

# DCM for Evoked Responses

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Department of Computer Science  
University College London

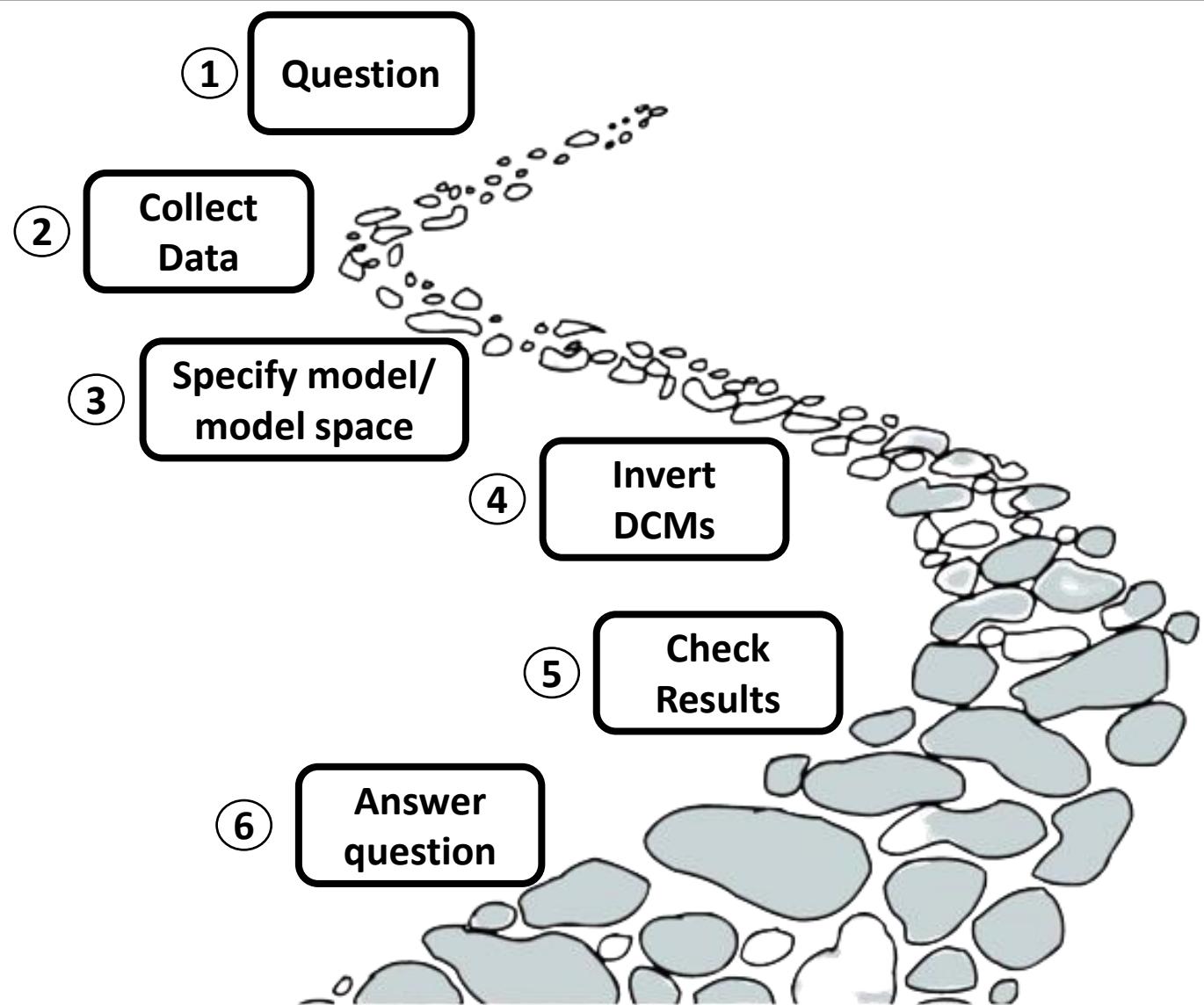
# SPM for MEG/EEG Course

## Course Agenda – Thursday 1<sup>st</sup> June

9.30-10.00	Day 3: Welcome and Registration  Chair: Benjy Barnett	Registration (Conservatory) Refreshments (Boardroom)
10.00 - 11.30	The principles of dynamic causal modelling  Multimodal DCM	Amirhossein Jafarian
11.30 - 12.35	DCM for evoked responses	Daniel Hauke
12.35 -13.35	DCM for Cross-Spectral Densities	Dimitris Pinotsis
13.35-14.35	Lunch	Boardroom
	Chair: Mansoureh Fahimi	
14.35 – 15.20	DCM demo	Julia Rodriguez-Sanchez
15.20 -16.35	Bayesian model selection and averaging	Peter Zeidman
16.35-17.35	Q&A clinic	Karl Friston

# Outlook - The DCM analysis path

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# Which questions can DCM for evoked responses answer?

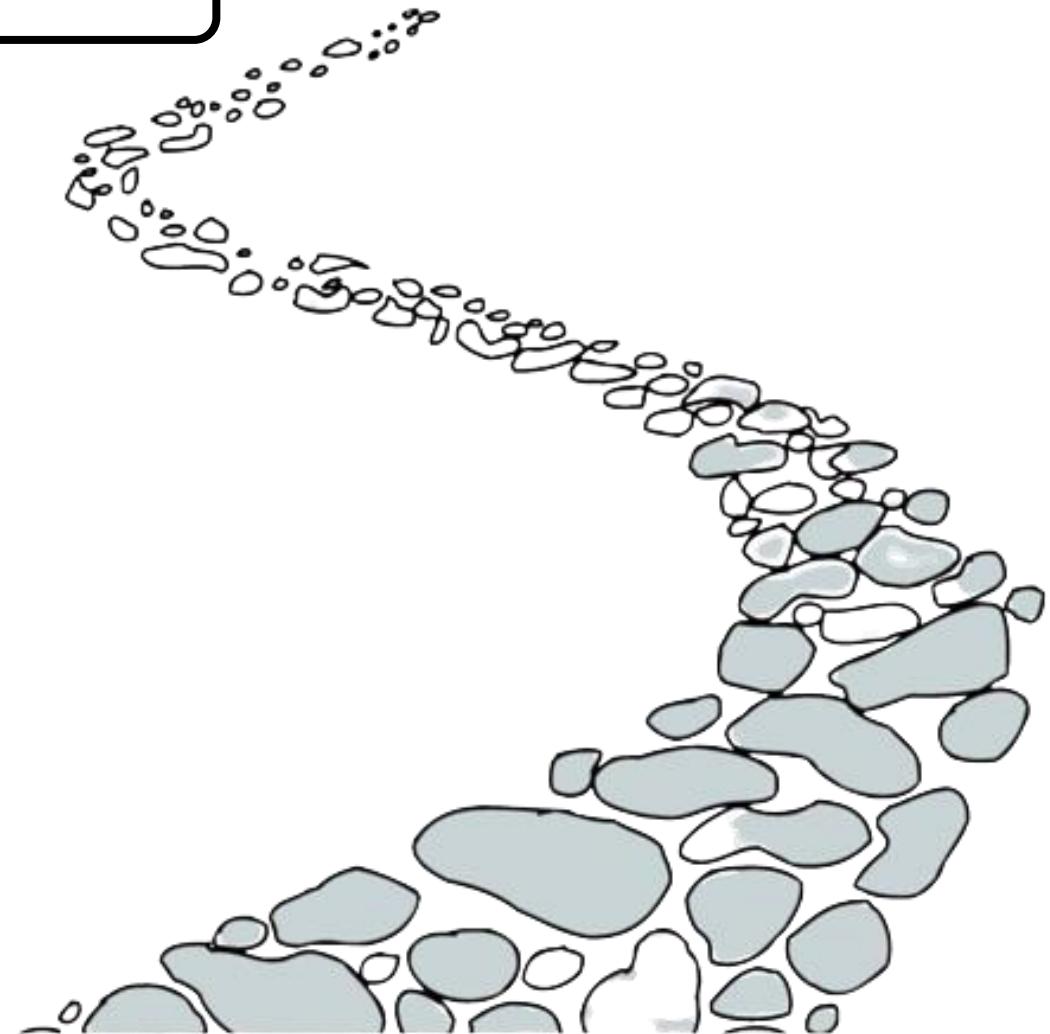
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## Good questions

- Does the network with regions A, B and C explain my data better than the network with regions A and B?
- Are regions A and B linked in a bottom-up, top-down or recurrent manner?
- How does my experimental manipulation change the effective connectivity between regions?
- Or within a region?
- What EEG signal would I expect if I increase the connectivity even further?

