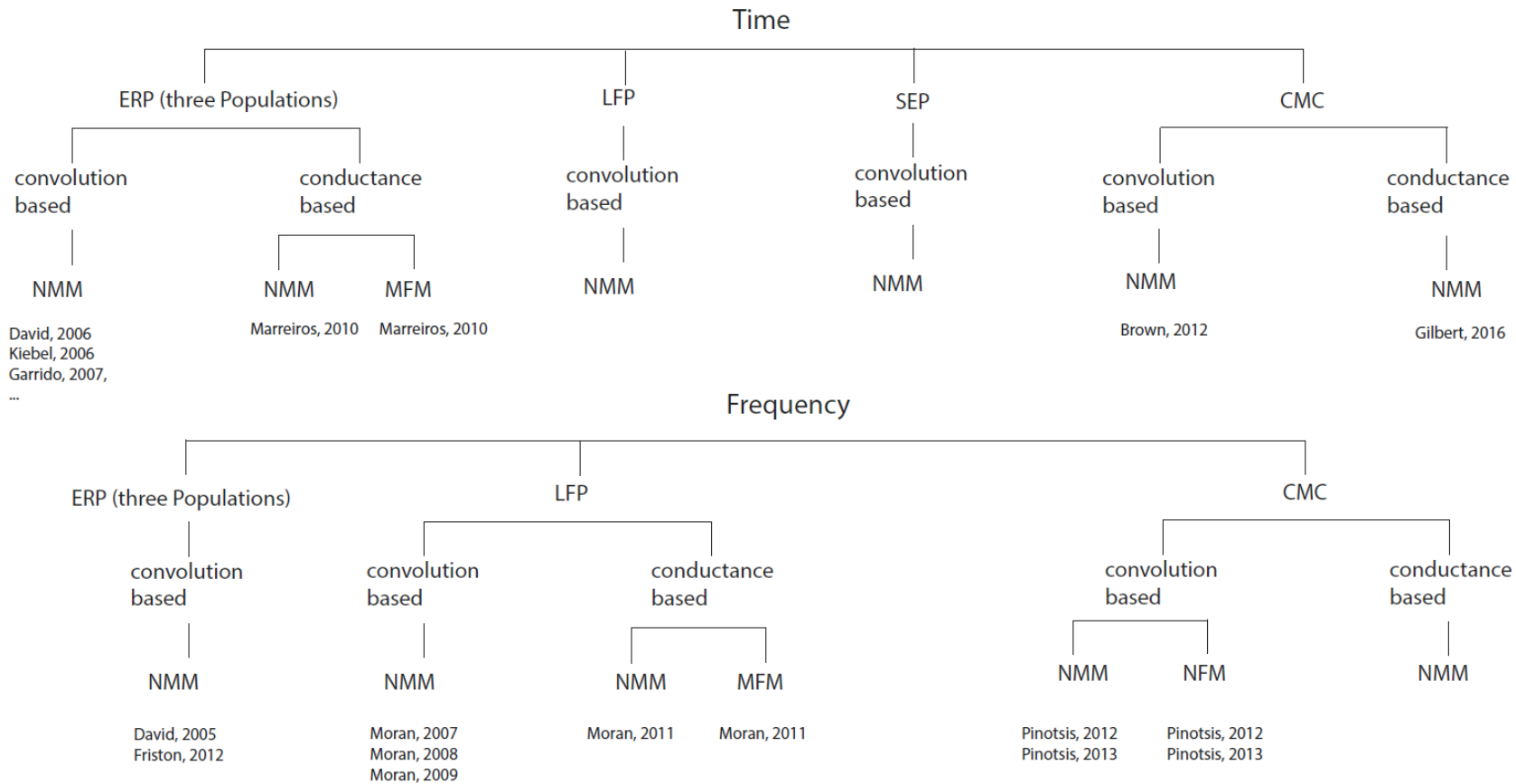
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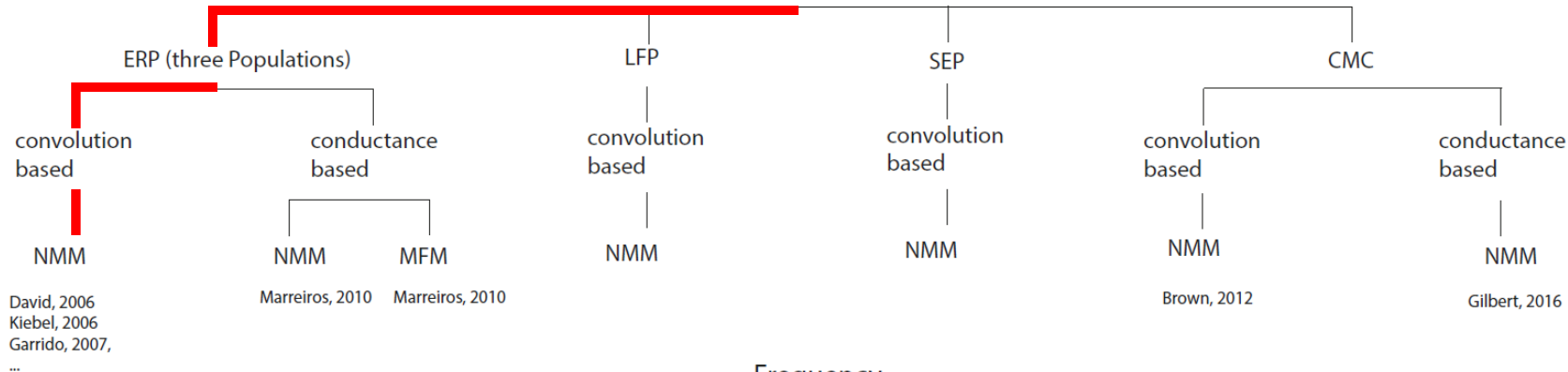


Glossary:

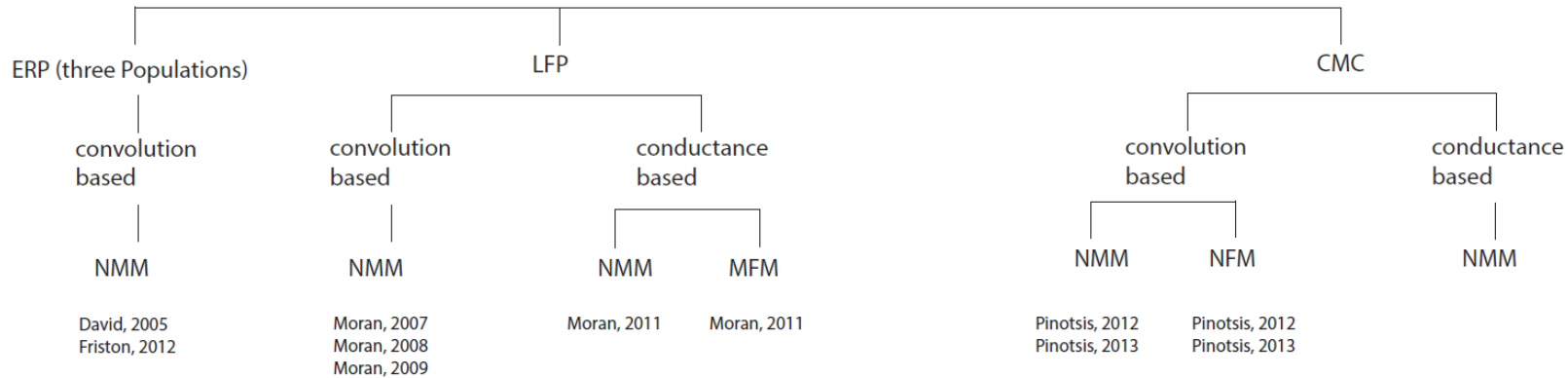
- ERP (Evoked Response Potential)
- LFP (Local Field Potential)
- SEP (Somatosensory Evoked Potential)
- CMC (Canonical Microcircuit)
- NMM (Neural Mass Model)
- MFM (Mean Field Model)
- NFM (Neural Field Model)

focus of today

Time



Frequency



Glossary:

- ERP (Evoked Response Potential)
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I. Macroscale

Scalp maps, dipoles and connected networks



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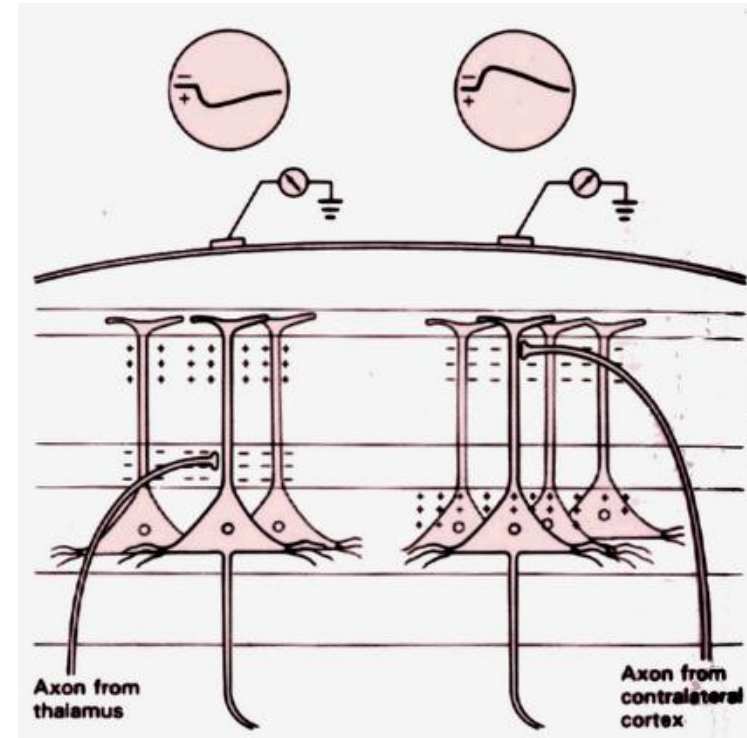
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EEG

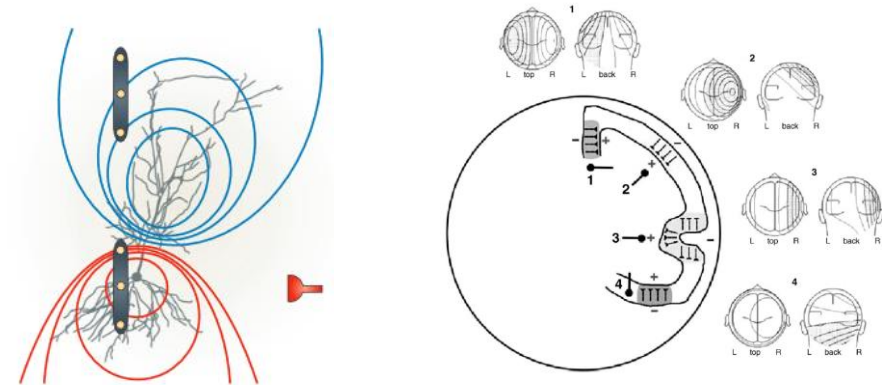
- Temporary accumulation of positive and negative charge during the generation of action potentials
- Resulting Electrical Potential (Energy) is a **scalar** quantity. It depends on the **medium**, and the location wrt the '**source**', thus leading to Potential Energy differences on the scalp.



Buzsaki et al., 20012, Nature Reviews, Neuroscience

Forward Model

- Local Field Potentials (LFP)
 - Measure the activity much closer to the source.
 - 'Simpler' forward model, because we have a much more direct measure of source activity.
- Sensor-Level Data
 - Electrical Potential 'travels' from source to the scalp.
 - This mapping is usually referred to as **Leadfield matrix**



Aguiar et al, 2000, Symposium on Applied Computing

Ebersole, Handbook of Clinical Neurophysiology, 2004

Forward Model

> ECD

- Sensor-Level Data
 - Given some neural activity modeled as a **dipole** at some **location** in the cortex and some **momentum** (vector), the **leadfield** matrix computes the projection of this activity onto the scalp.

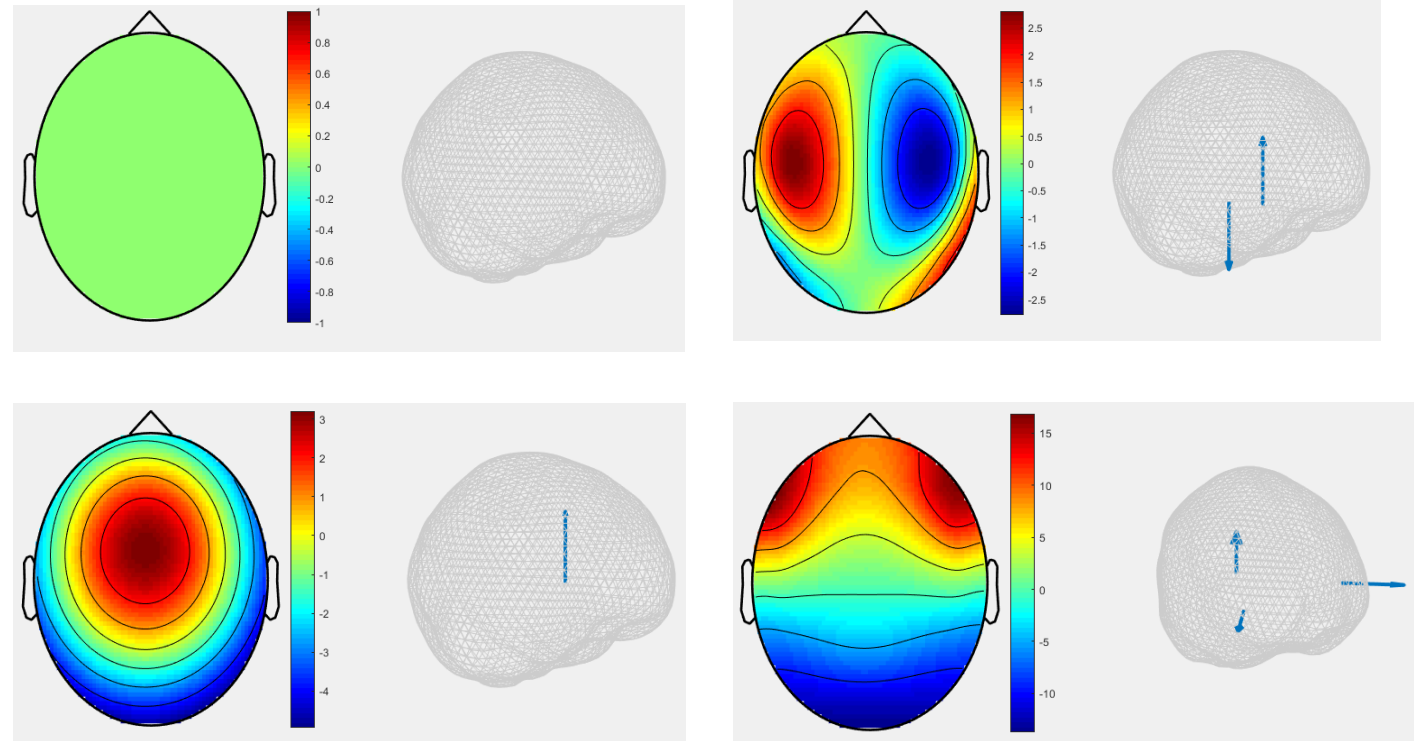


Figure | Different possible dipole configurations that lead to different scalp potentials.

Almost Dynamic Causal Model

- We want to model the full sensor x time space
- Electrode Activity , Scalp Activity and Source Activity
- In DCM, the dipole moments are **constant** quantities, but act as **gain factor** on the **neural dynamics**!

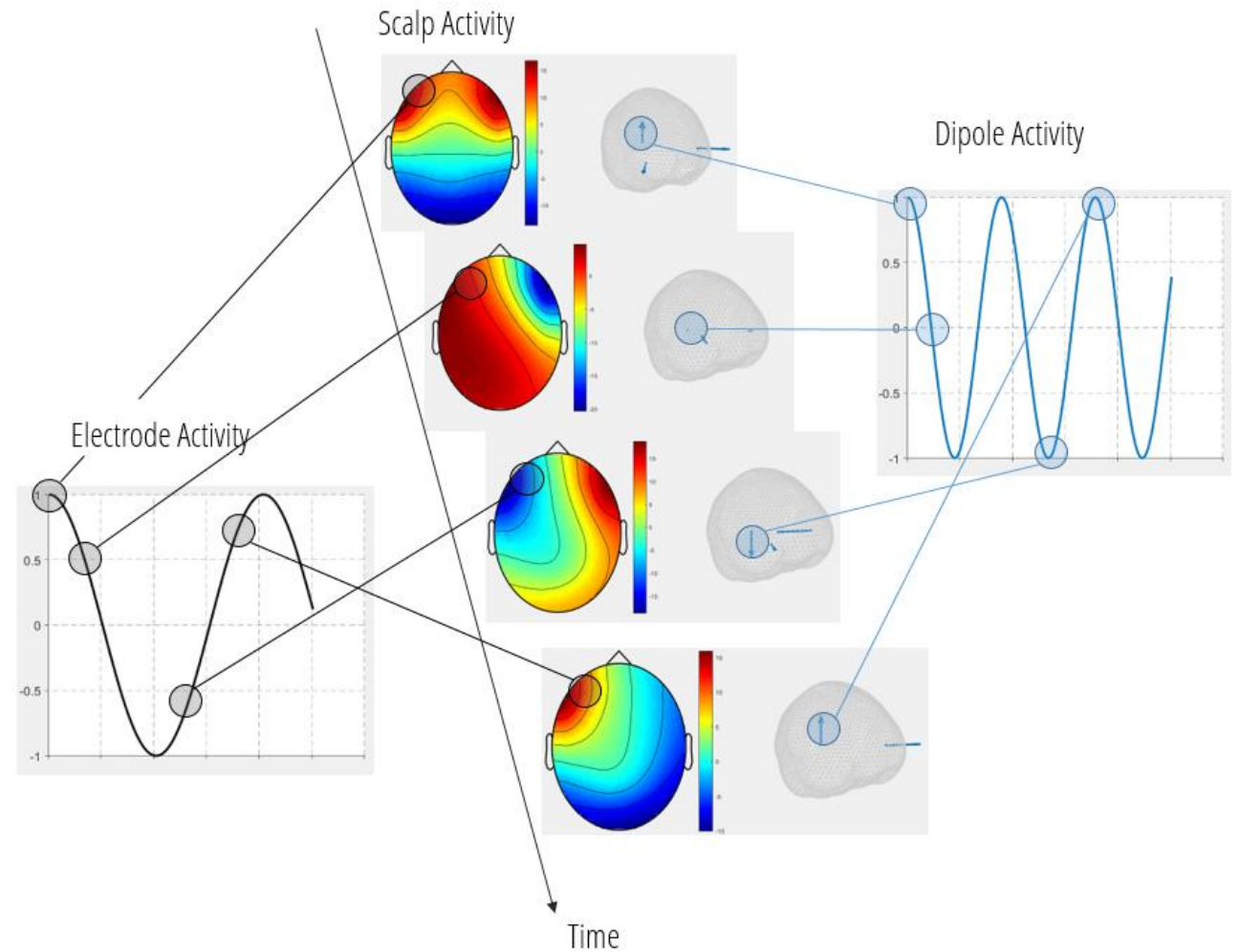


Figure | Changes in the scalp and electrode potential, as the dipole moments change over time.