①

Question

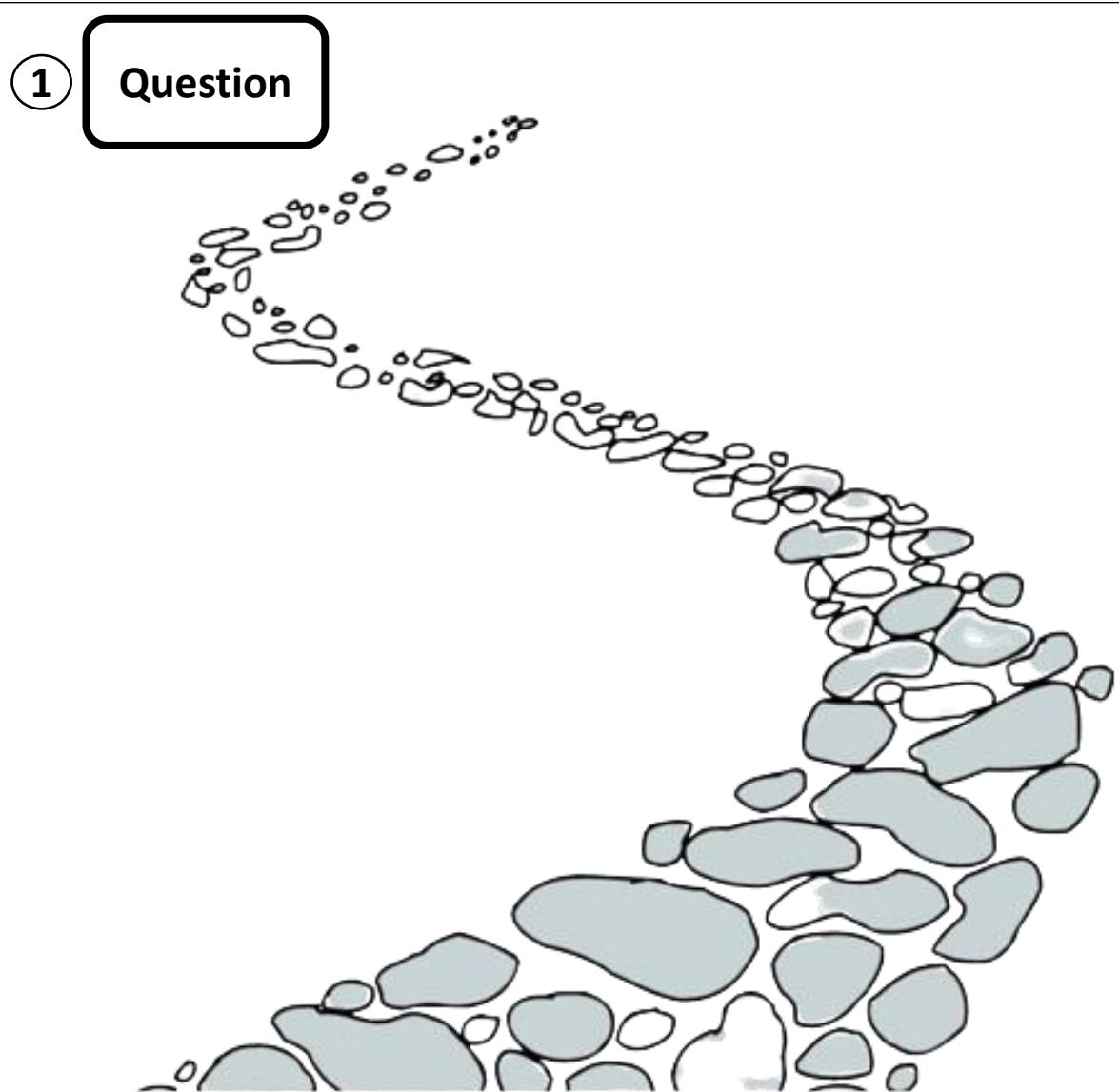


# Which questions can DCM for evoked responses answer?

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## Not-so-good questions

- We did not find an effect in our ERP analysis - Can you model the data with DCM to find some effects to publish a cool paper?
- How does the connectivity change within our 200-region network?
- How does the connectivity change within our 20-region network?

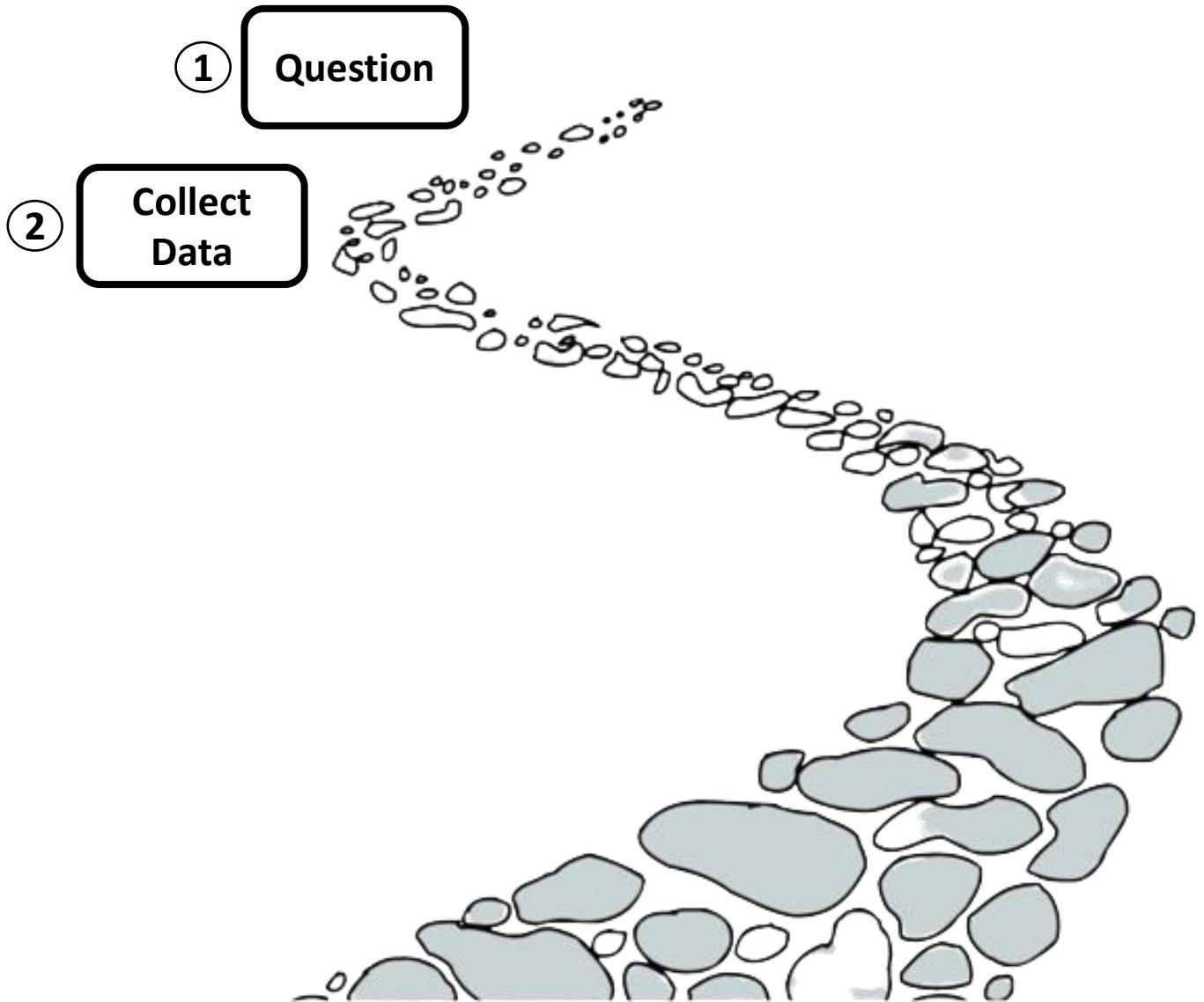


# Data collection

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## Preprocessing

- Downsample (e.g., 100 Hz)
- Filter (e.g., 1-40 Hz)
- Epoch
- Remove artefacts
- Average
  - Per subject
  - Grand average

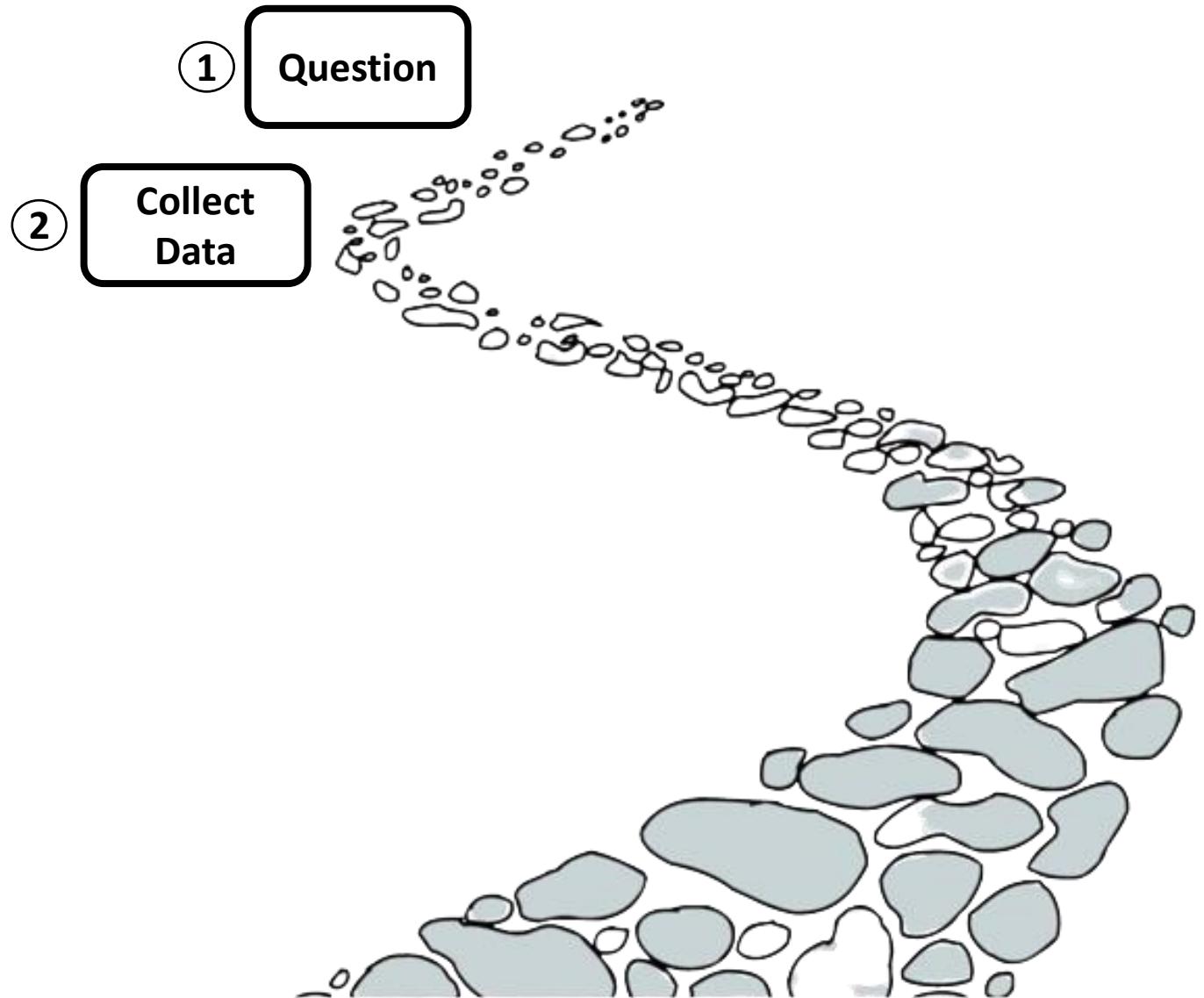


# Data collection

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## Classical analysis

- Make sure there are effects!
- DCM is used to **explain** these effects

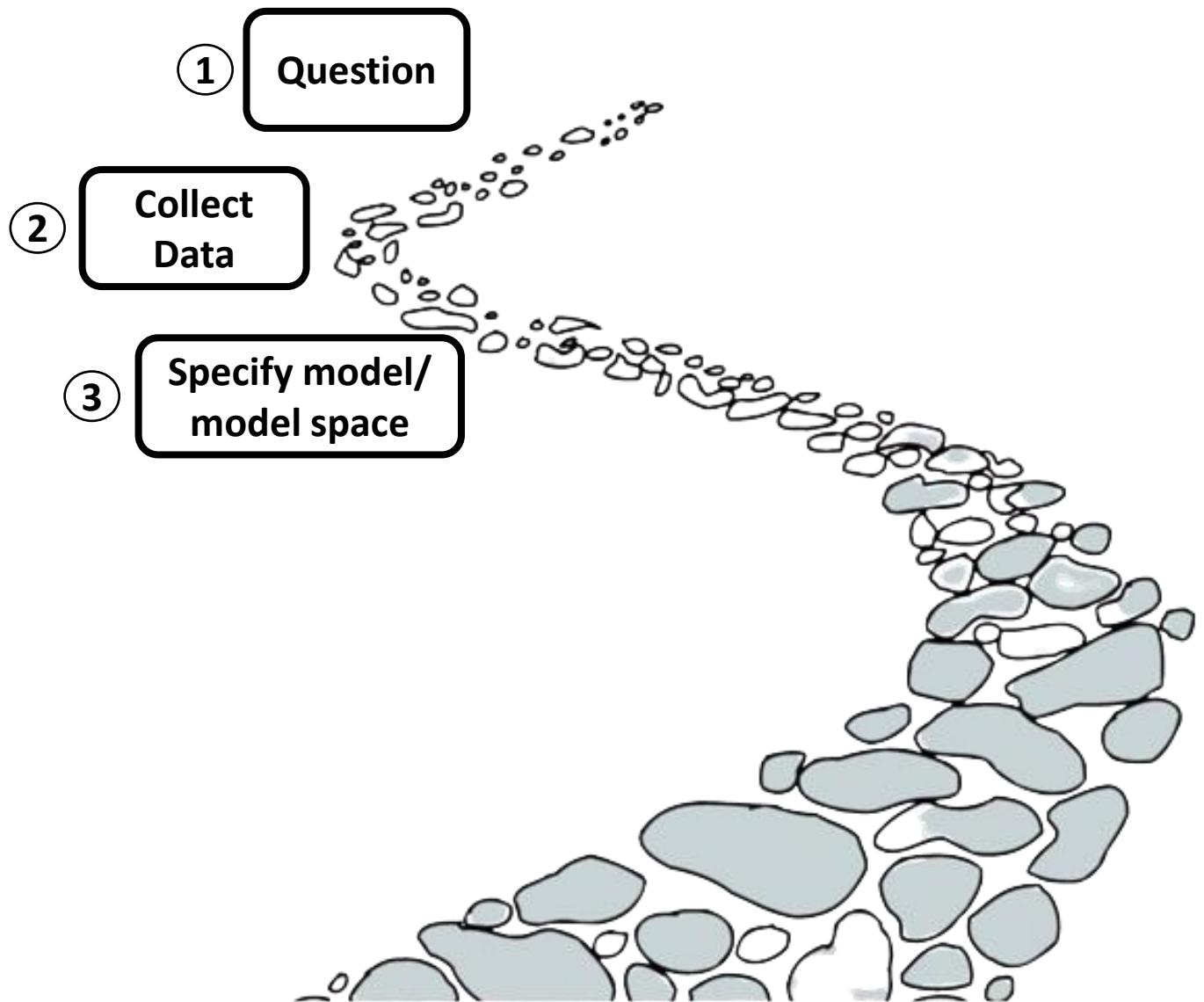


# Specify model/model space

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## Steps

- Translate your question into a model comparison or a parameter inference problem
- Select regions
- Select a variant of DCM
- Example: The “ERP” model
- Specify connectivity architecture

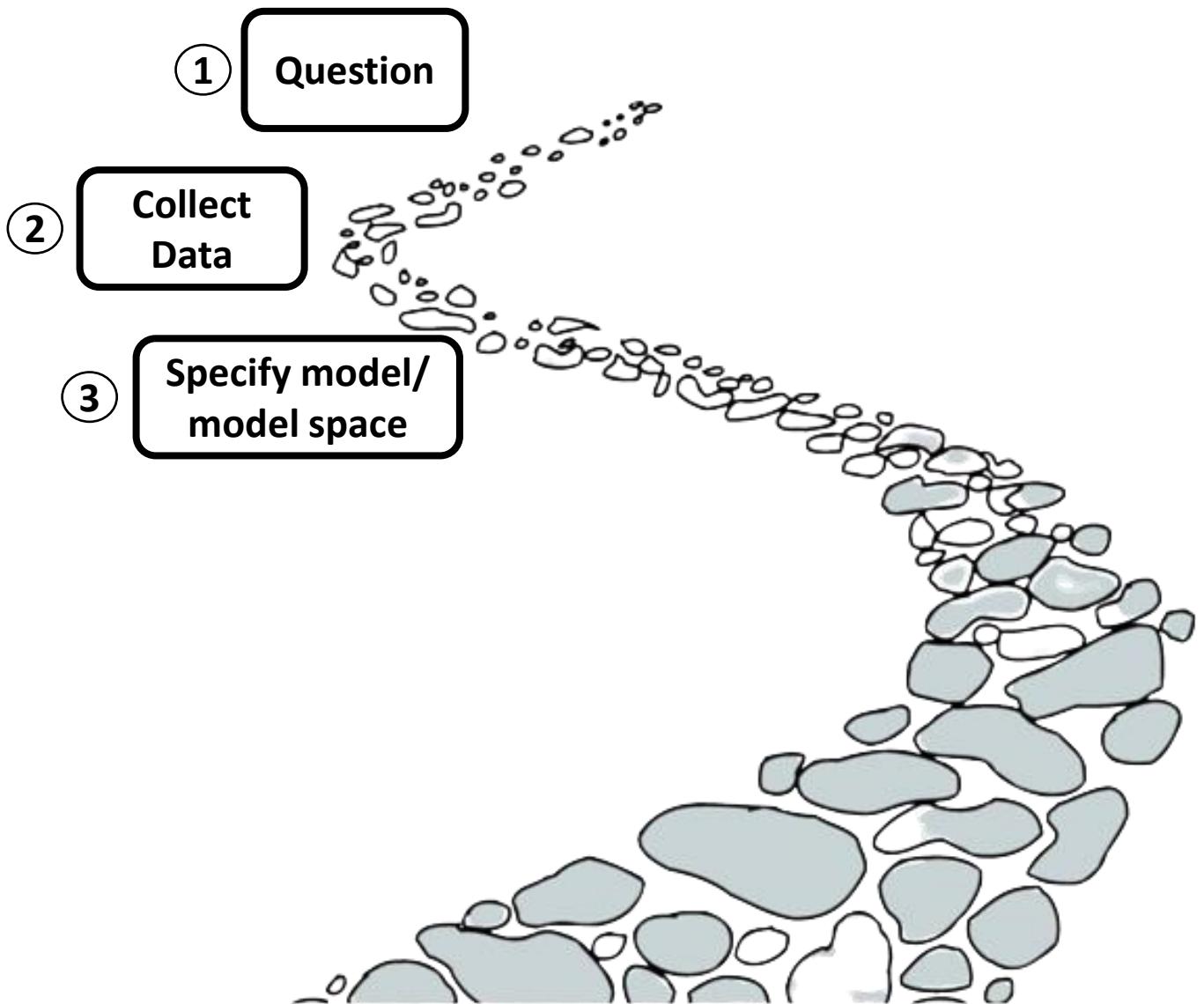


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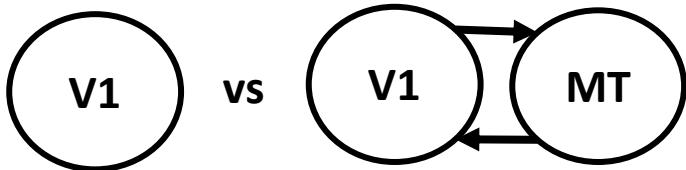


# Translate your question

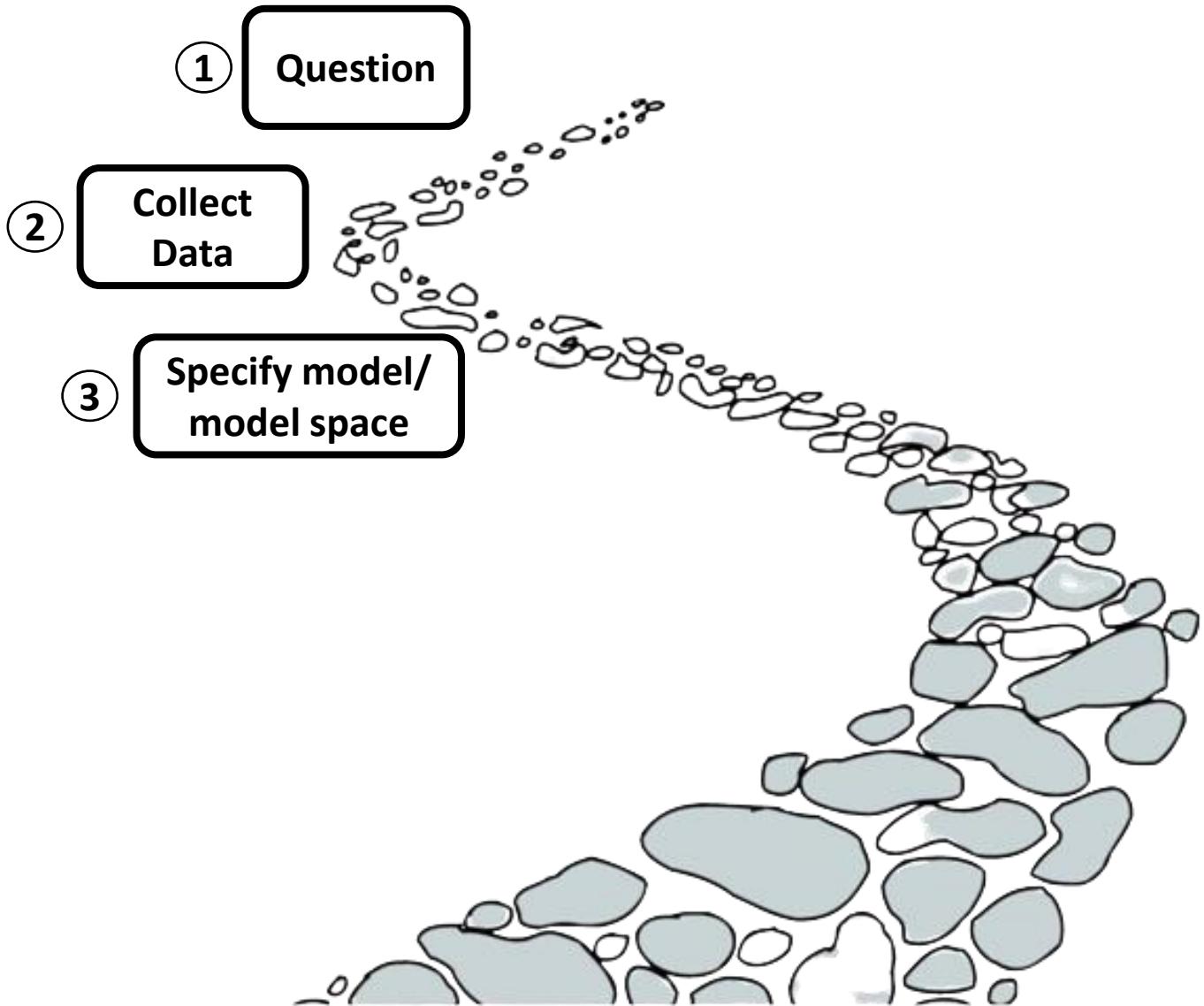
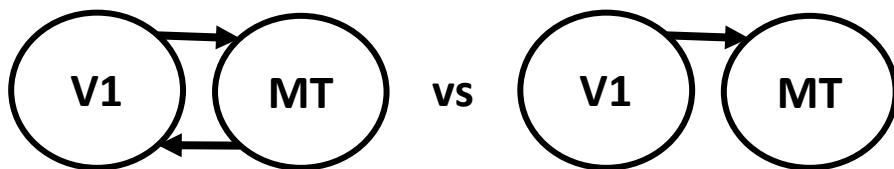
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## Model comparison example