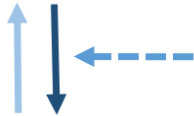
Connected Network

Sources / Regions



Connections

- Forward
- Backward
- Lateral



Modulatory effects



Driving Input

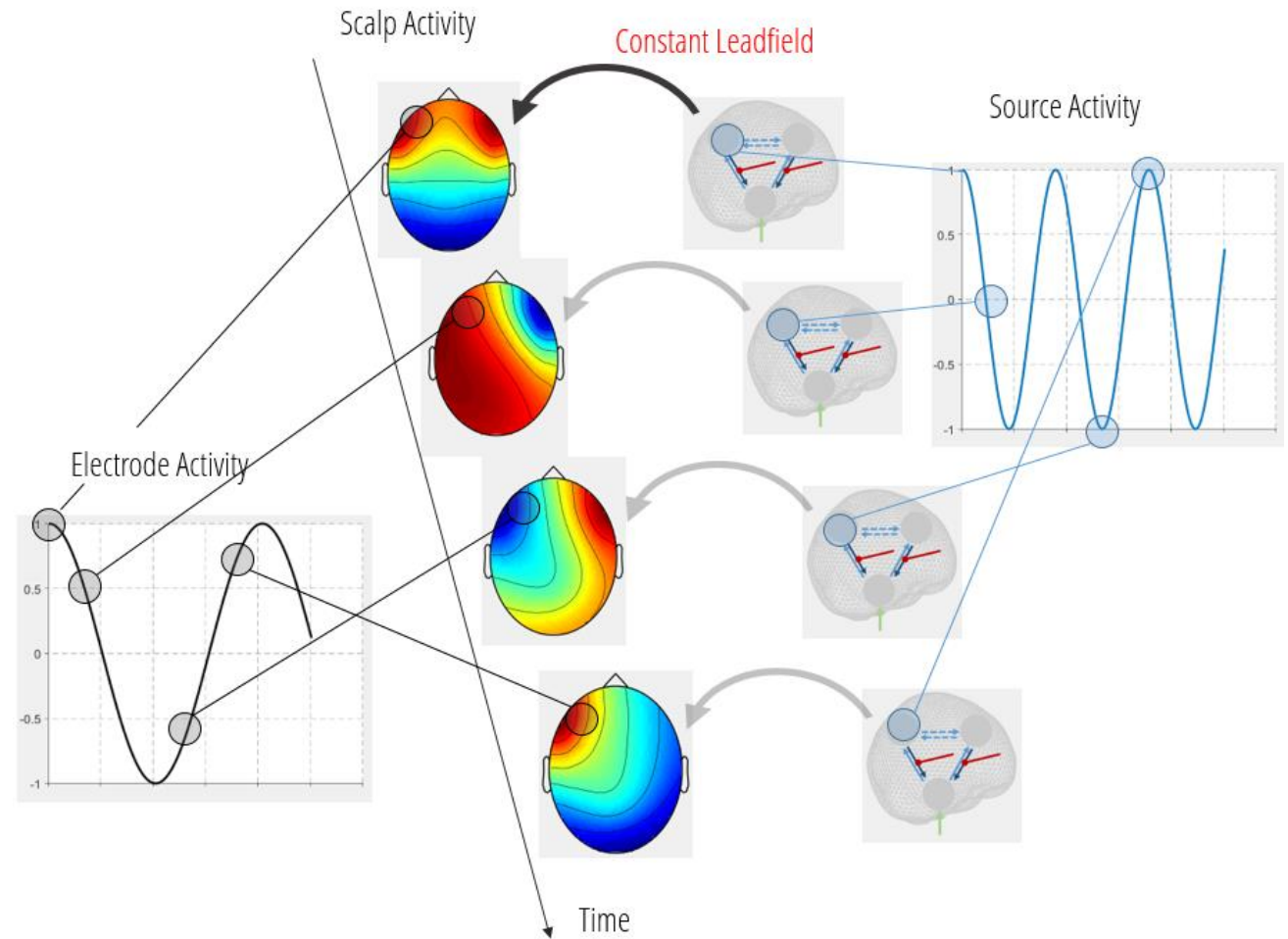
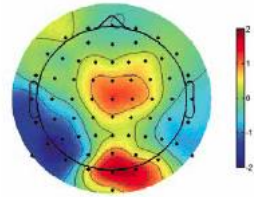


Figure | Changes in the scalp and electrode potential, as the underlying source activity changes over time. The moments of the leadfield act as constant gain factors.

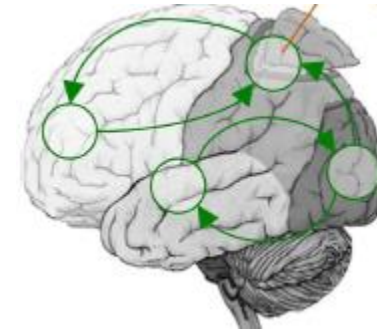
Hypothesis Testing

- Macroscale view (similar to DCM for fMRI)
- Framework to test multiple hypotheses as Bayesian Model Selection (BMS -> Stefan Frässle) questions:

Data



(Hidden) Neuronal Model



Inverse Problem: Inference

$$p(\theta, \Sigma | y)$$

Forward model: Prediction

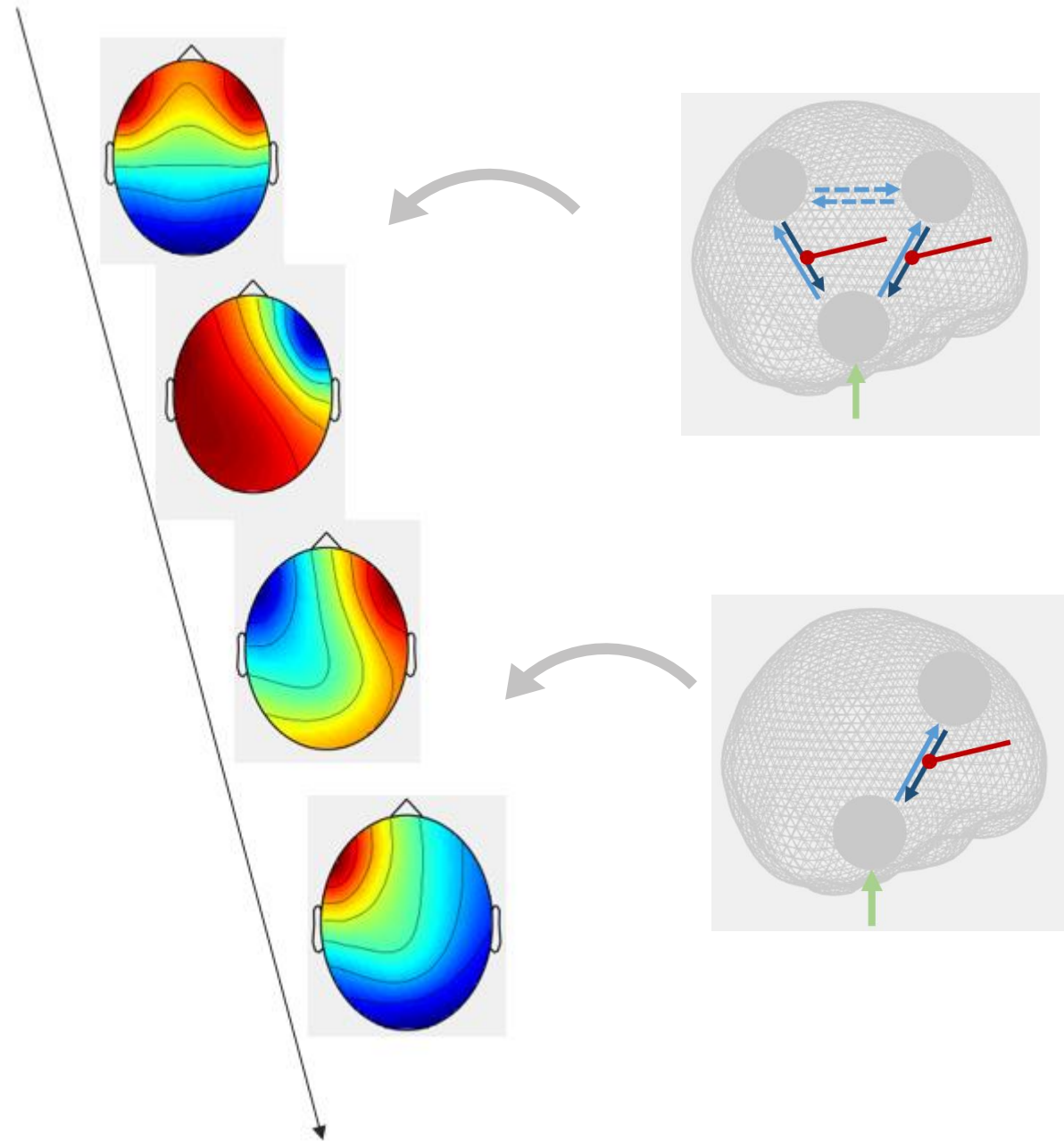
$$y = g(x, \theta) + \varepsilon$$

Dynamic Equations

$$\frac{dx}{dt} = f(x, u, \theta)$$

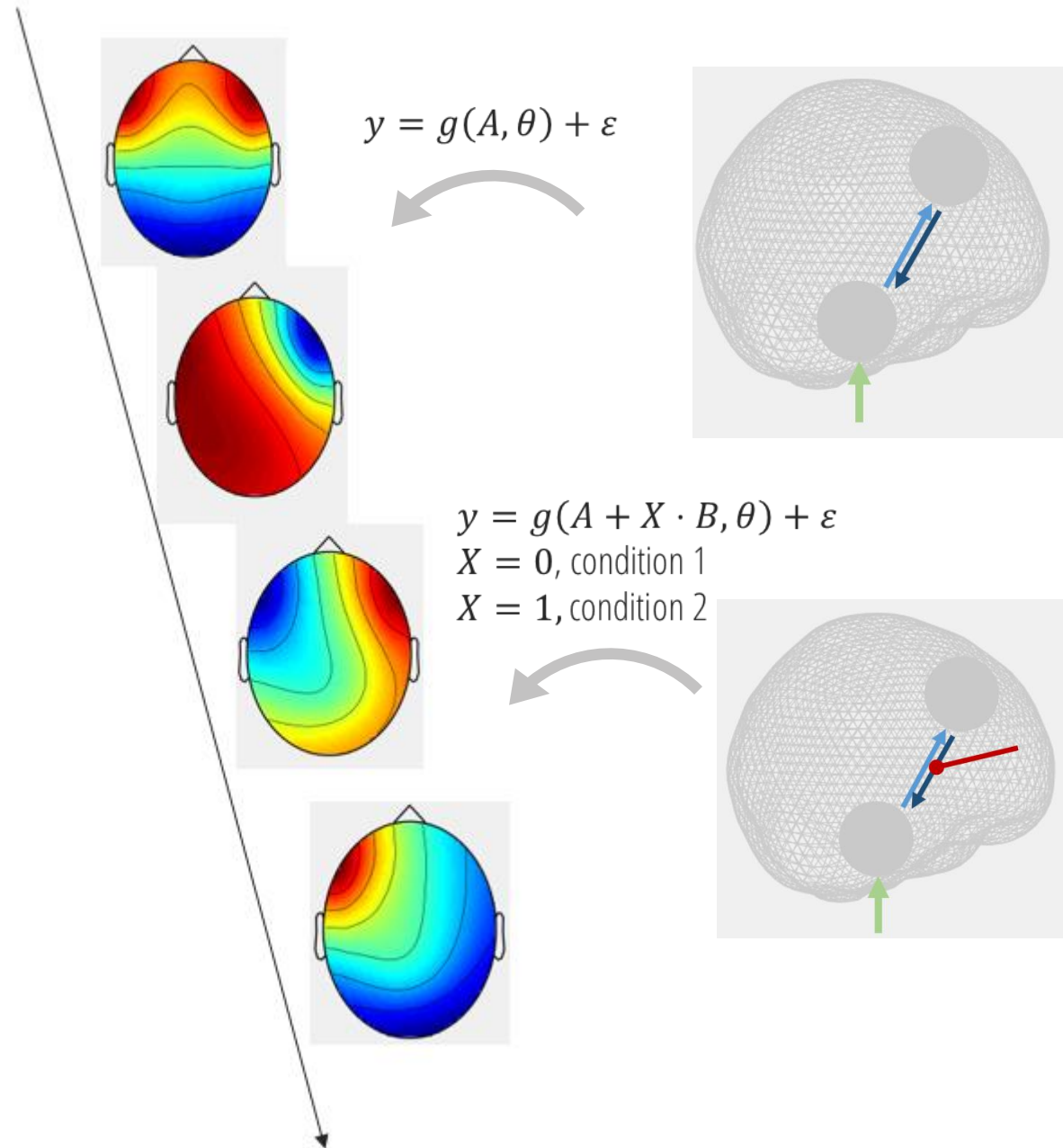
Hypothesis Testing

- Macroscale view (similar to DCM for fMRI)
 - Framework to test multiple hypotheses as Bayesian Model Selection (BMS -> Stefan Frässle) questions:
 - Does a model including regions A, B and C explain the data better than a model including only A and B.
- Only possible for scalp data (not LFP or fMRI)!



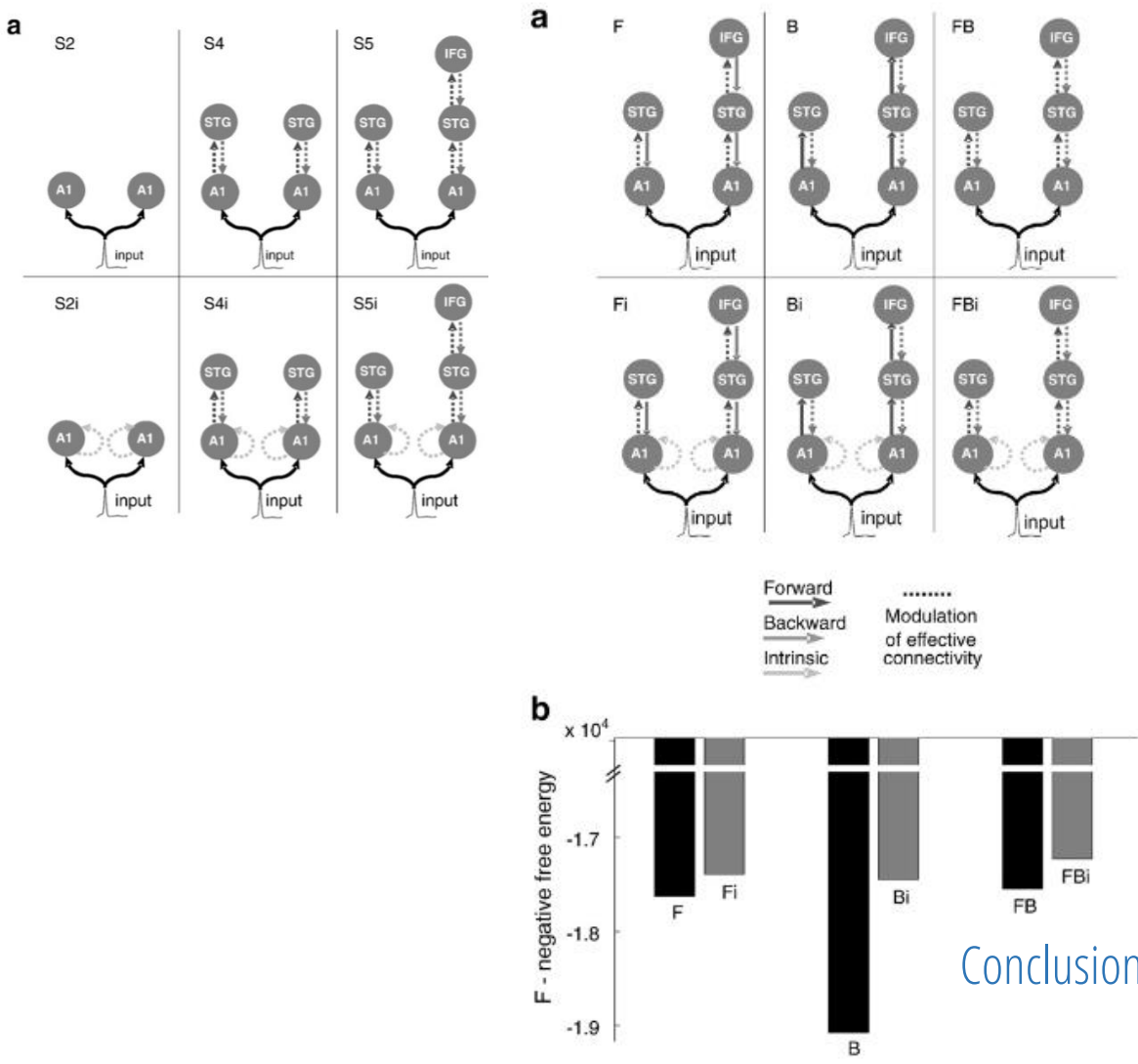
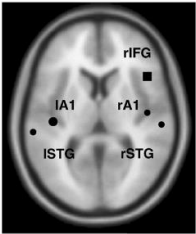
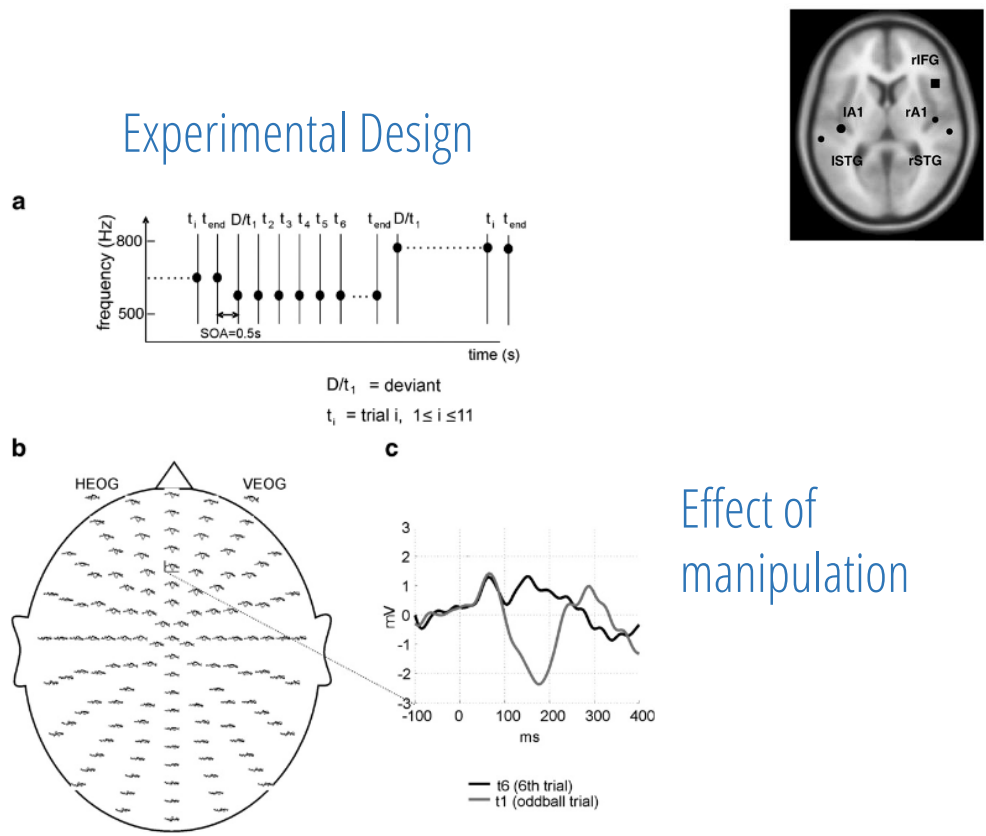
Hypothesis Testing

- Macroscale view (similar to DCM for fMRI)
- Framework to test multiple hypotheses as Bayesian Model Selection (BMS -> Stefan Frässle) questions:
 - Does a model including regions A, B and C explain the data better than a model including only A and B.
 - Can we explain a difference in activation between conditions as a condition specific modulation of one of the connections?