- Is my task activating MT and V1 or only V1?



- Are backward connections switched off when individuals are sleeping?

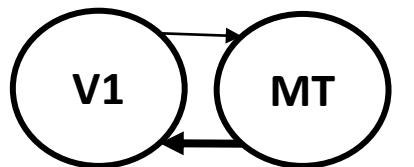


# Translate your question

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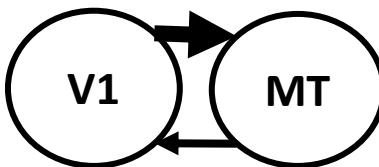
## Parameter inference example

- Is attention increasing forward connectivity?

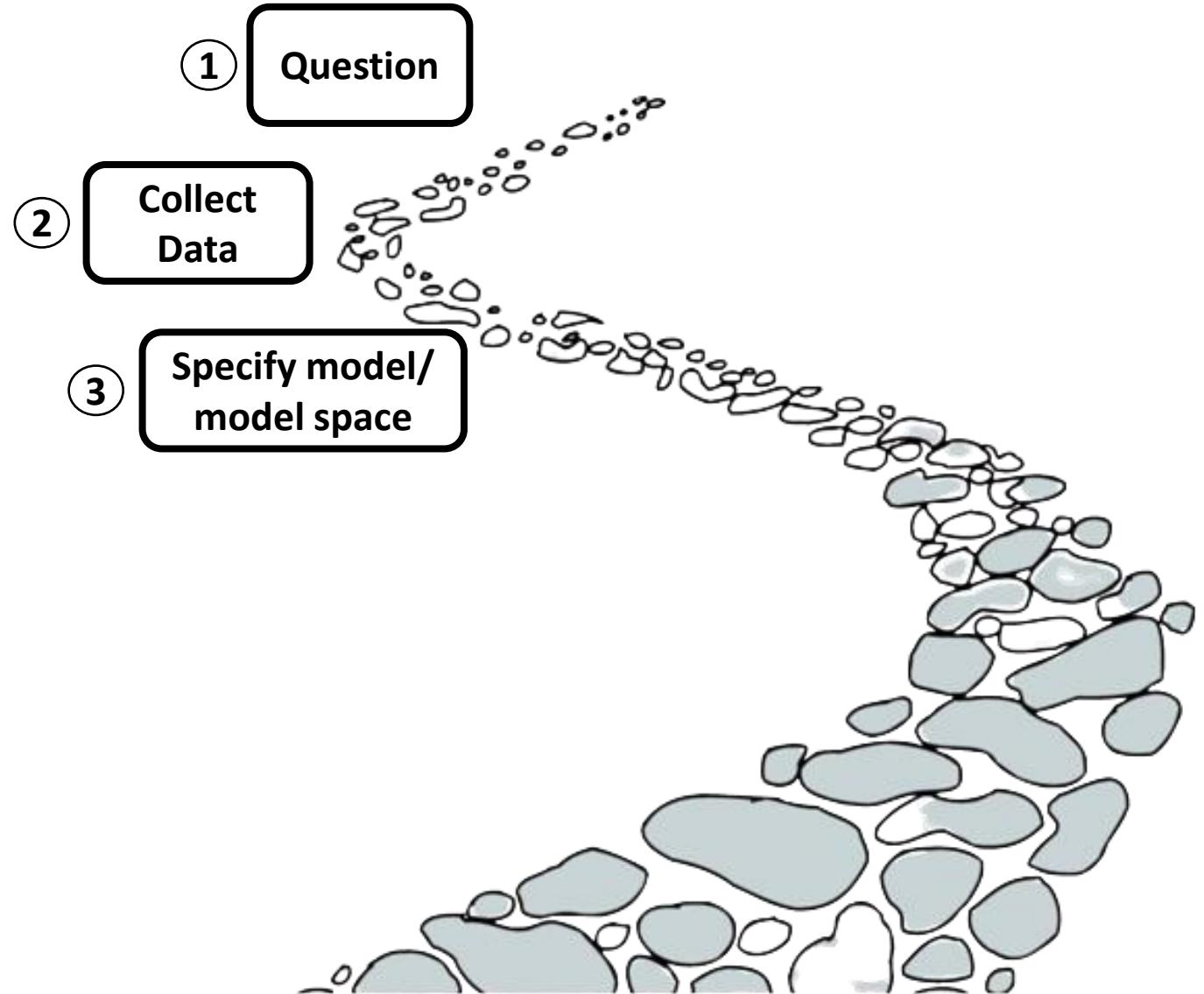


No attention

vs



Attention

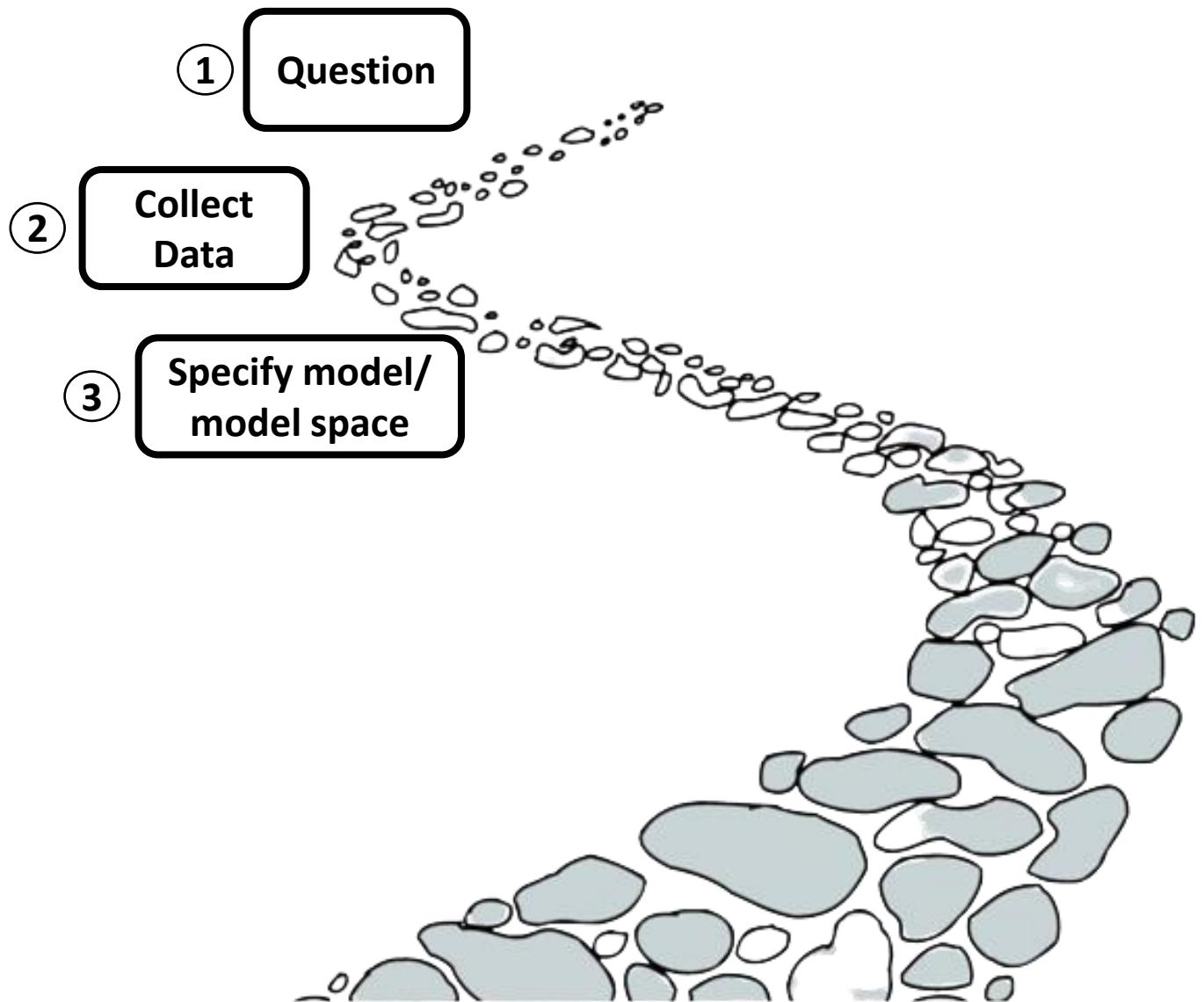


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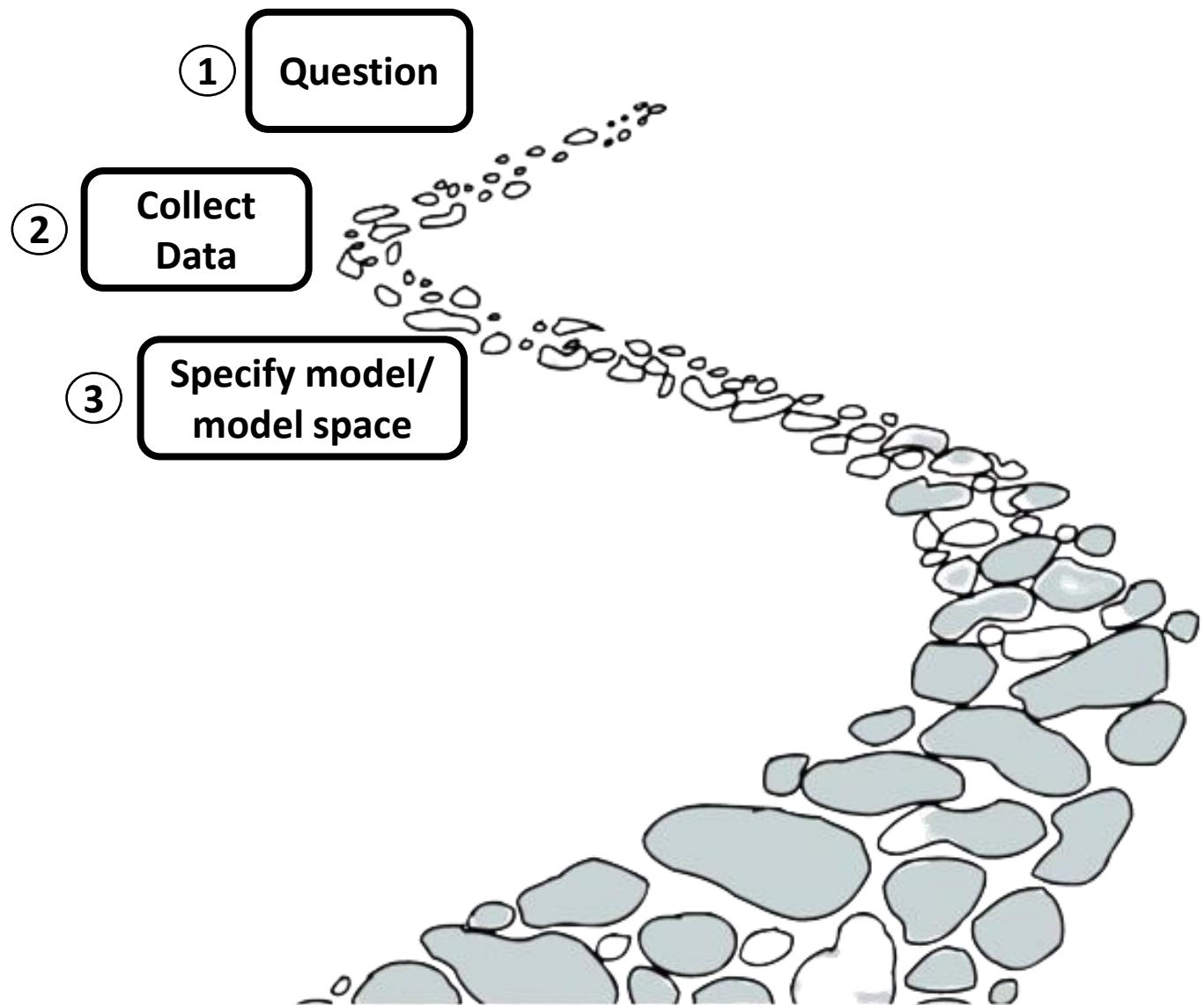


# Select regions

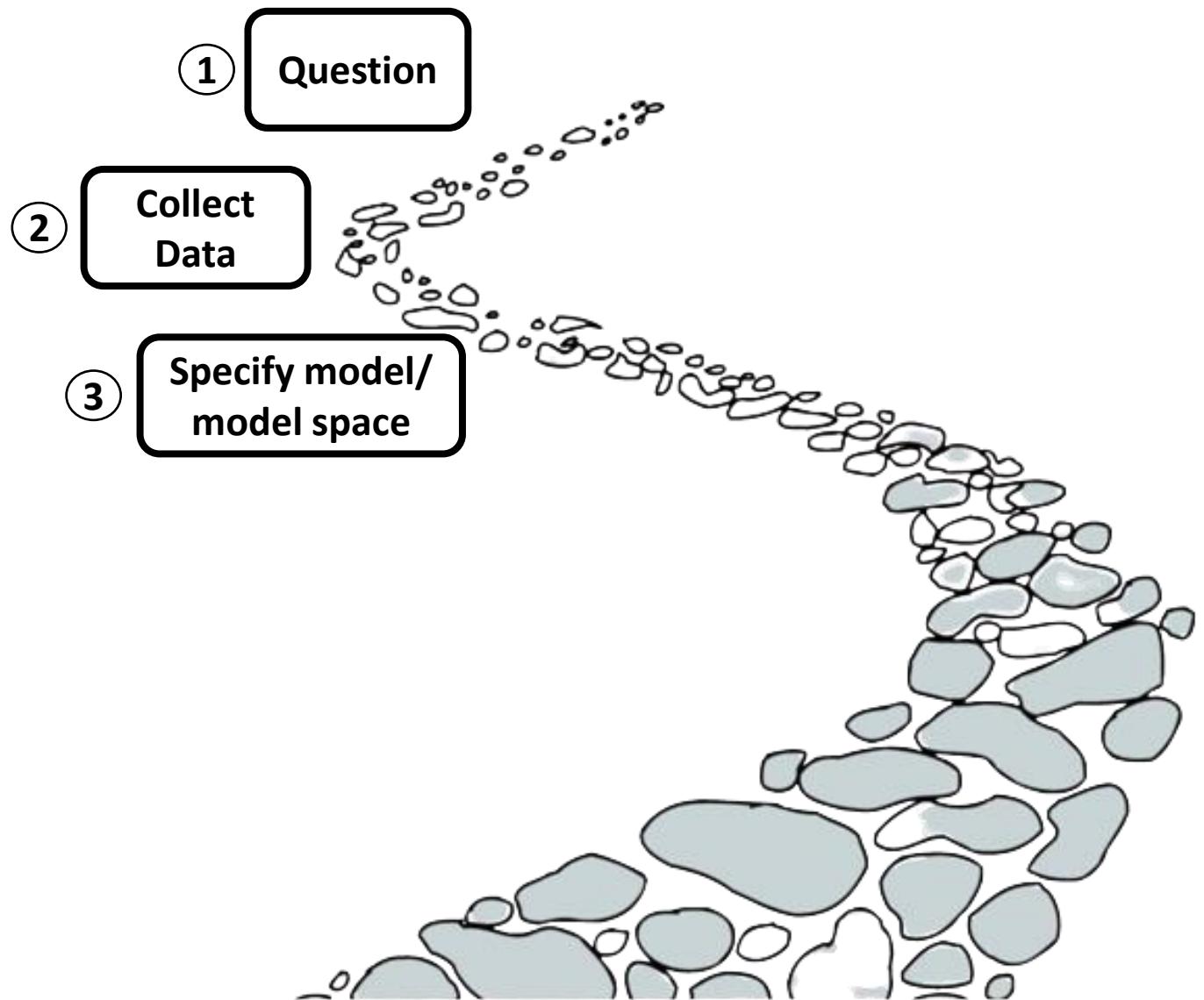
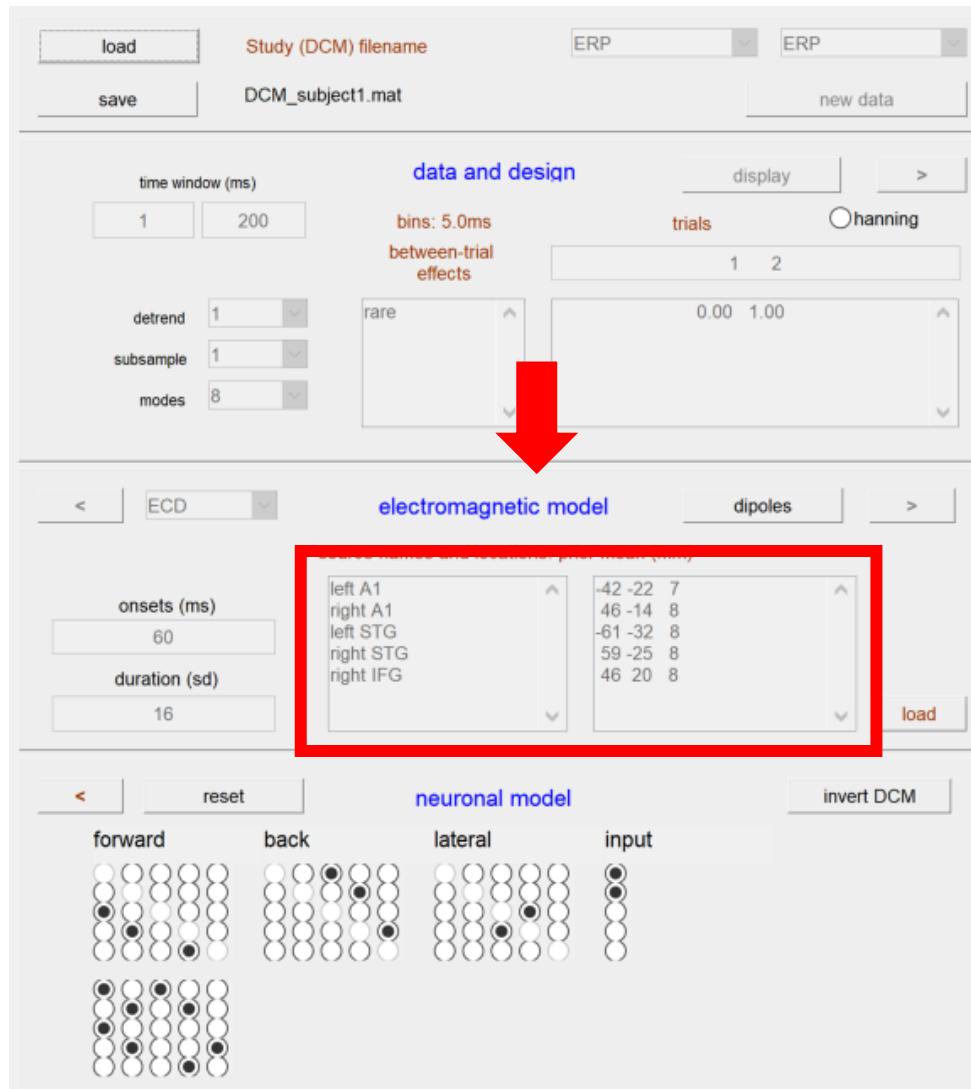
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## Ways to select regions

- Literature
- MRI
- Source reconstruction
- Dipole fitting



# The DCM analysis path

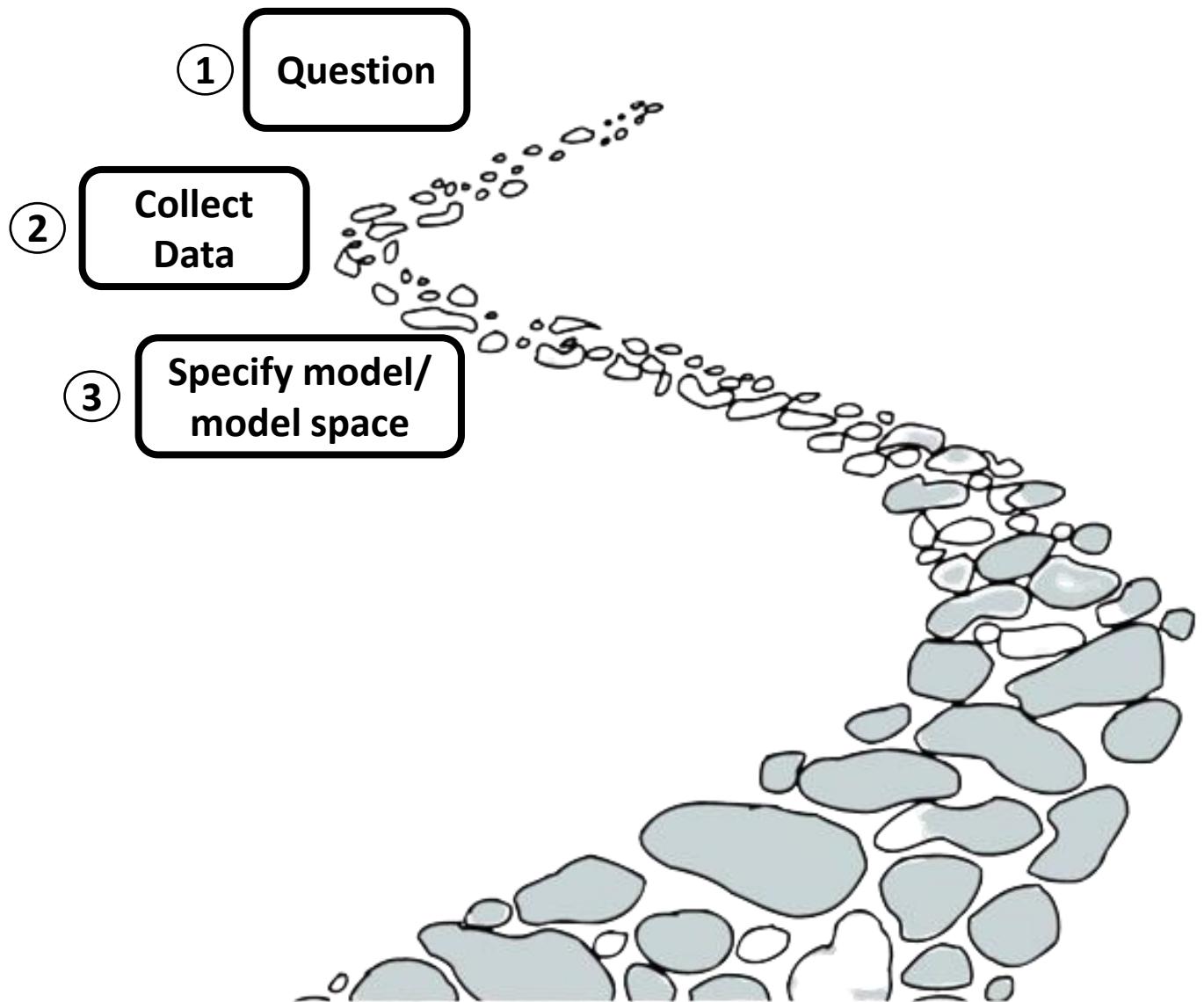


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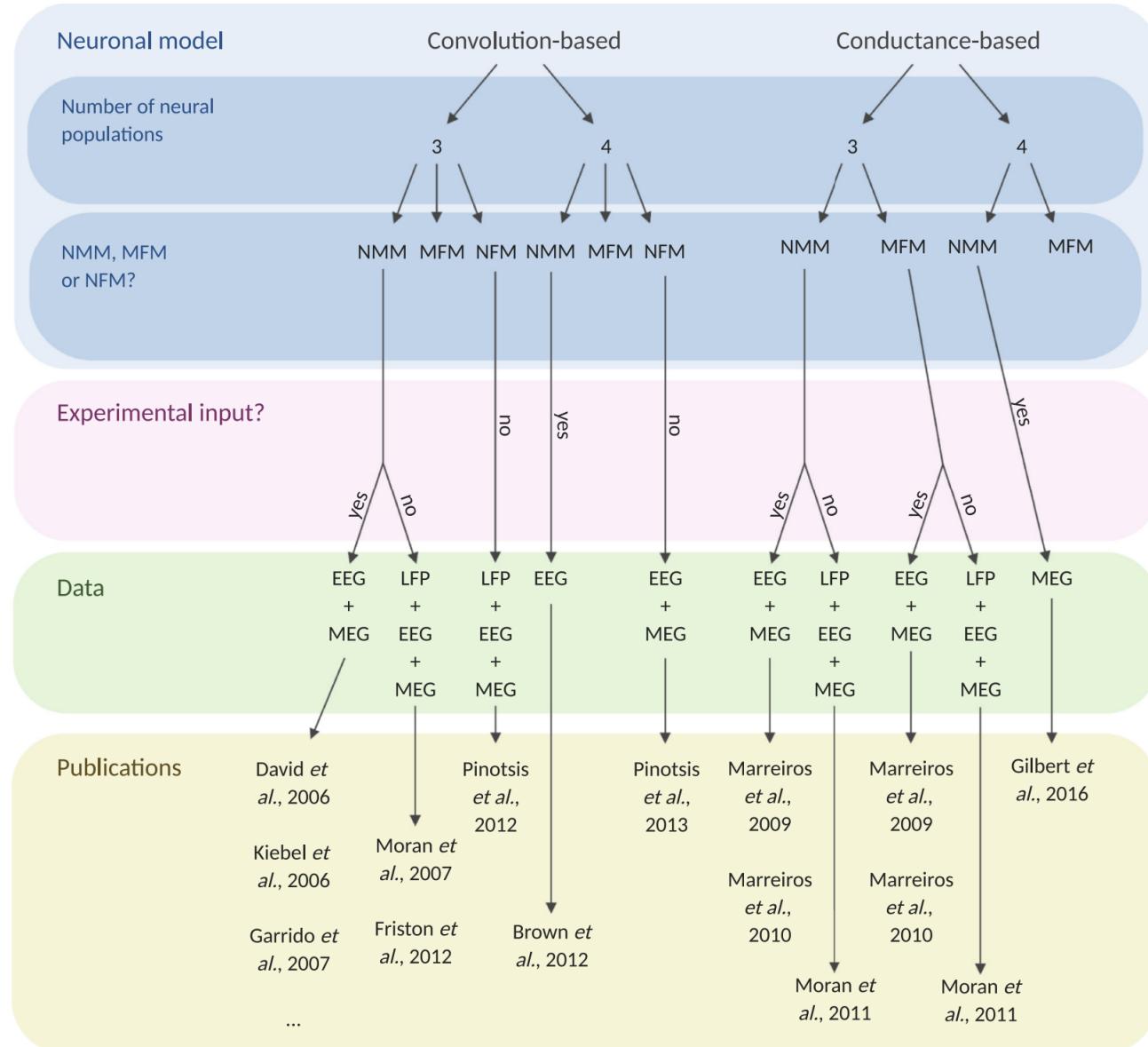
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## Steps

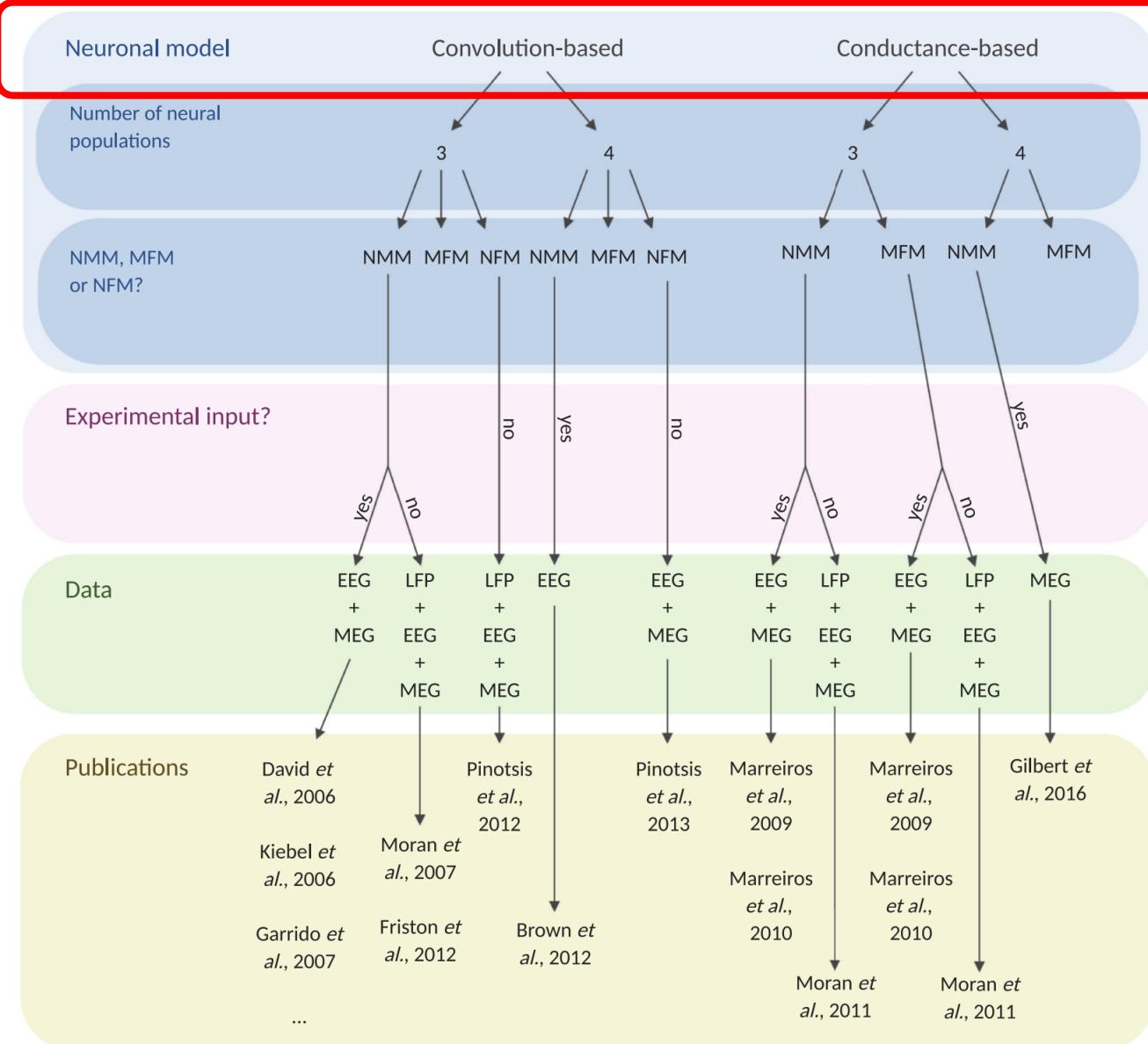
- Translate your question into a model comparison or a parameter inference problem
- Select regions
- **Select a variant of DCM**
- Example: The “ERP” model
- Specify connectivity architecture