STUDY: IDENTIFYING MECHANISMS

Competing Hypotheses with regard to mechanisms expressed through different modulation structure.



Conclusion

II. Mesoscale

Layered Structure of the cortical column



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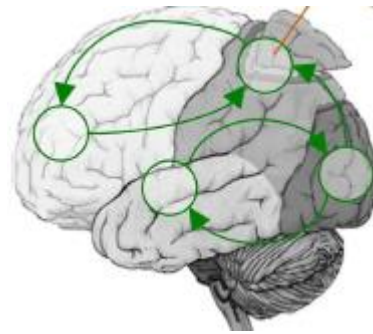
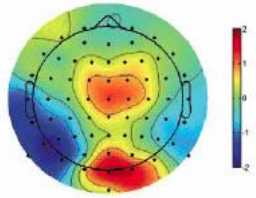
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Swiss Federal Institute of Technology Zurich

Data

(Hidden) Neuronal Model

Inverse Problem: Inference

$$p(\theta, \Sigma | y)$$



Forward model: Prediction

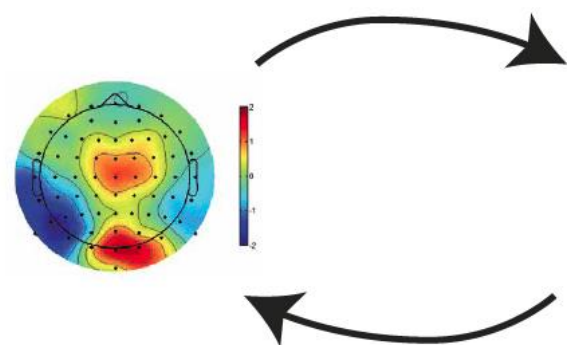
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Dynamic Equations

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Data

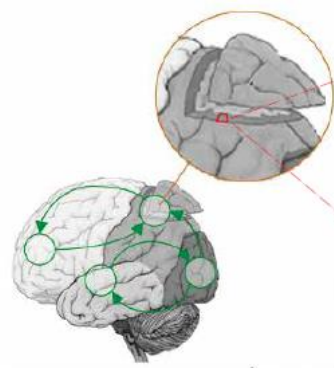
(Hidden) Neuronal Model



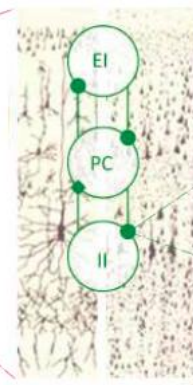
macroscale

mesoscale

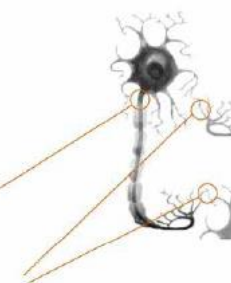
microscale



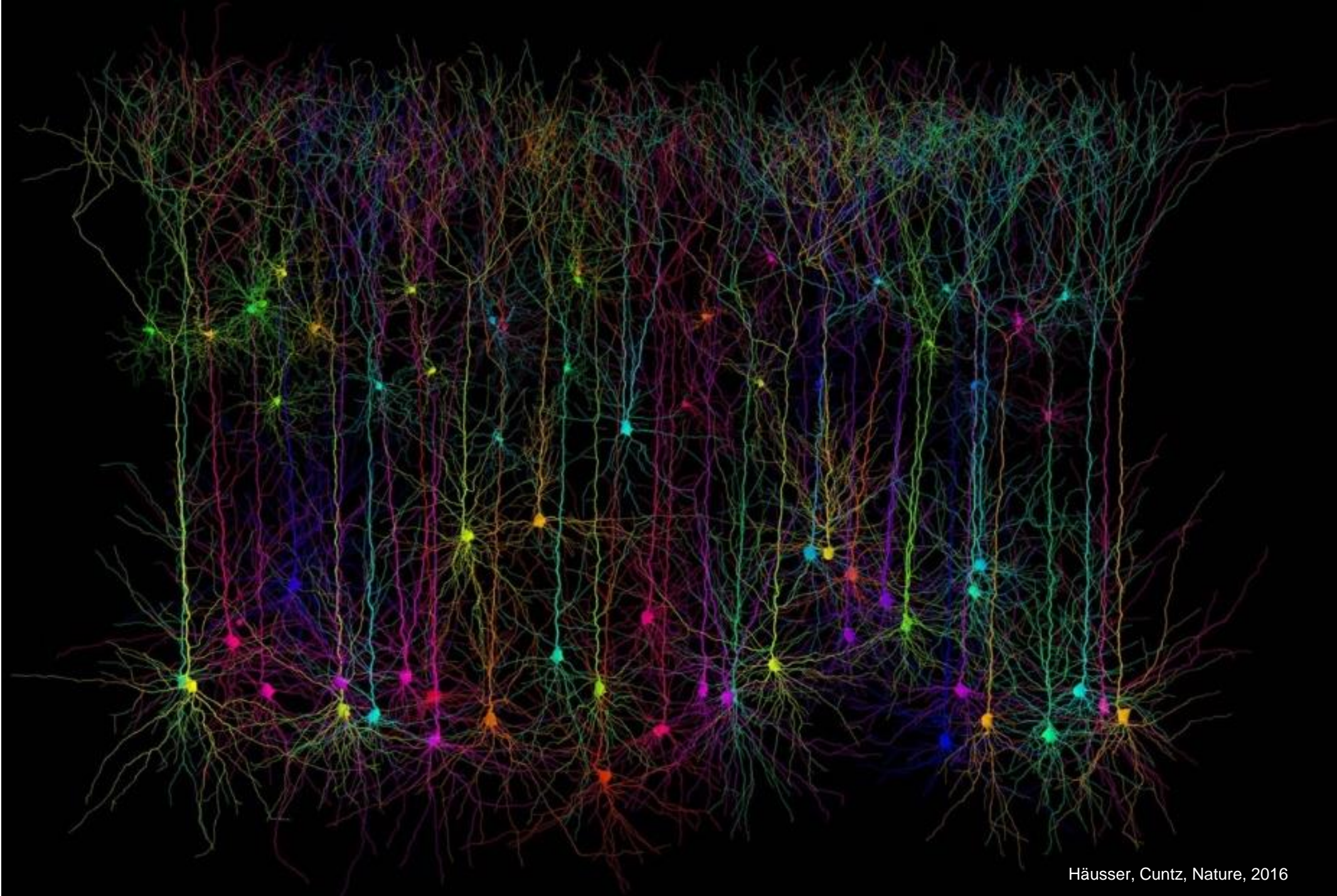
Recurrent network of cortical sources

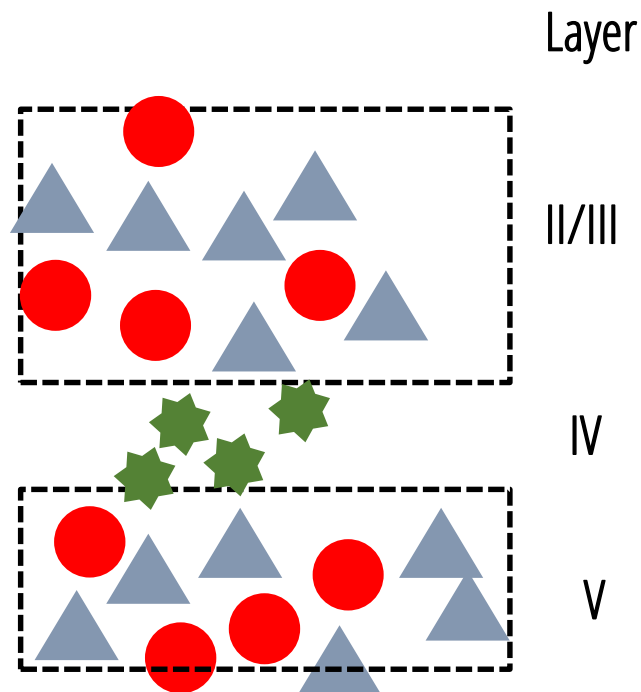
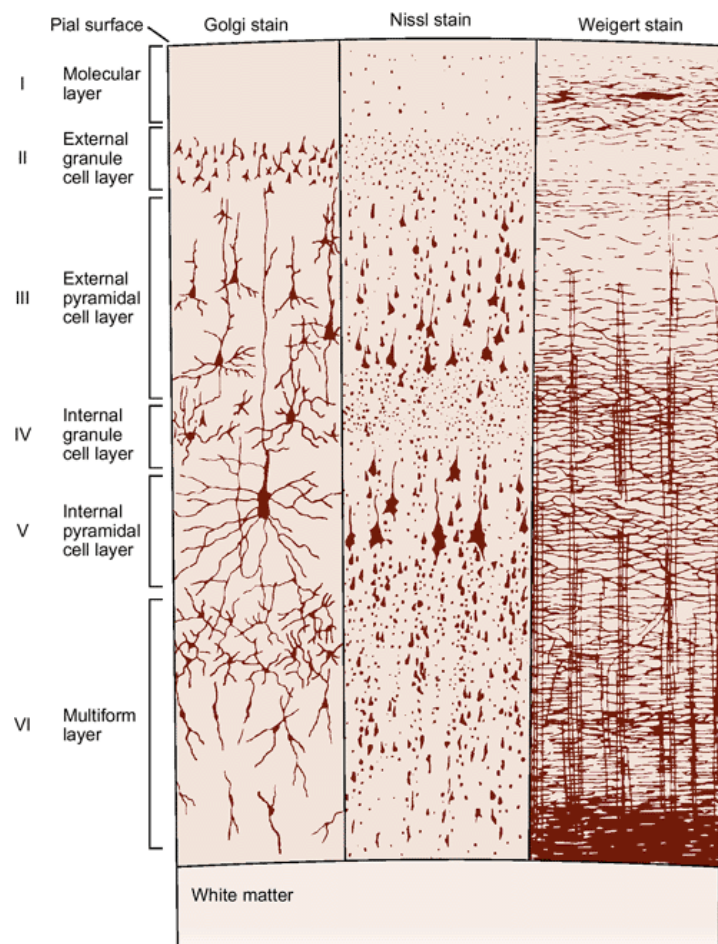


Layered Structure of the cortical column



$$\frac{dx}{dt} = f(x, u, \theta)$$





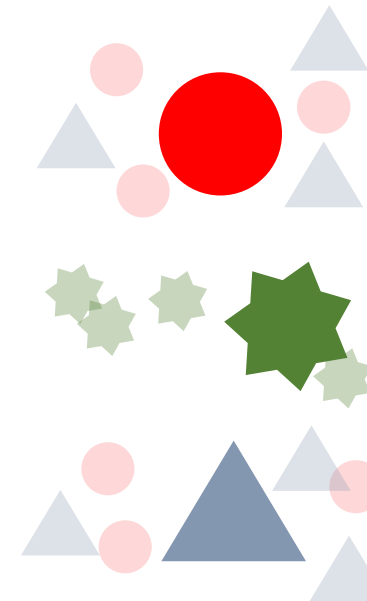
Layer

II/III

IV

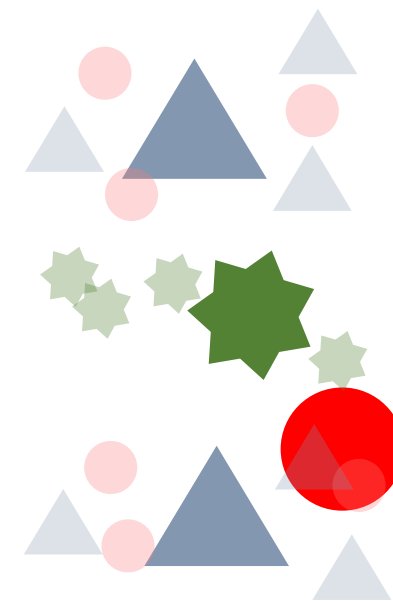
V

ERP



Superficial and deep pyramidal and inhibitory cells are combined in a single population

CMC



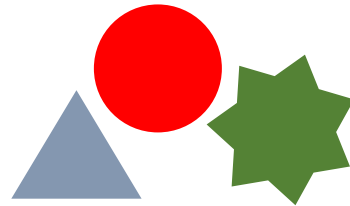
Superficial and deep pyramidal cell populations are modeled individually

Kandel et al. 2000 (from Heimer 1994)

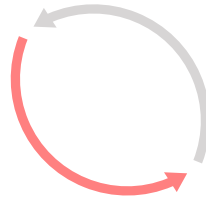
Mesoscale

Three types of Cell Populations:

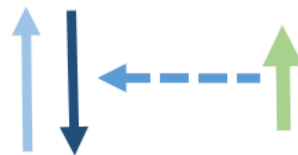
Pyramidal, Inhibitory, Stellate



Inhibitory / Excitatory effects on different populations



Extrinsic Connectivity / Driving Input

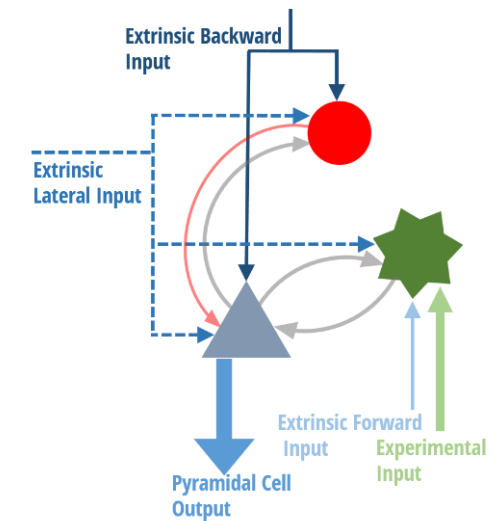
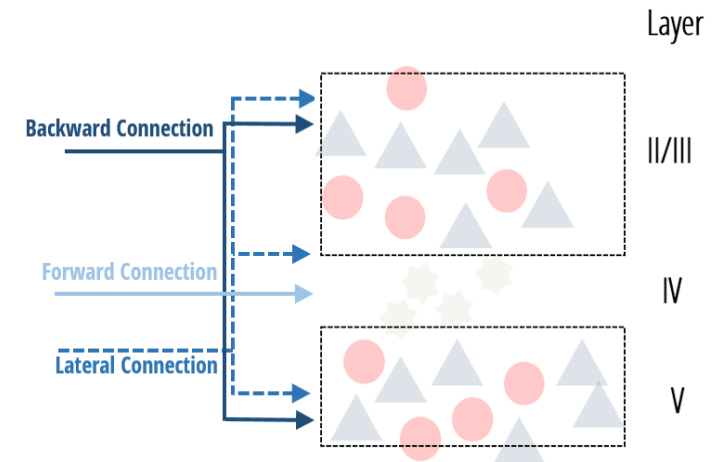


Mesoscale

- Name of a between source connection refers to the layer, which a connection targets
Felleman & van Essen 1991
- Output from the Pyramidal cell population
- Pyramidal Cell contribute most to the EEG signal
- Driving input into stellate cell layer IV

Considerations:

- Importance of distinguishing layer III and V pyramidal cells to model task (e.g. Predictive coding)
- Modeling of particular data features, i.e. Oscillations of particular frequencies.