# Select a variant of DCM for evoked responses



# Select a variant of DCM for evoked responses



## Convolution-based

SPM: ERP, SEP, LFP, CMC, NFM

- Wilson & Cowan (1973), Jansen & Ritt (1995)

## Conductance-based

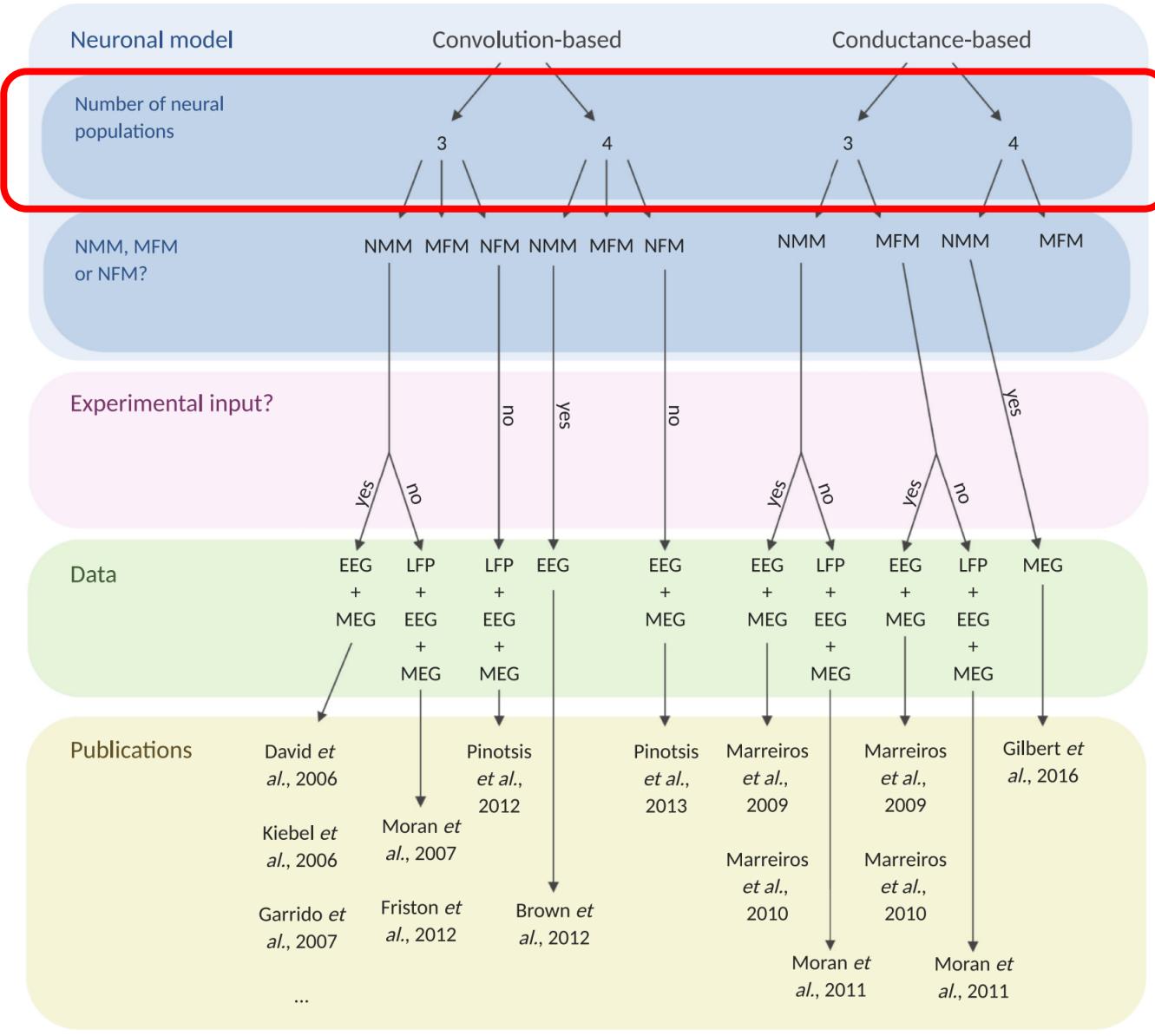
SPM: NMM, NFM, CMM, NMDA, CMM\_NMDA

- Hodgkin & Huxley (1952), Morris & Lecar (1981)

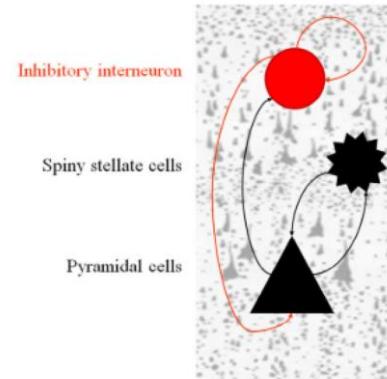
## How to choose?

- Are you interested in ion channels: conductance-based
- Otherwise: Convolution

# Select a variant of DCM for evoked responses



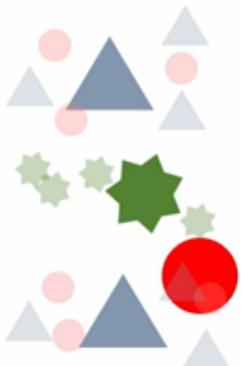
## **3 populations** (ERP, SEP, LFP, NFM)



Moran et al. (2013),  
*Front. Comput. Neurosci.*

## 4 populations

(CMM, CMC, CMM\_NMDA)



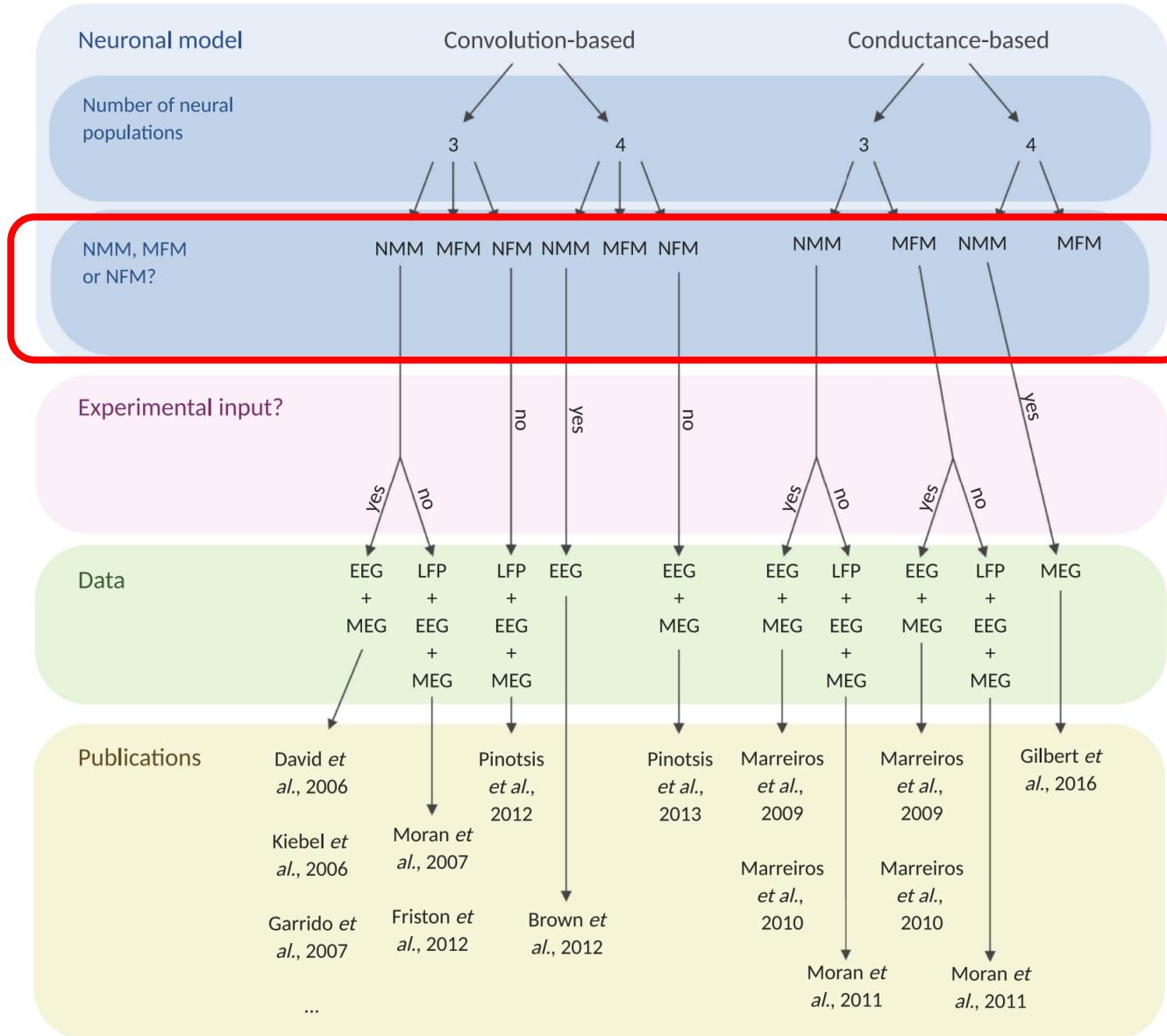
Schöbi (2019), CPC Lecture

# How to choose?

- Do you want to test predictive coding?
  - Do you expect specific effects in either deep or superficial pyramidal cells?