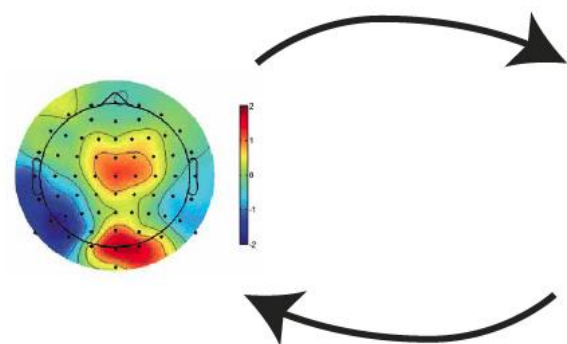
III. Microscale

Mechanisms governing the generation of average post-synaptic potentials



Data

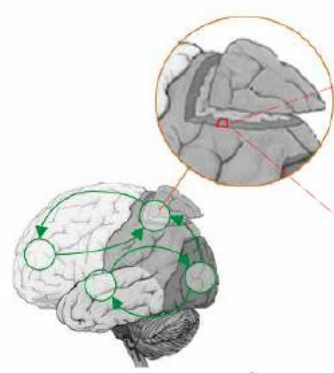
(Hidden) Neuronal Model



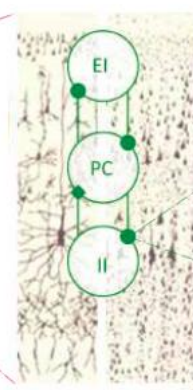
macroscale

mesoscale

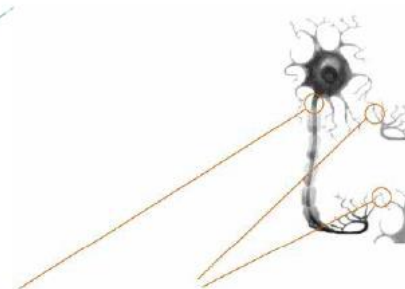
microscale



Recurrent network of cortical sources



Layered Structure of the cortical column



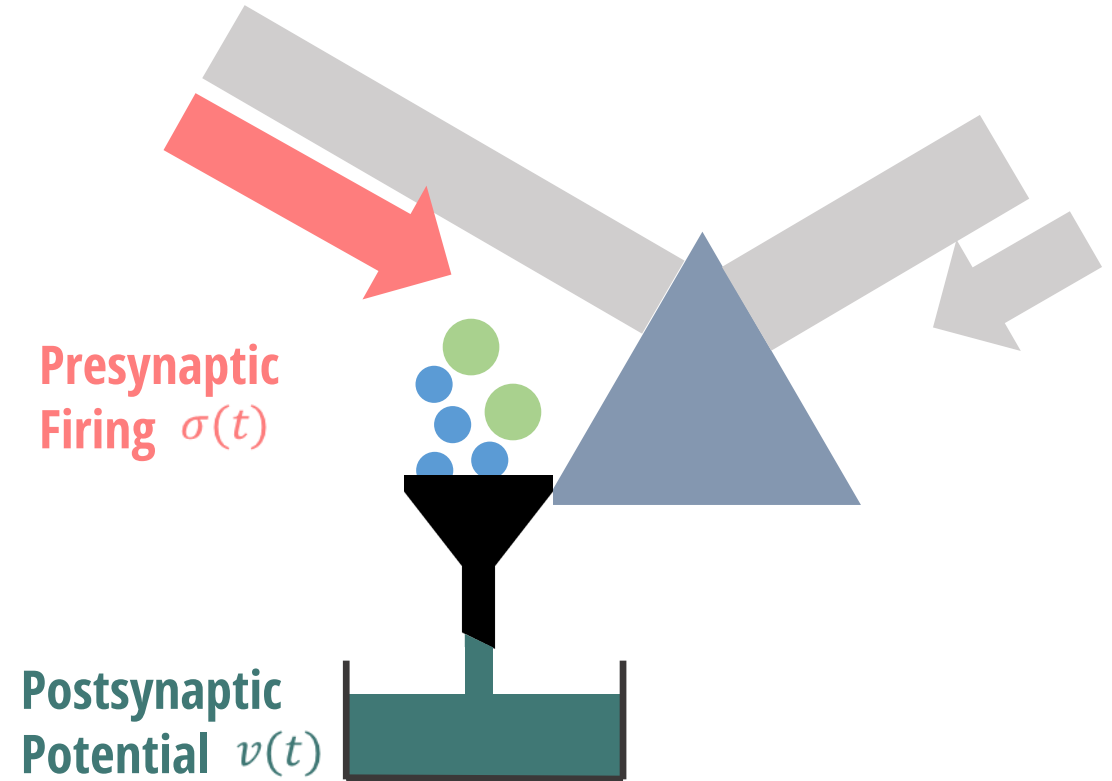
Mechanisms governing generation of average post-synaptic potentials:

$$\frac{dx}{dt} = f(x, u, \theta)$$

Microscale > convolution based DCM

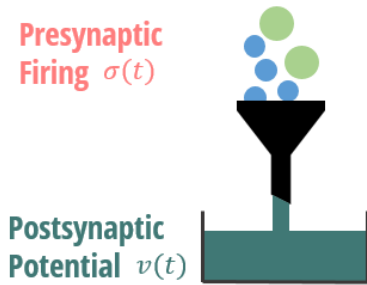
- Jansen and Rit (1995)
- A convolution kernel transforms the presynaptic firing rate into postsynaptic potential
- Kernel parametrized by two parameters

$$v(t) = \int_{-\infty}^t h(t - \tau, H, \kappa) \sigma(\tau) d\tau$$



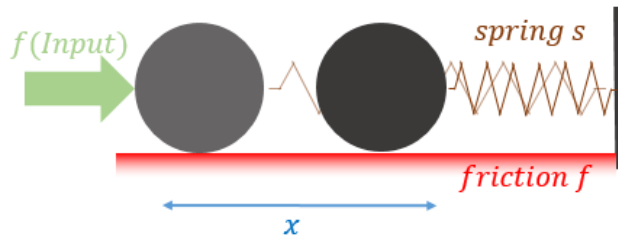
Voltage over time shows similarities with Harmonic Oscillator.

(This is a consequence of the convolution operation)



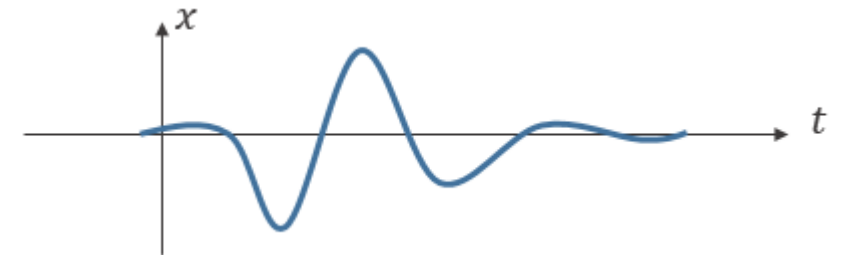
$$\ddot{v} = f(\text{Input}) - \frac{2}{\tau} \dot{v} - \frac{1}{\tau^2} v$$

Equation describing the post synaptic potential in the convolution based DCM.



$$\ddot{x} = f(\text{Input}) - f\dot{x} - sx$$

Equation describing the behavior of a mass attached to a spring (H.O.)



Microscale > convolution based DCM

- Jansen and Rit (1995)
- A convolution kernel transforms the presynaptic firing rate into postsynaptic potential
- Kernel parametrized by two parameters
- For mathematical convenience, the second order differential equations are transformed into first order differential equations.

Current state of the population. How much firing arrives (and from where).

$$\frac{dx}{dt} = f(x, A, H, C, \sigma, \theta_{kernel})$$

What the cell population is connected to. The size of the kernel.

$$\ddot{v} = f(\text{Input}) - \frac{2}{\tau} \dot{v} - \frac{1}{\tau^2} v$$

●

 $k_e A_{back} \triangle + k_e A_{lateral} \triangle + k_e G_{\bullet} \triangle$

★

 $k_e A_{Forward} \triangle + k_e A_{lateral} \triangle + k_e G_{\star} \triangle + C \text{ (trace)}$

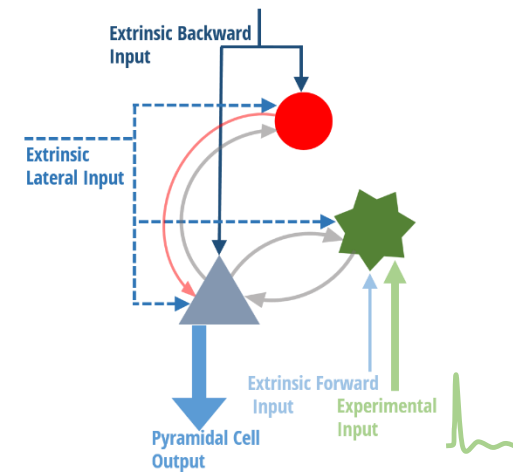
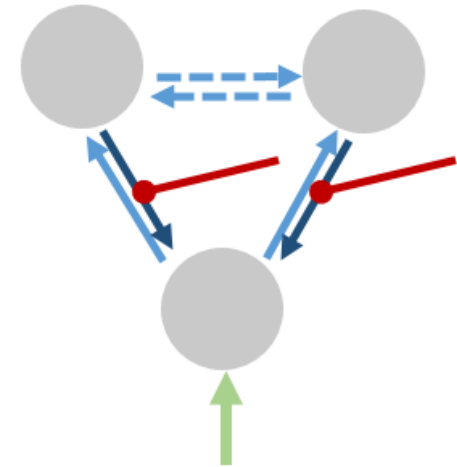
▲
↑

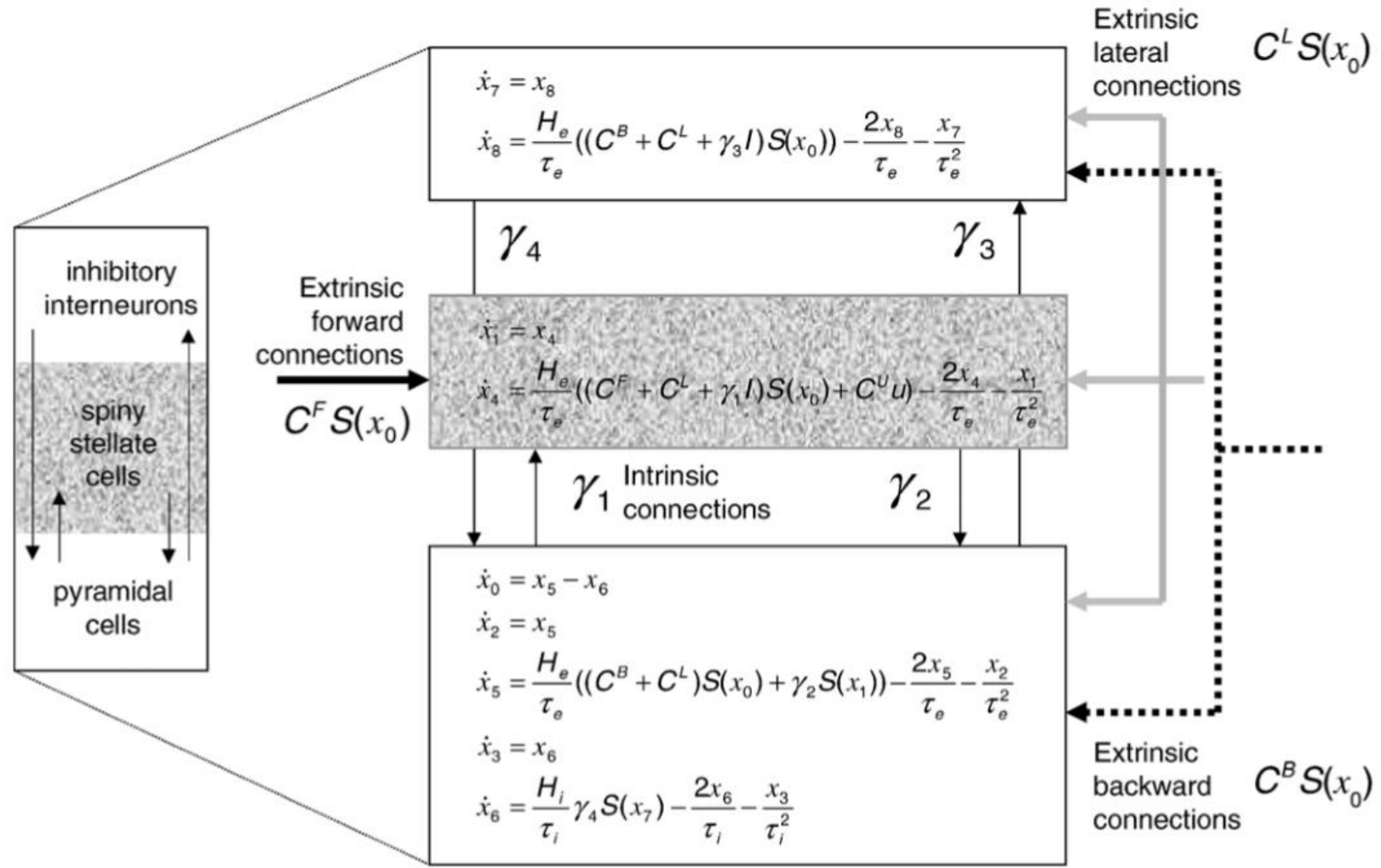
 $k_e A_{back} \triangle + k_e A_{lateral} \triangle + k_e G_{\star} \star$

▲
↓

 $k_i G_{\bullet} \bullet$

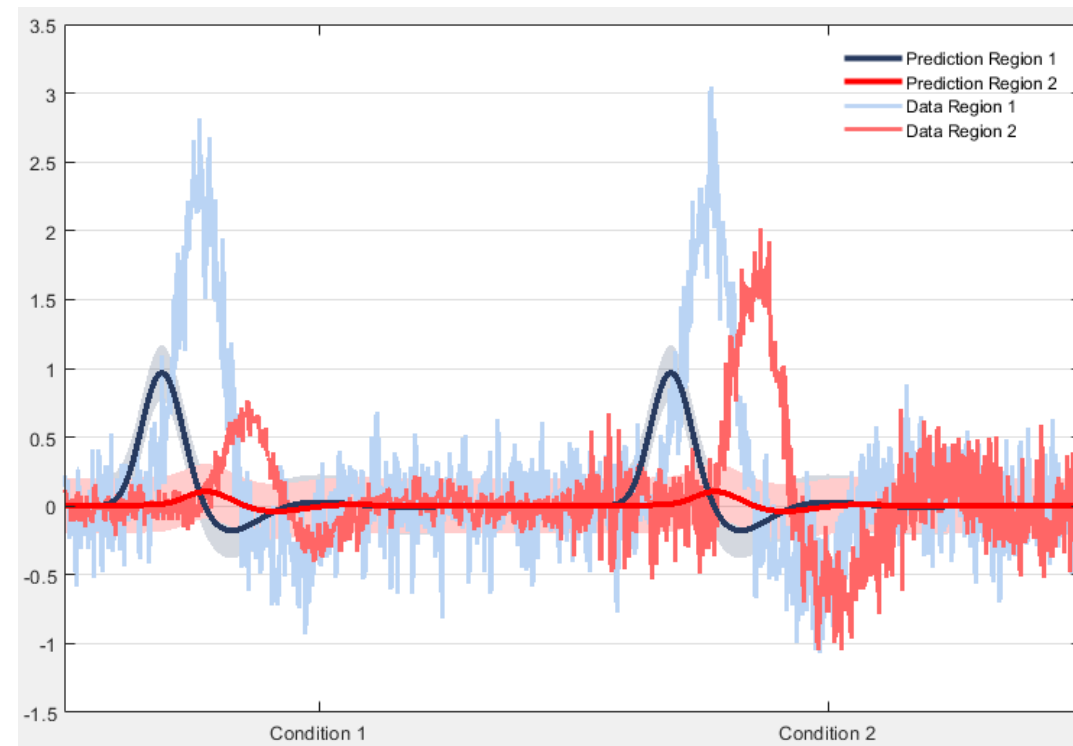
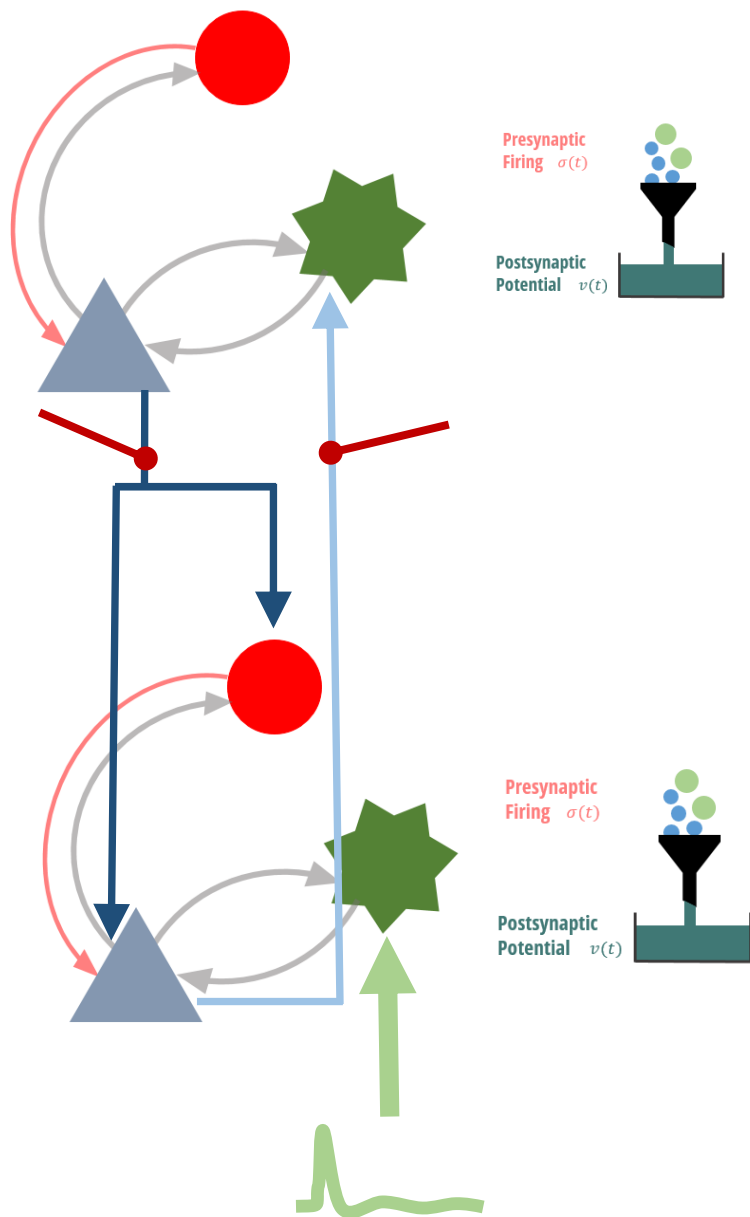
G_{\bullet}^{\triangle} : Coupling strength from \triangle to \bullet .

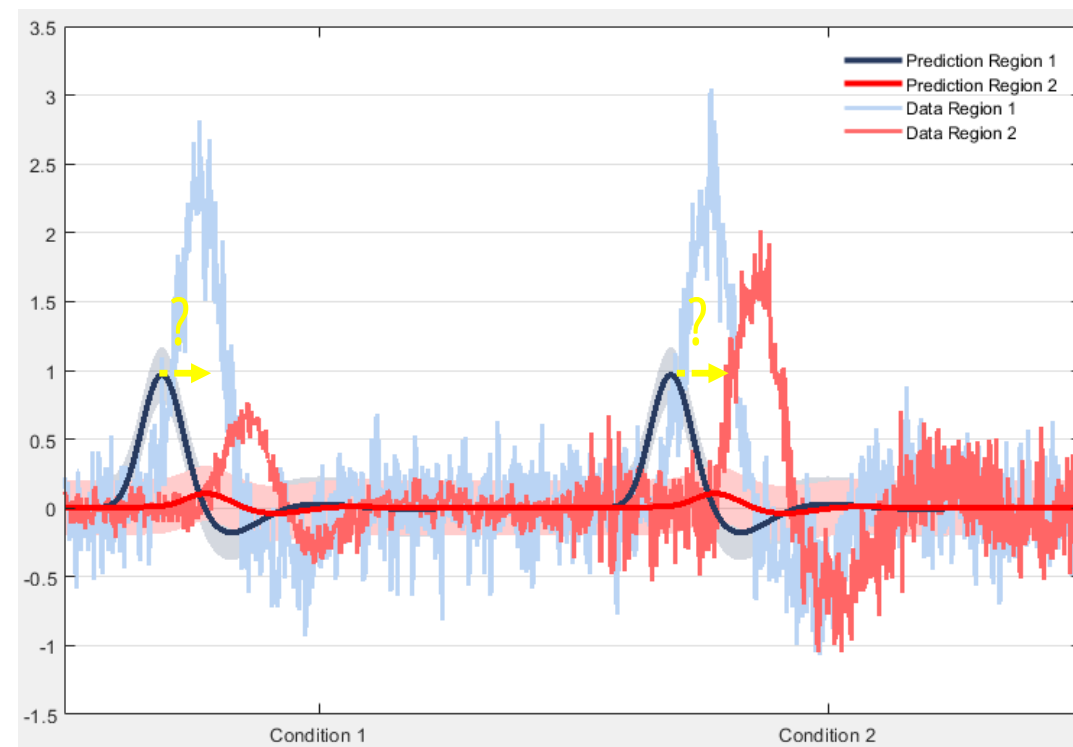
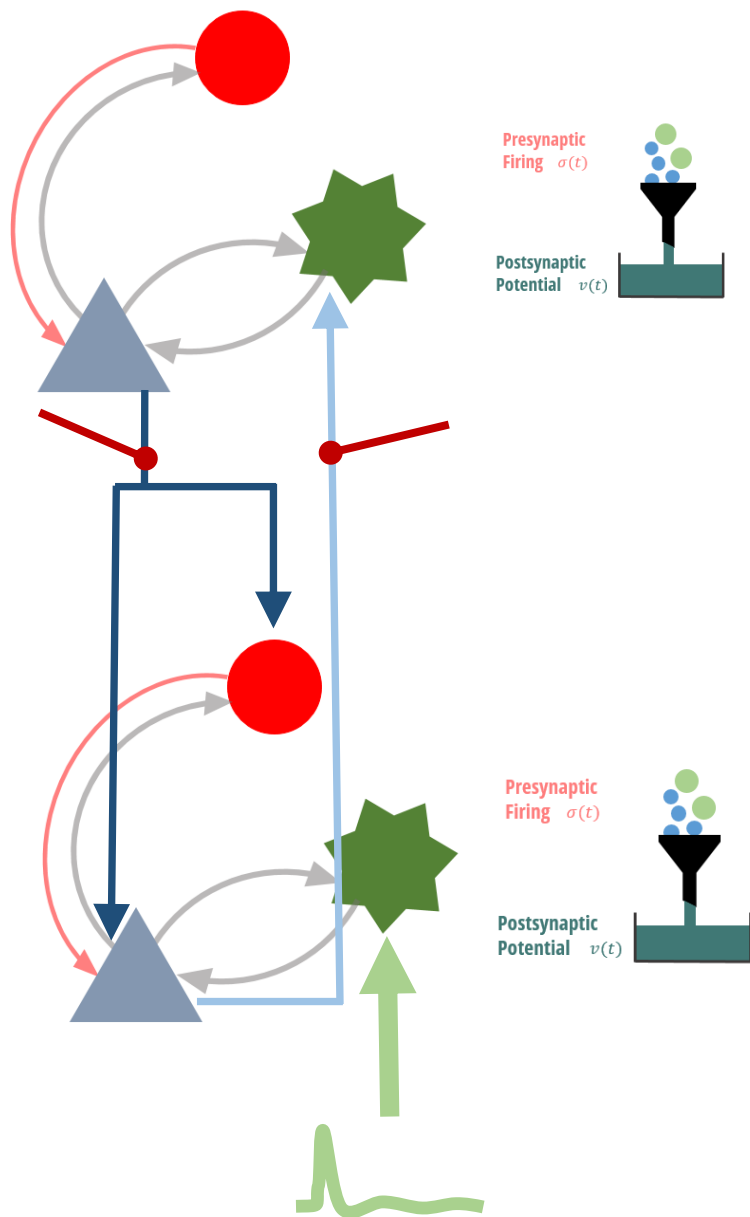


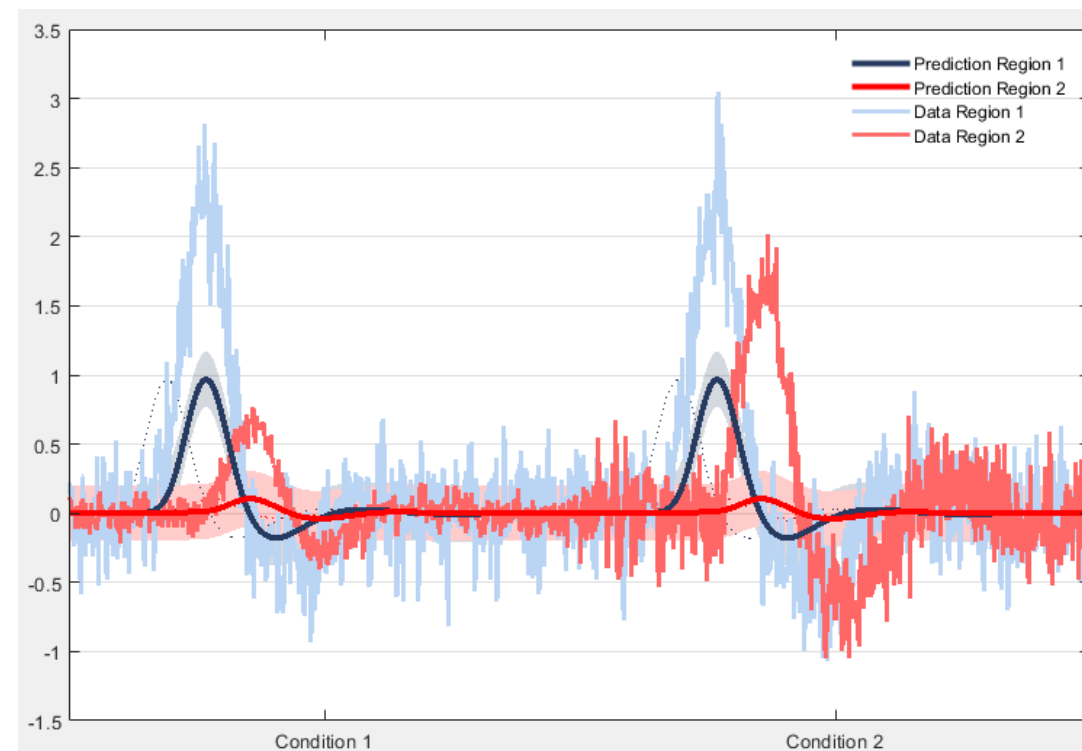
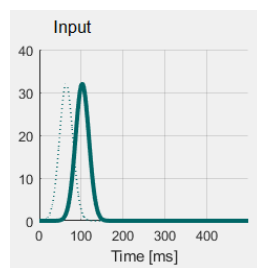
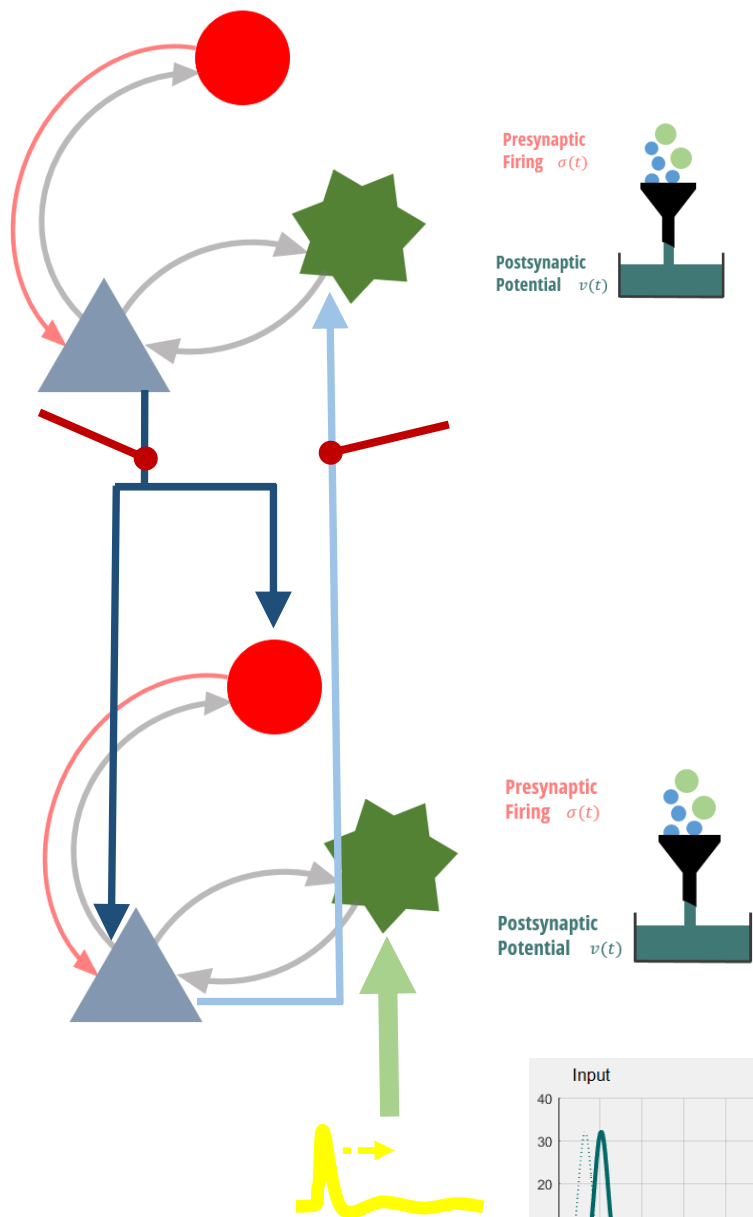


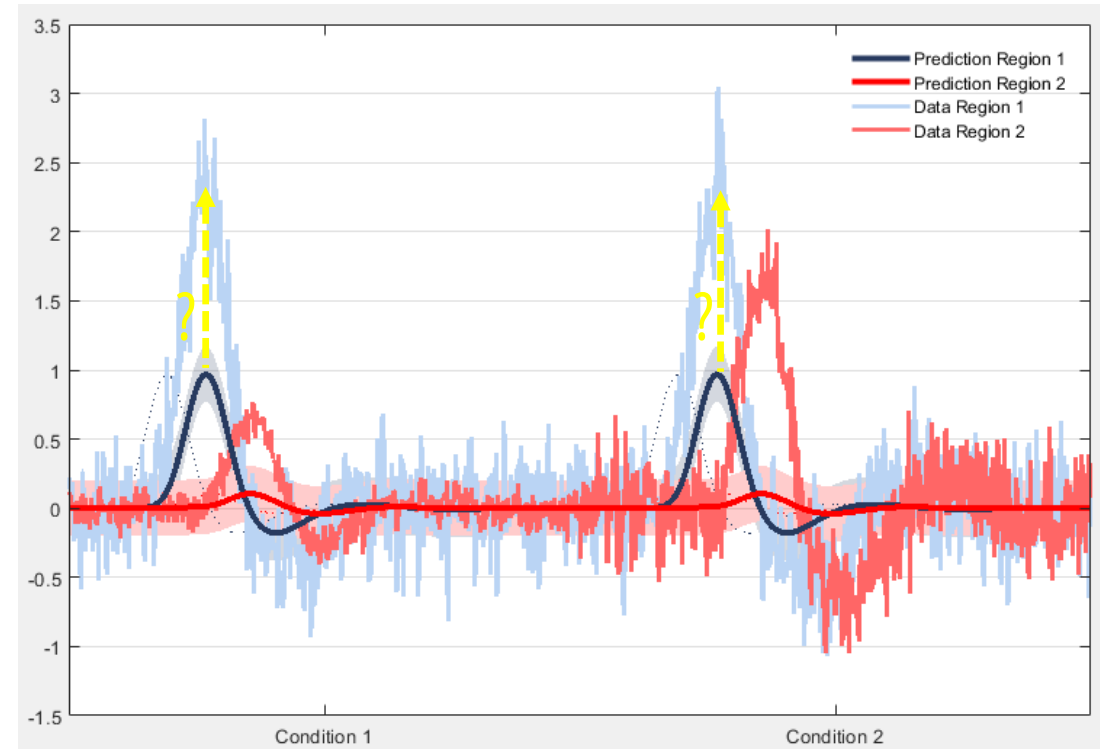
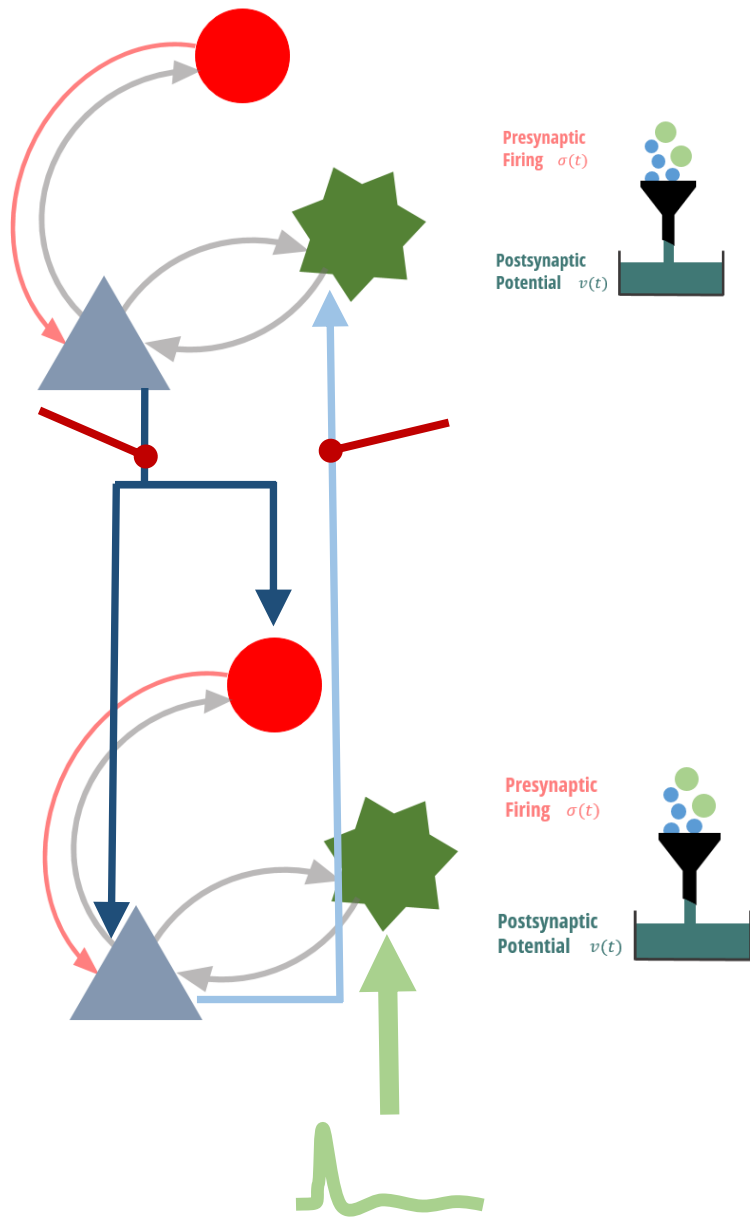
IV. Inferring on parameters

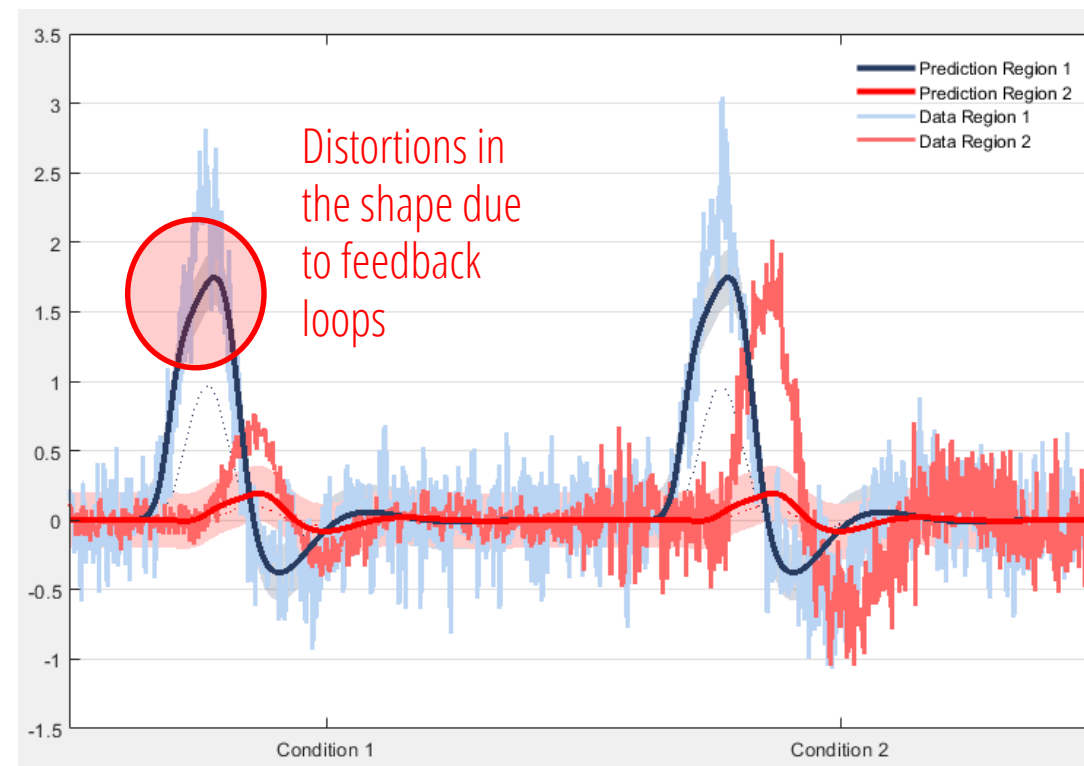
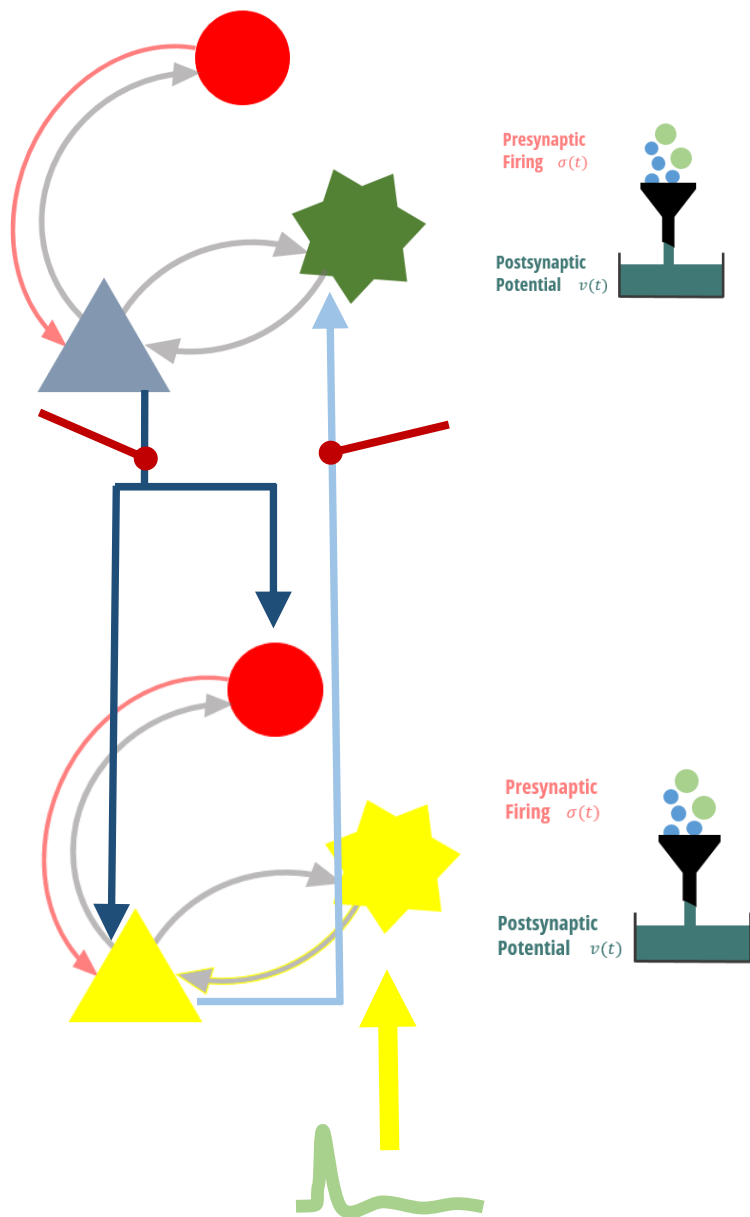
Getting a feeling for how different parameters affect different aspects generated data

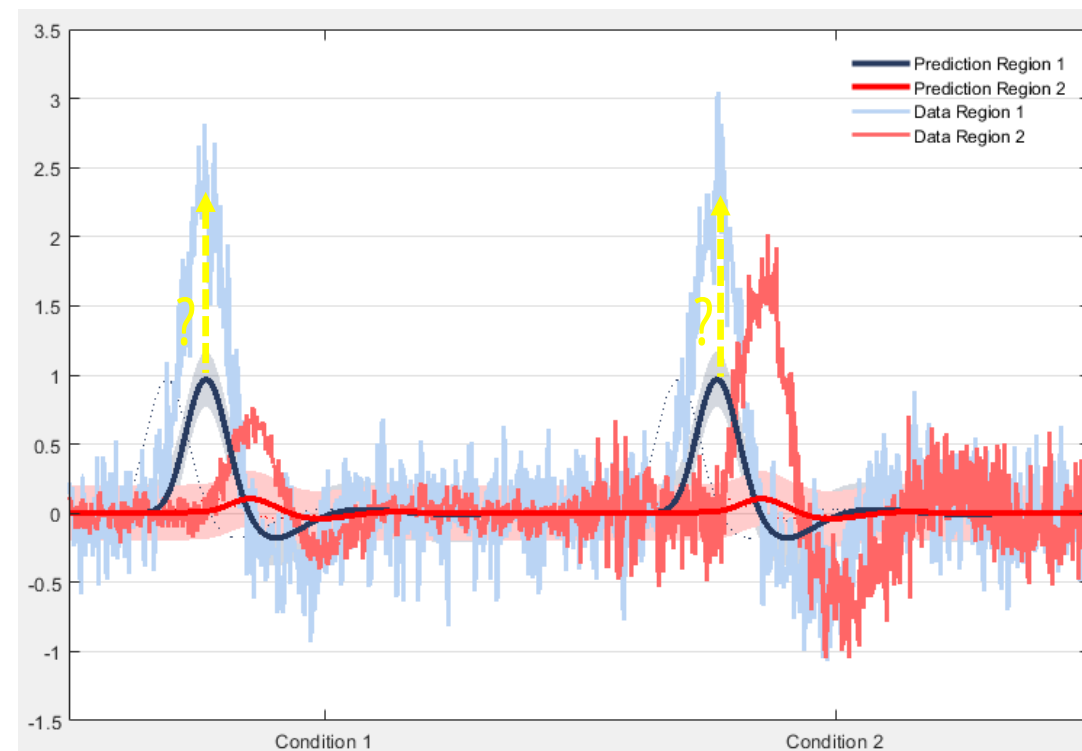
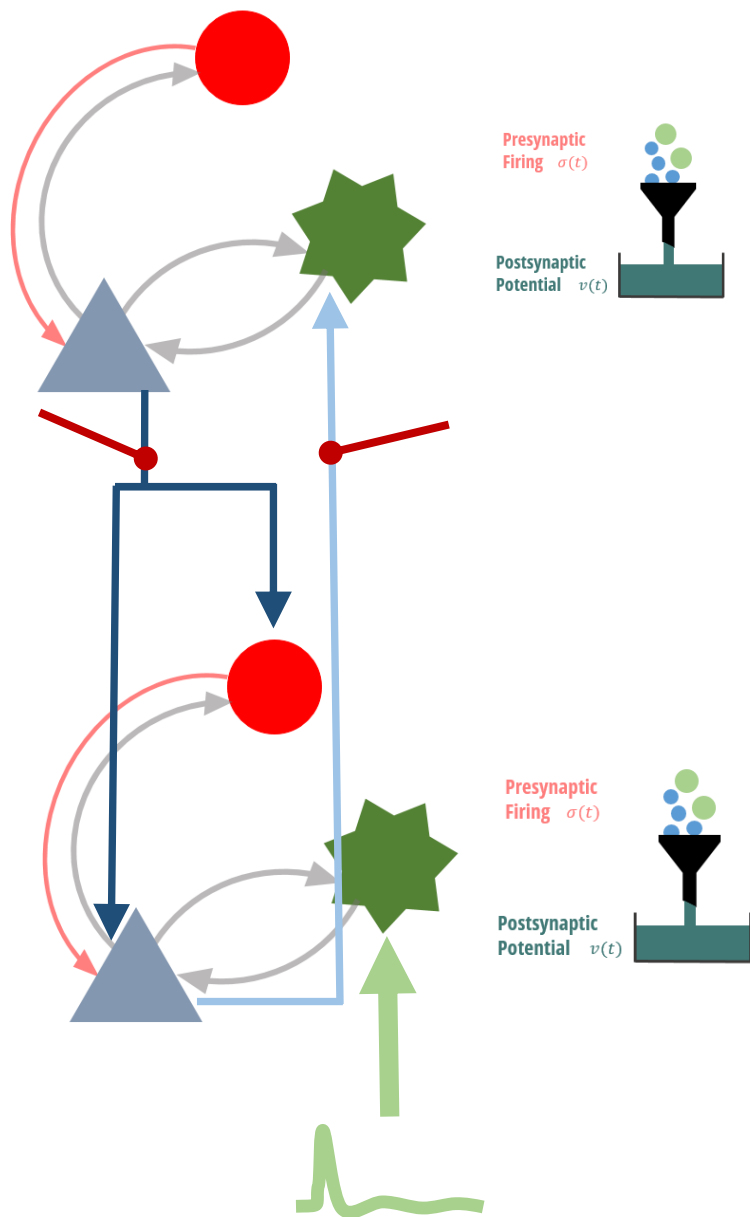


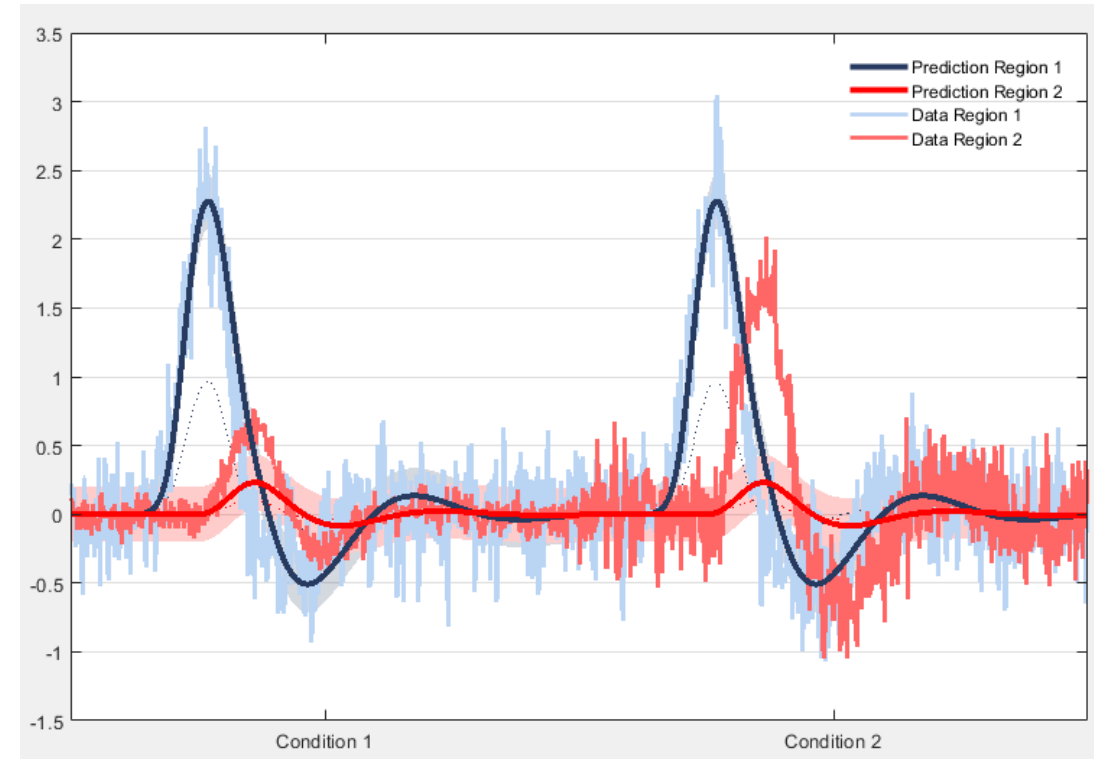
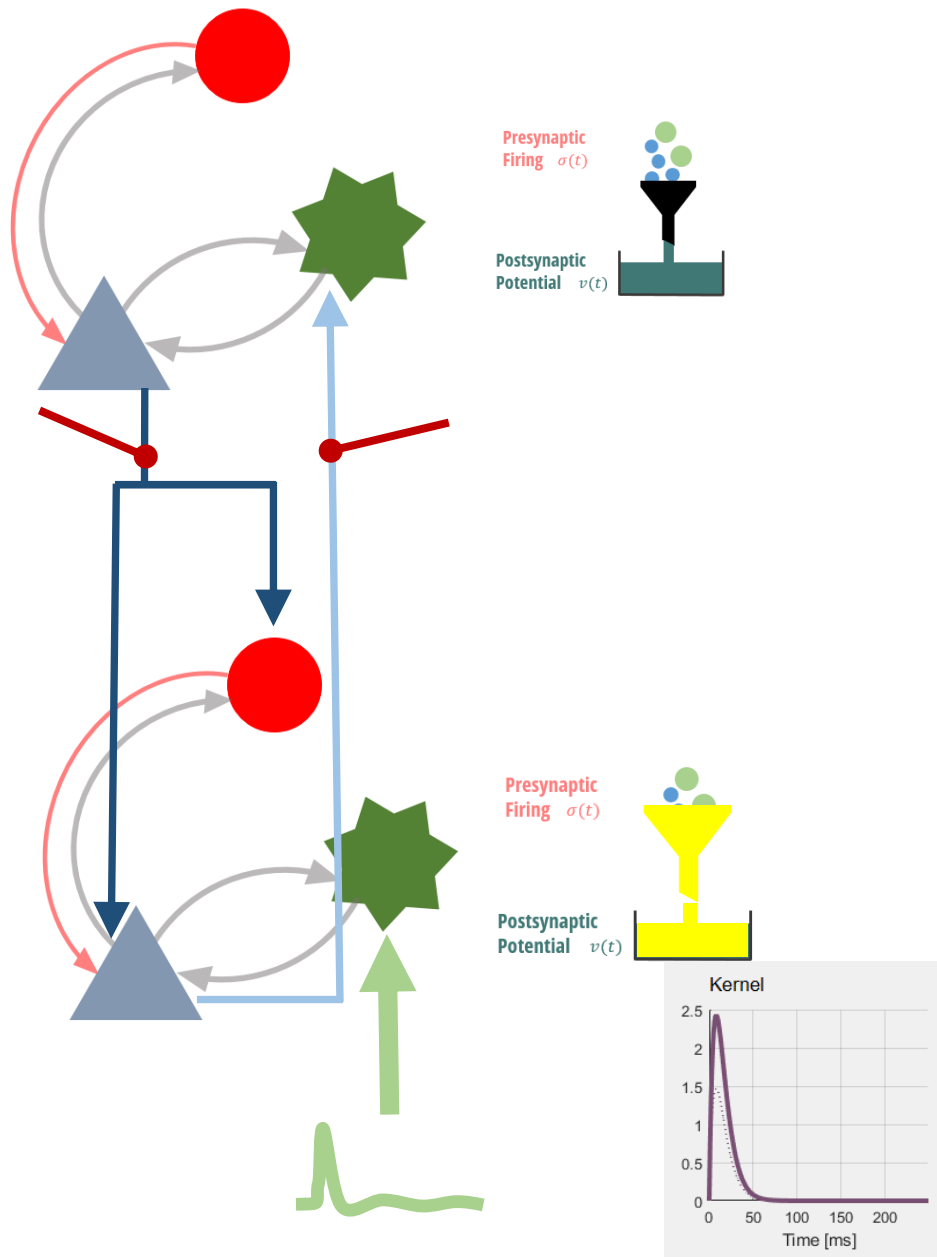


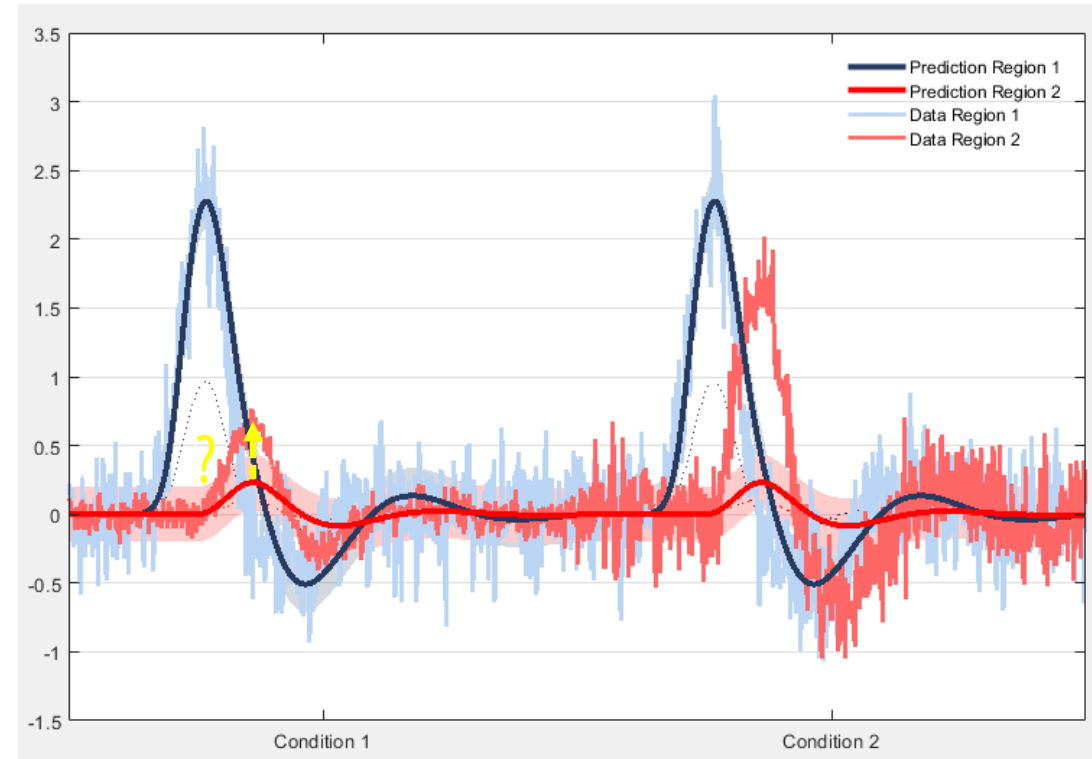
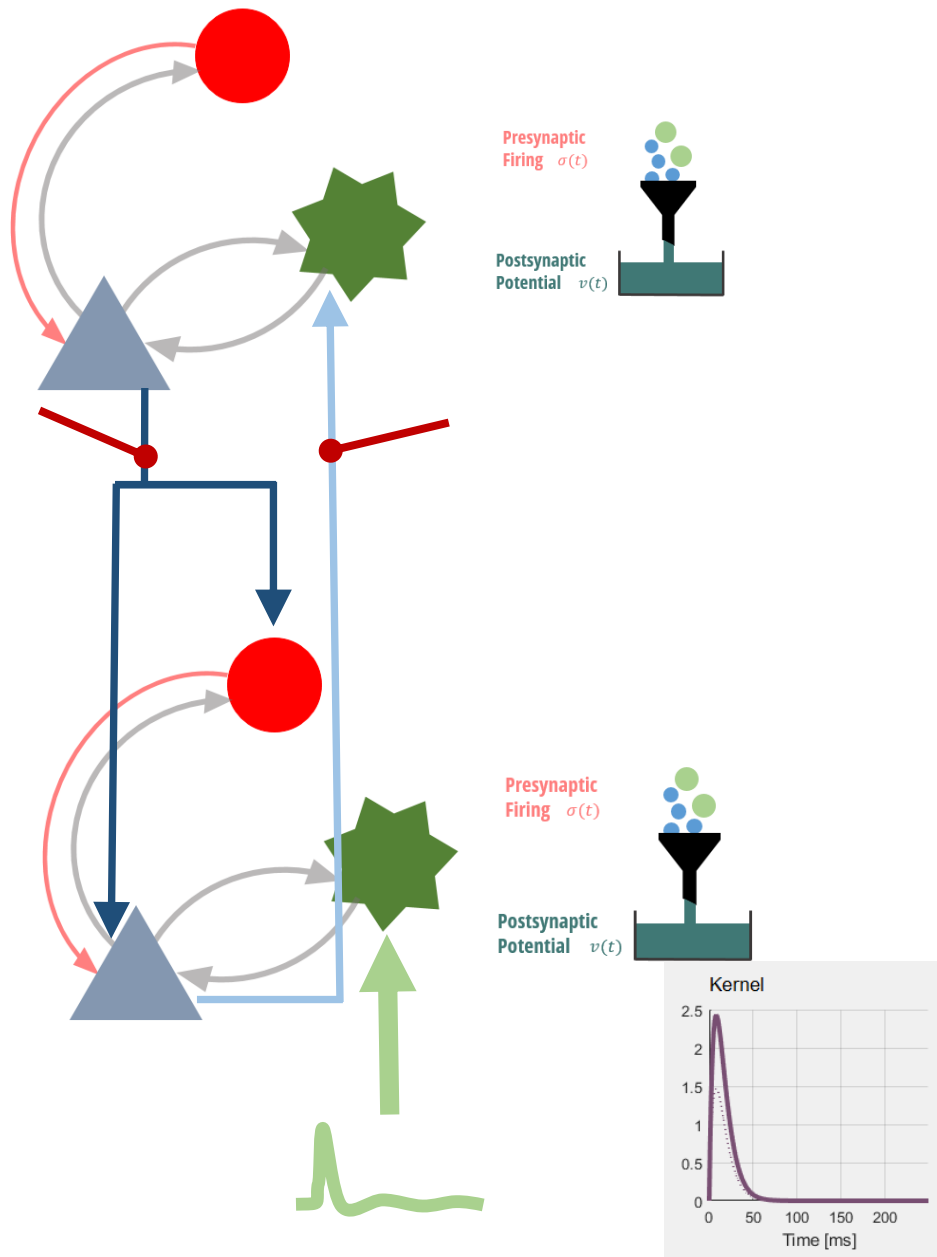


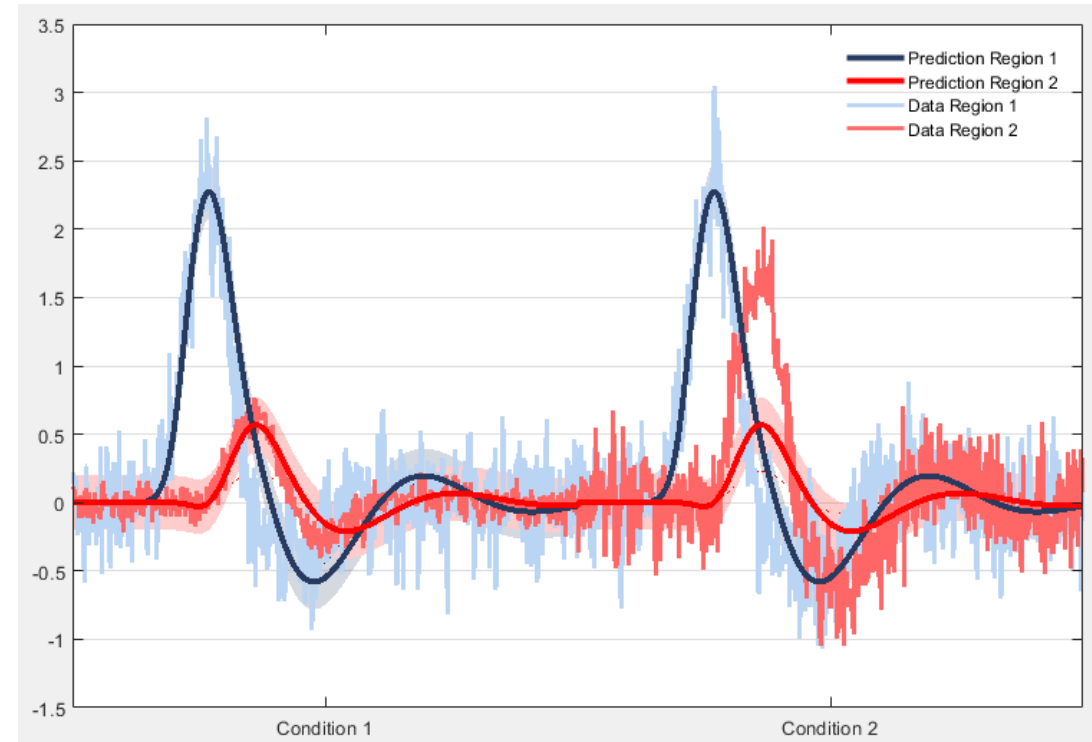
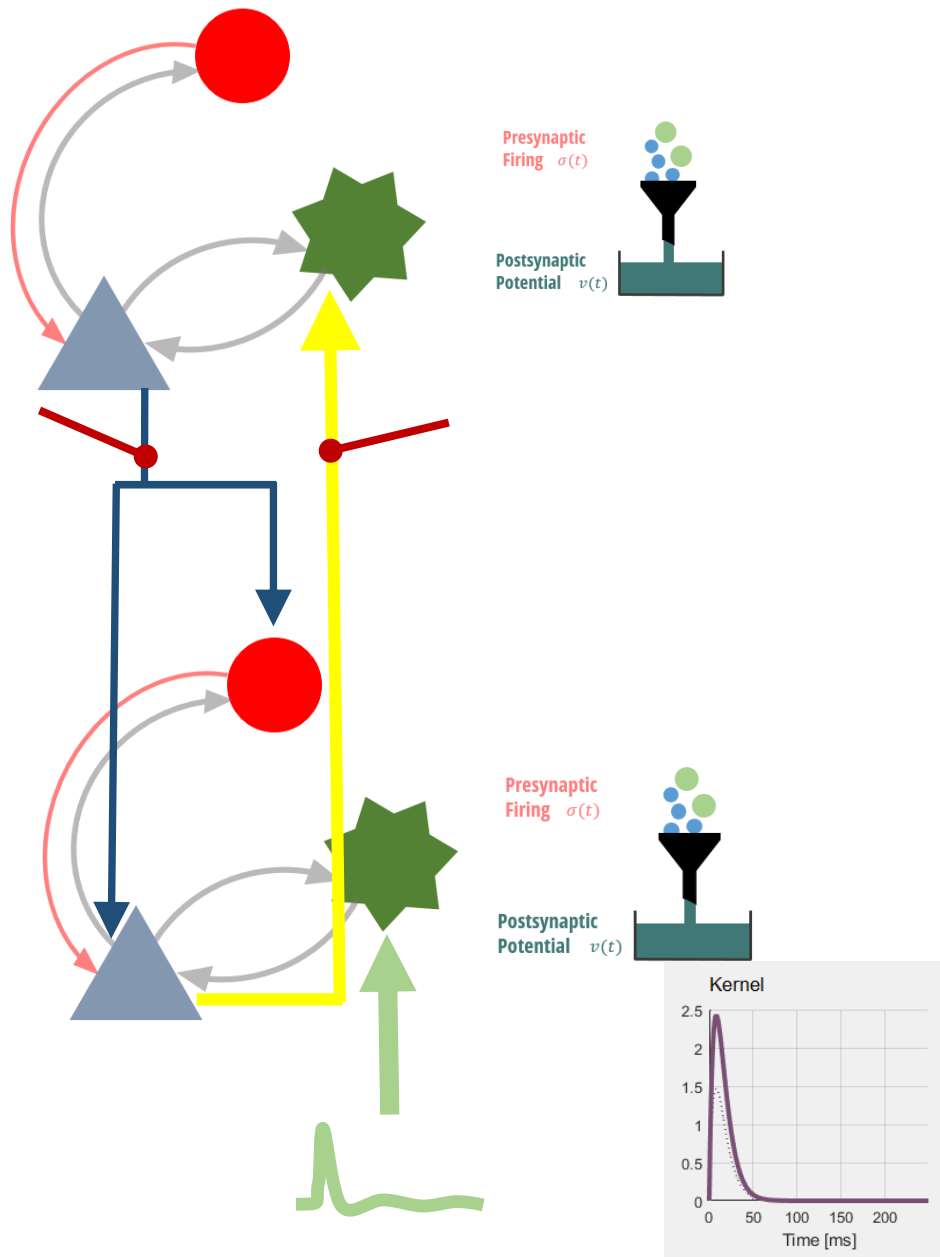


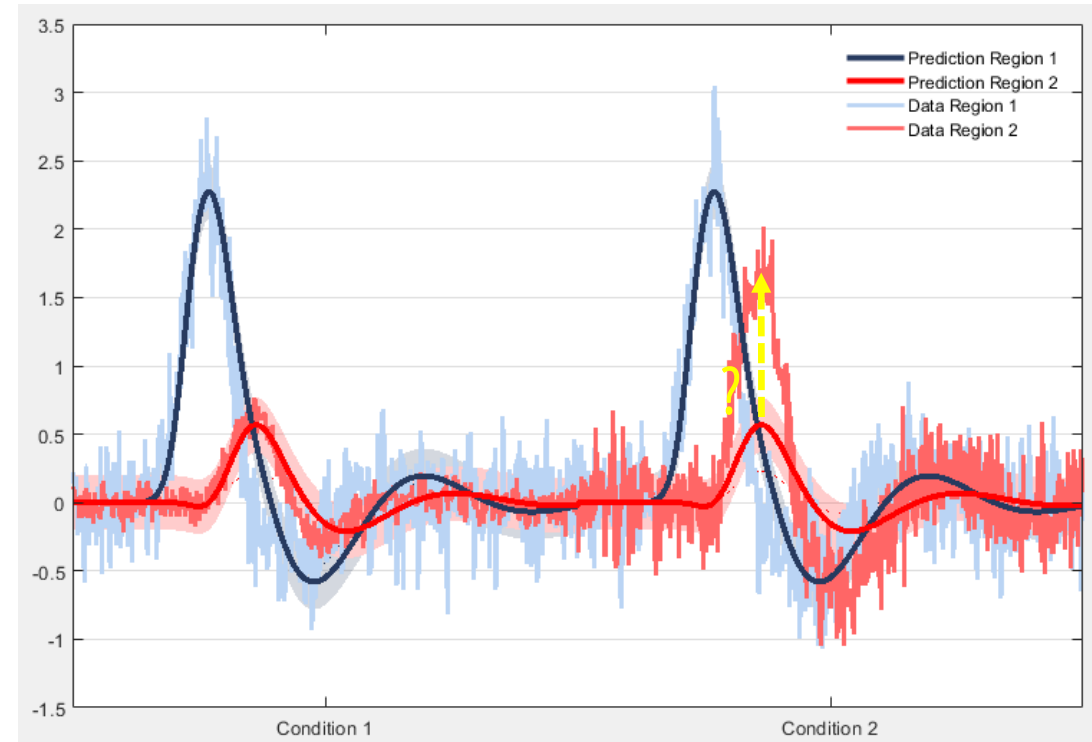
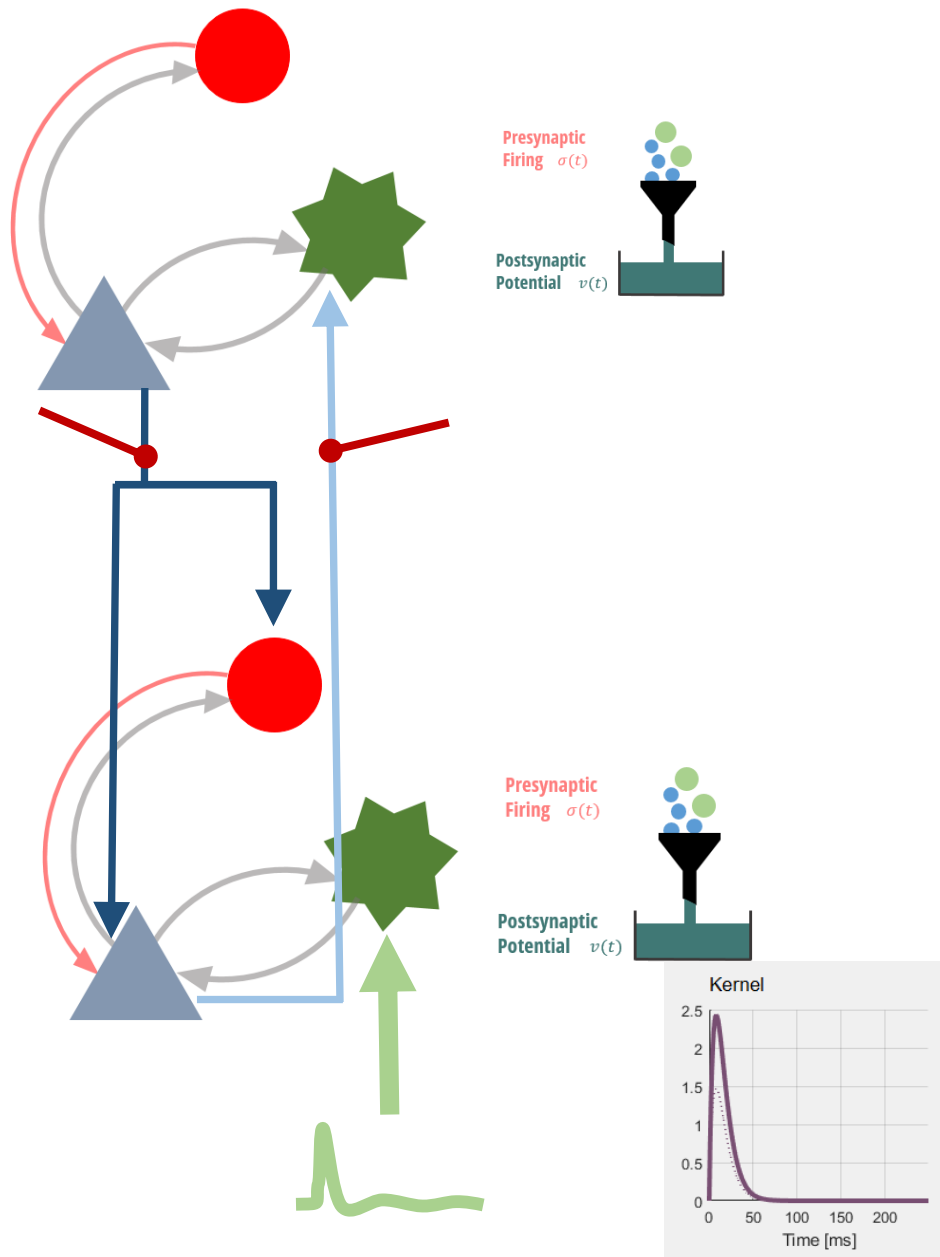


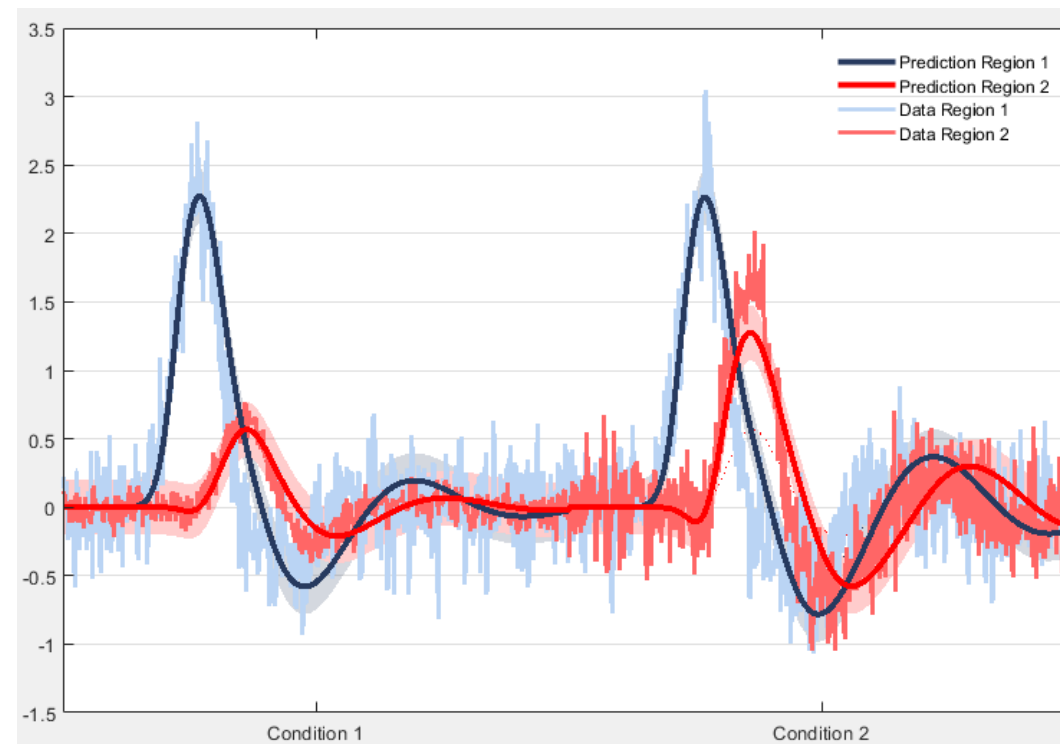
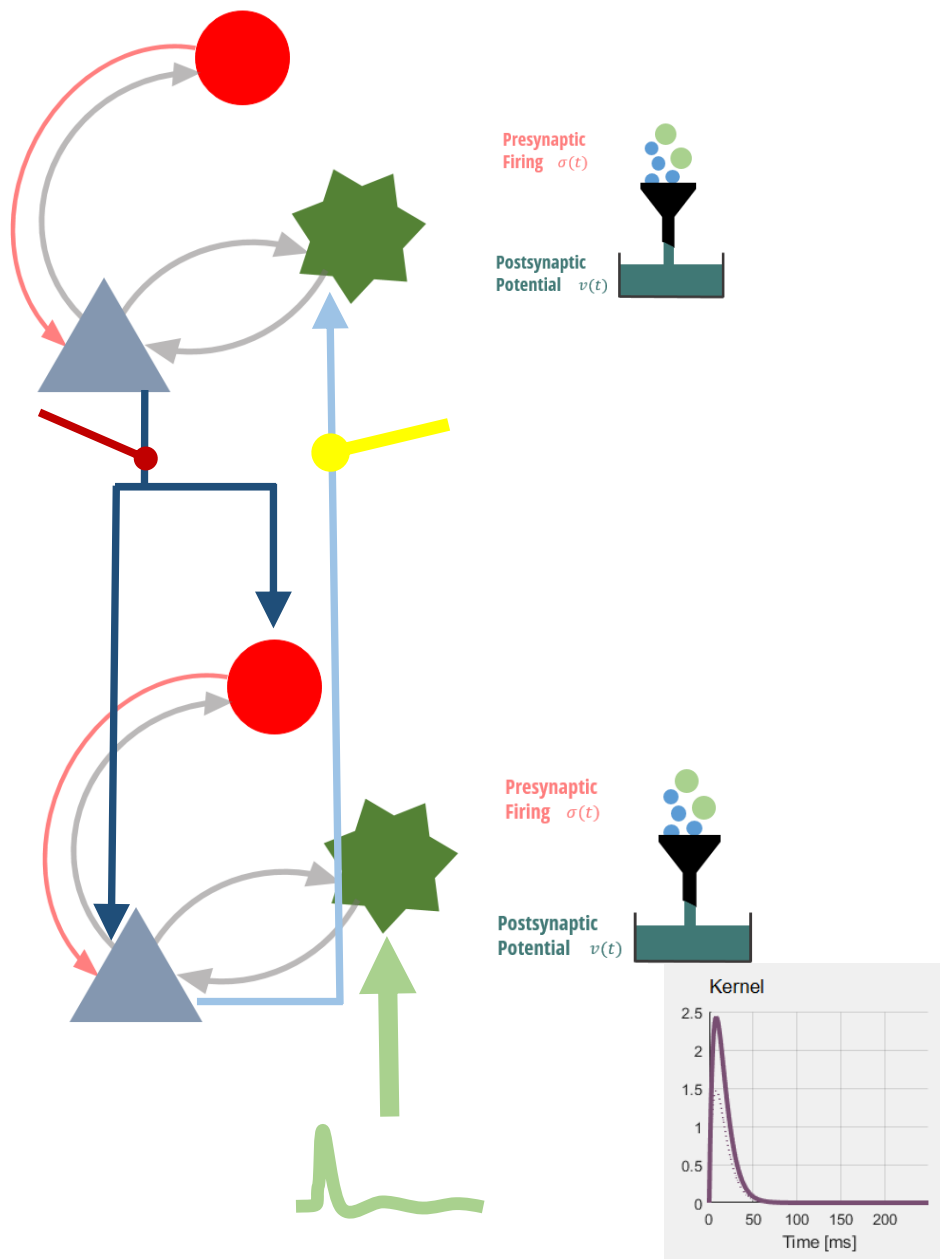












Inferring on parameters

- Multiple manipulations might lead to similar changes in data features
- One objective measure, which of the competing hypotheses is the best, is Bayesian model selection
- Inference will only be as good as your model and inversion machinery
- Always check results for pitfalls



Inferring on parameters

- Multiple manipulations might lead to similar changes in data features
- One objective measure, which of the competing hypotheses is the best, is Bayesian model selection
- Inference will only be as good as your model and inversion machinery
- Always check results for pitfalls:
 - 'Blindly' trusting inversion results
 - 'Overinterpretation' of single parameter estimates
 - Priors



V. Conductance Based DCM

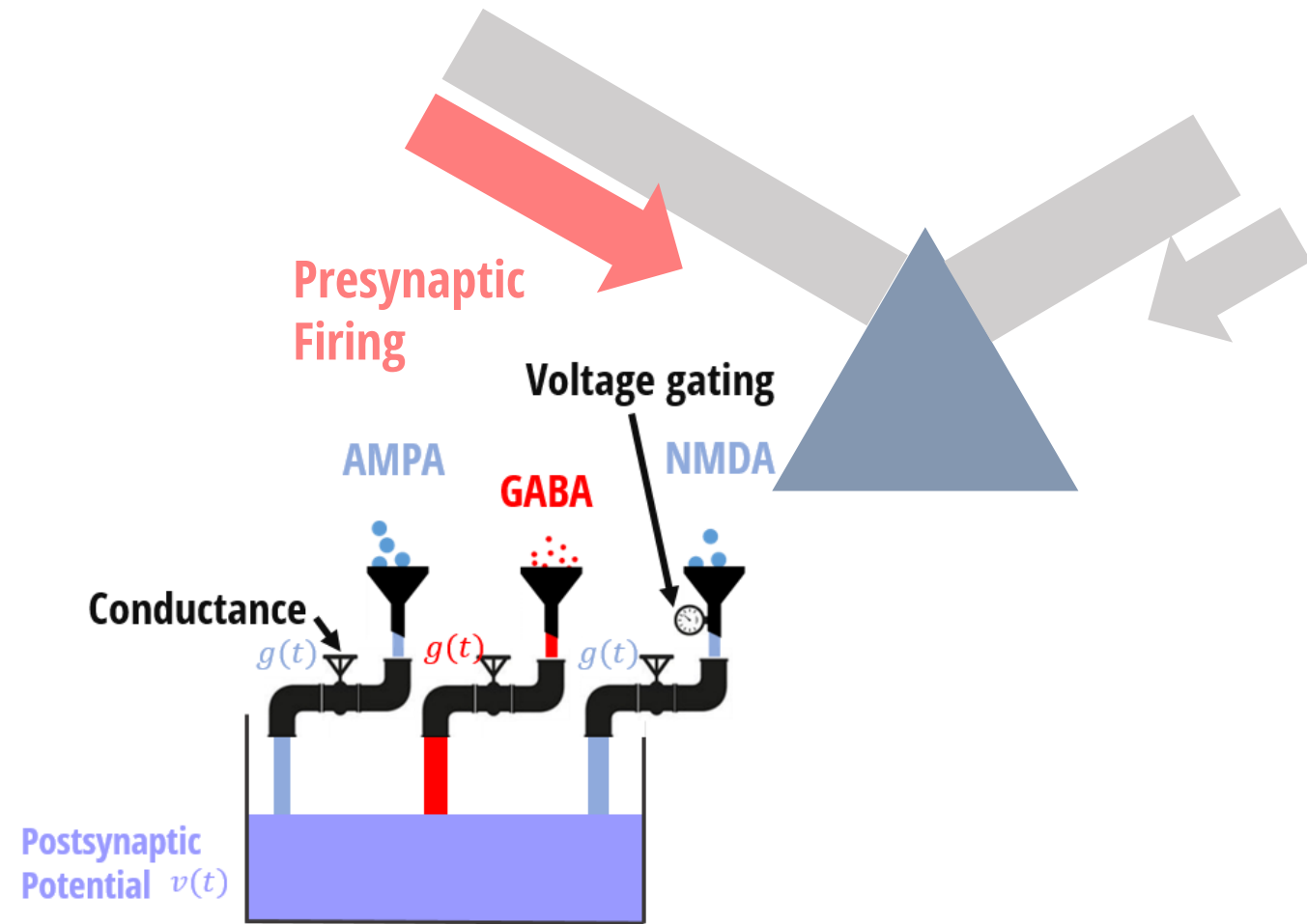
A very brief outlook on modeling synaptic mechanisms much more explicitly

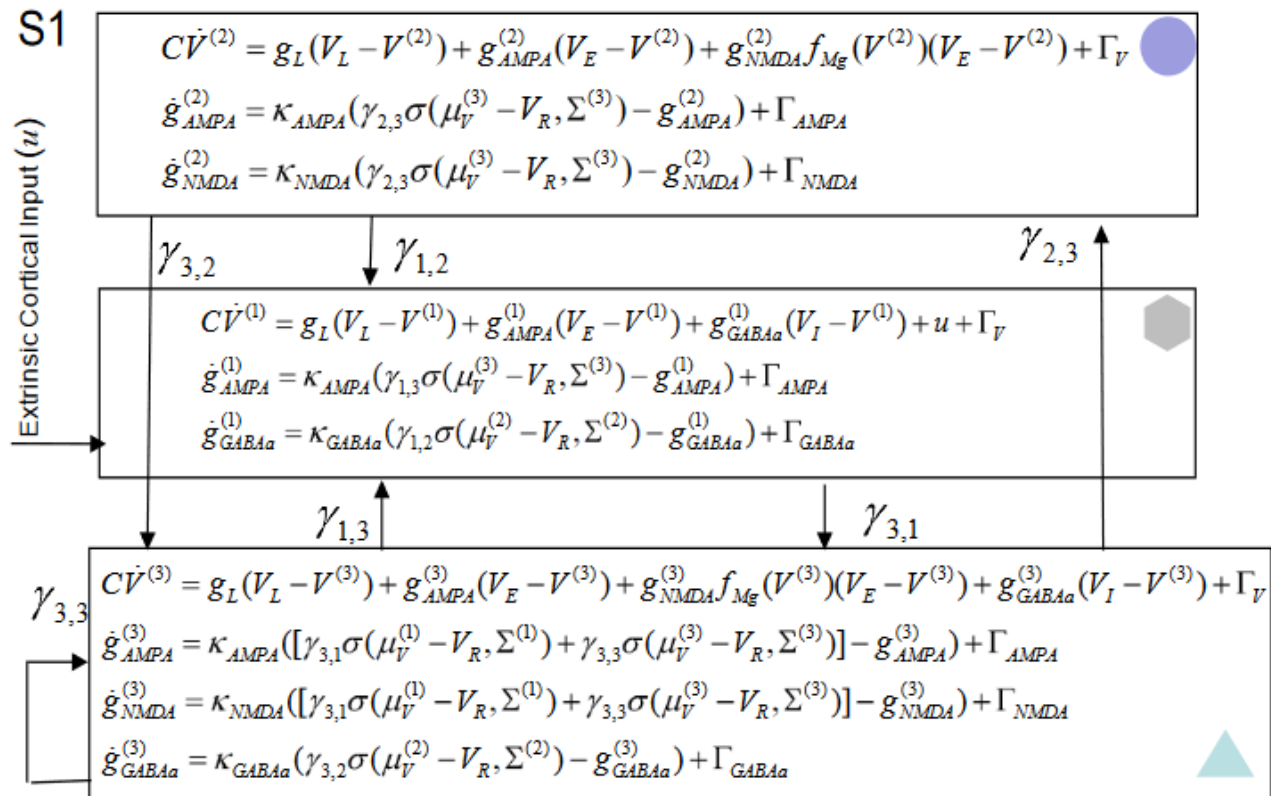


Conductance based DCM

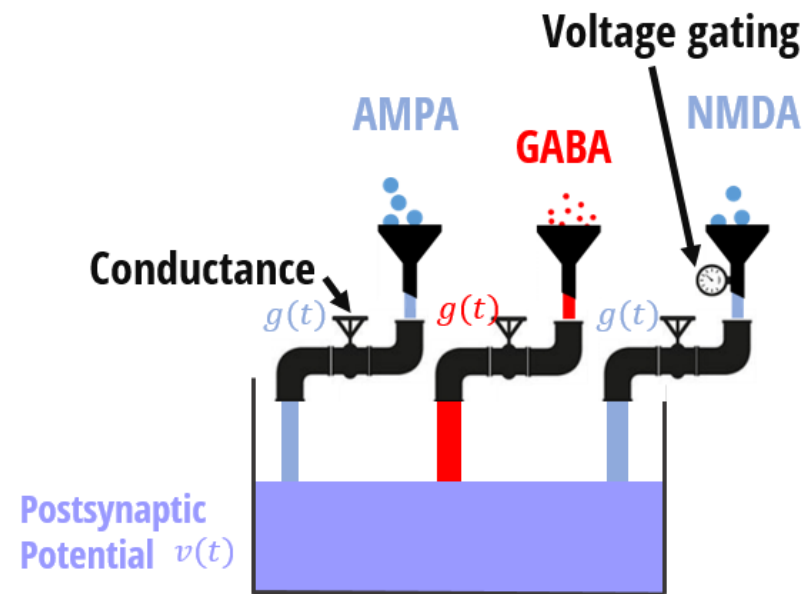
Morris- Lecar (1998)

Hodgkin and Huxley:
Current discharging the capacitor = Current passing through the resistor





Moran et al, Current Biology, 2011



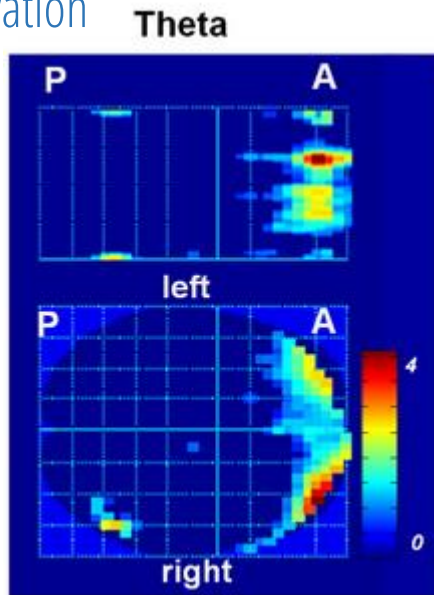
Current state of the population. How much firing arrives (and from where). NMDA Voltage Gate

$$\frac{dx}{dt} = f(x, A, H, C, \sigma, g(t), \Pi)$$

What the cell population is connected to. Resistance of the channels.

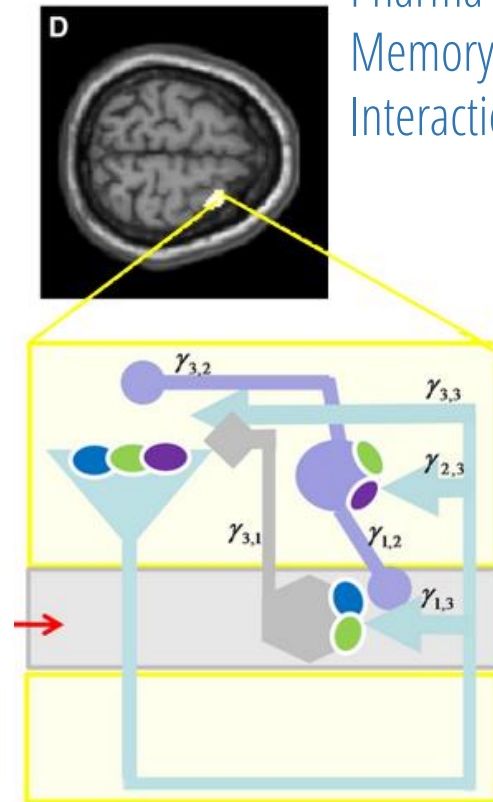
STUDY 1: INFERRING ON SYNAPTIC PARAMETERS

Working Memory
activation

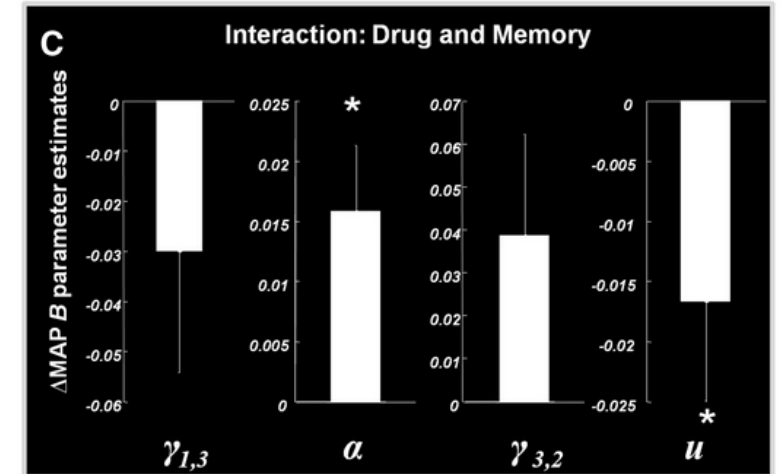


Moran et al, Current Biology, 2011

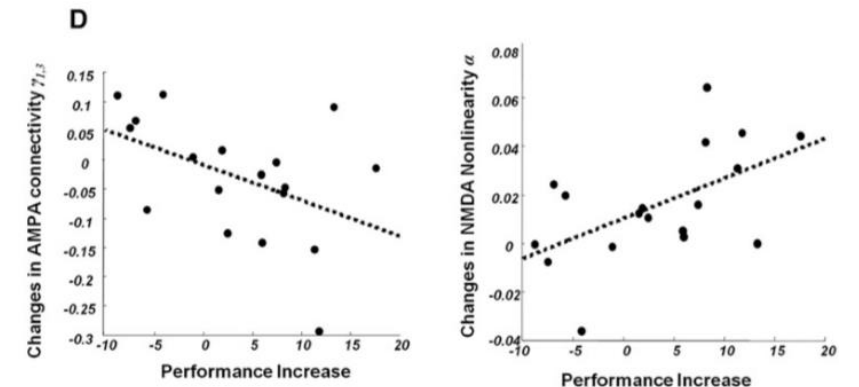
Pharma x
Memory
Interaction



Hypotheses about parameter
changes under pharma



Inferred parameters predict behavior

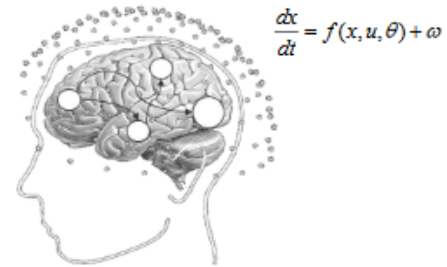


VI. Importance for computational psychiatry

Testable hypotheses from 'cheap', 'fast', 'simple' and non-invasive measurements



**1 Computational assays:
Models of disease mechanisms**



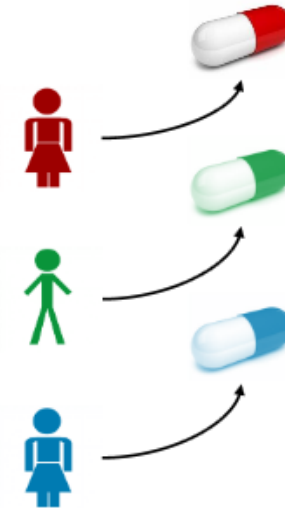
**2 Application to brain activity and
behaviour of individual patients**



**3 Detecting physiological subgroups
(based on inferred mechanisms)**



4 Individual treatment prediction



Stephan et al. 2015, *Neuron*

Data

(Hidden,) Neuronal Model

One can only infer on mechanisms, that have actually been modeled.

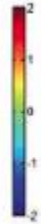
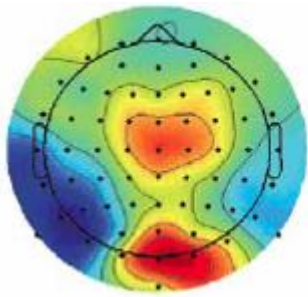
Inverse Problem: Inference

$$p(\theta, \Sigma | y)$$

macroscale

mesoscale

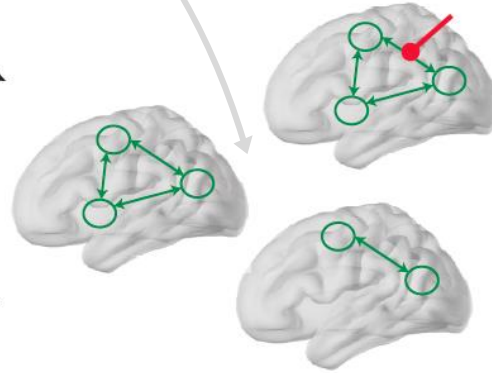
microscale



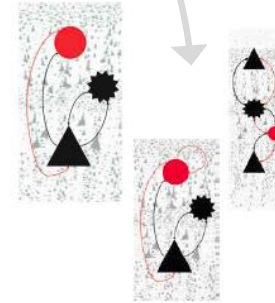
Forward model: Likelihood

$$y = g(x, \theta) + \varepsilon$$

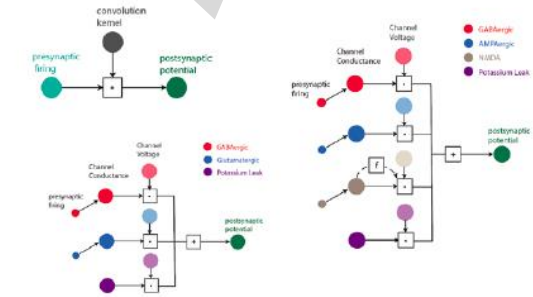
Data Features



Network of cortical sources and modulation of connection strength



Structure of the cortical column



Mechanisms governing generation of average post-synaptic potentials



Translational Neuromodeling Unit



University of
Zurich

ETH

Eidgenössische Technische Hochschule Zürich
Swiss Federal Institute of Technology Zurich

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JAKOB HEINZLE AND KLAAS ENNO STEPHAN
FOR SOME OF THE SLIDES!



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Zurich ^{UZH}

ETH

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Swiss Federal Institute of Technology Zurich