=> Canonical microcircuit

# Select a variant of DCM for evoked responses



## MFM: Mean-field model

- Considers mean and covariance

## NMM: Neural mass model

- Describes populations by their mean activity (special case of MFM)

## NFM: Neural field model

- Considers spatial dimension

## How to choose?

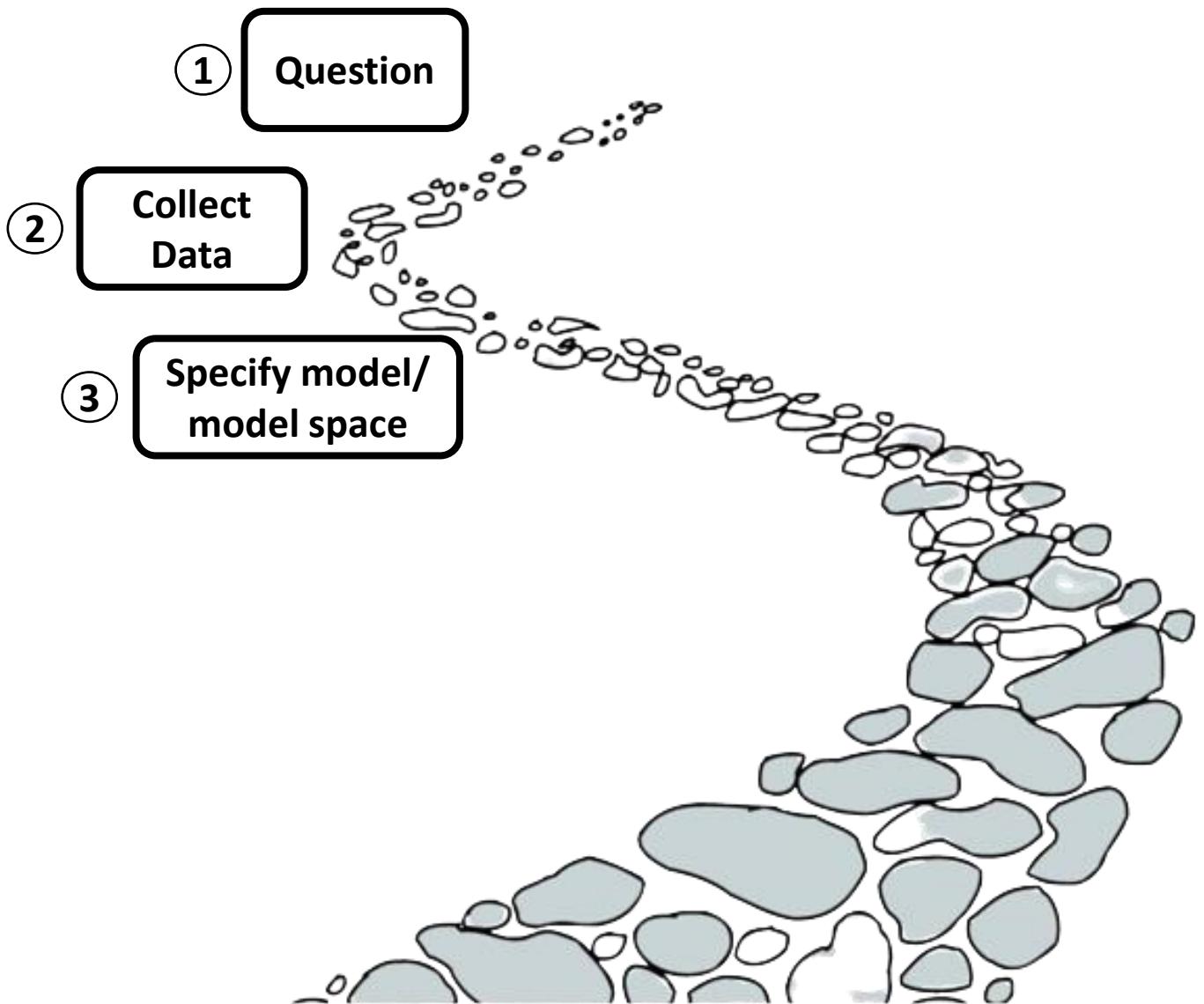
- MFM and NFM can express more complex dynamics (Marreiros *et al.*, 2009)
- Go there if this is needed, otherwise stick to the simpler model

# Specify model/model space

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## Steps

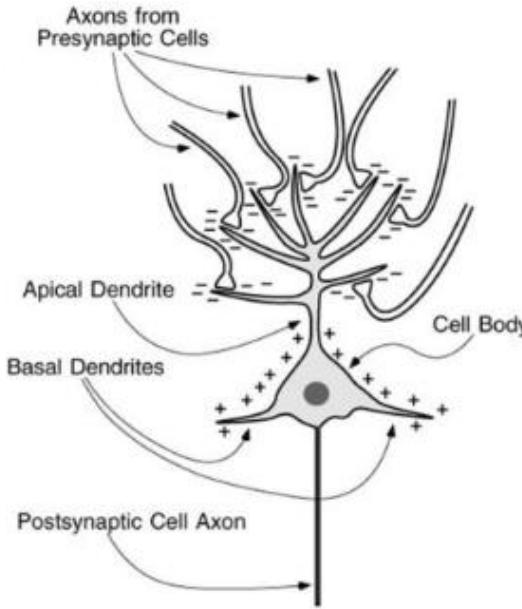
- Translate your question into a model comparison or a parameter inference problem
- Select regions
- Select a variant of DCM
- **Example: The “ERP” model**
- Specify connectivity architecture



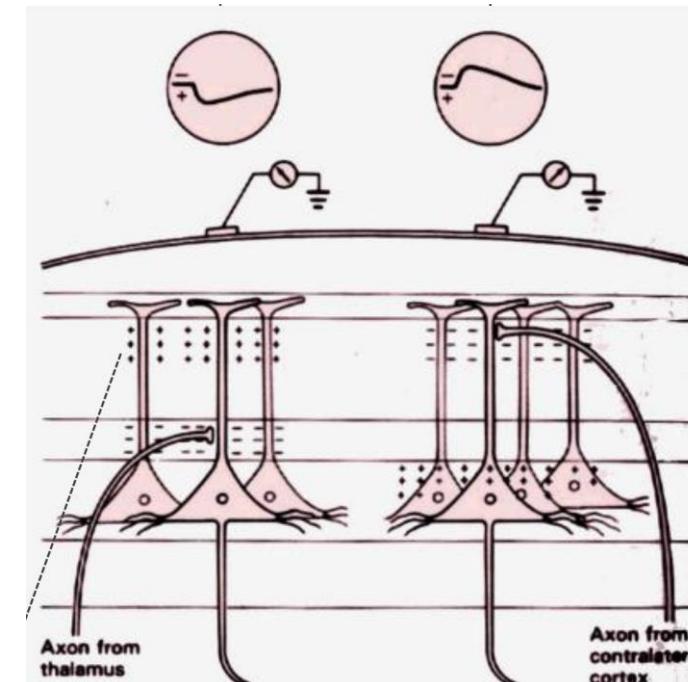
# What do we measure with EEG?

## Background

- Underlying mechanisms generating EEG



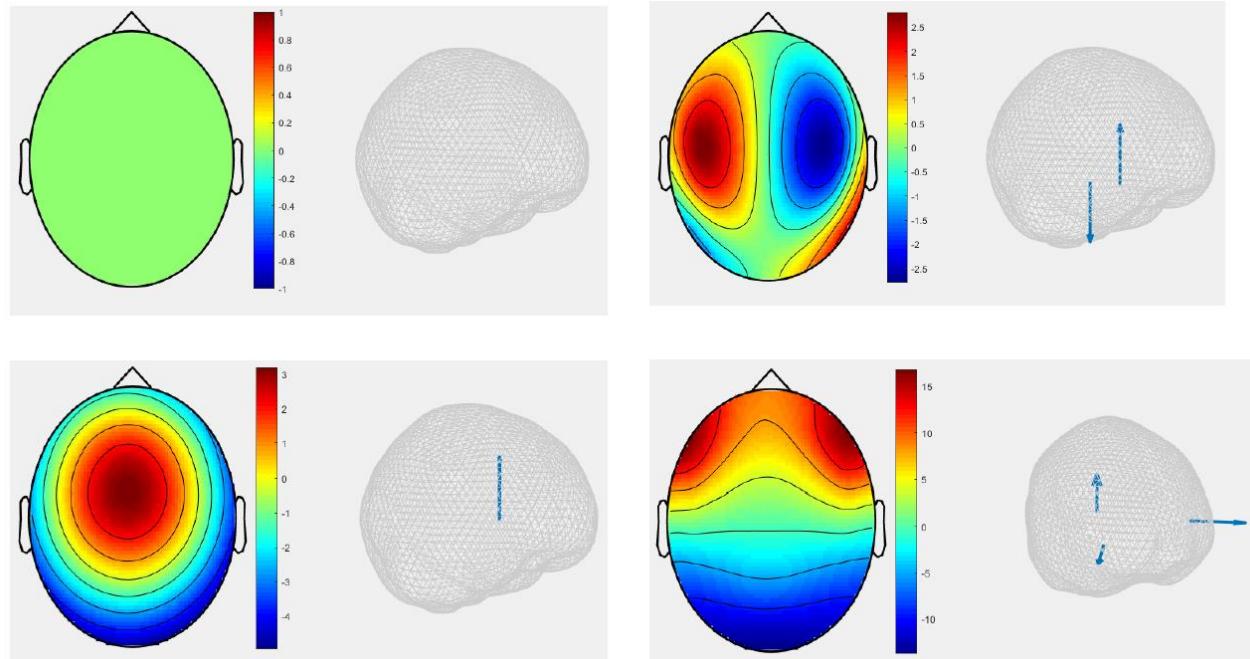
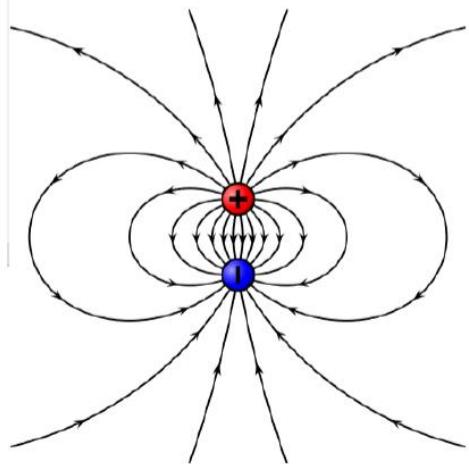
Measurable dipoles



# What do we measure with EEG?

## Background

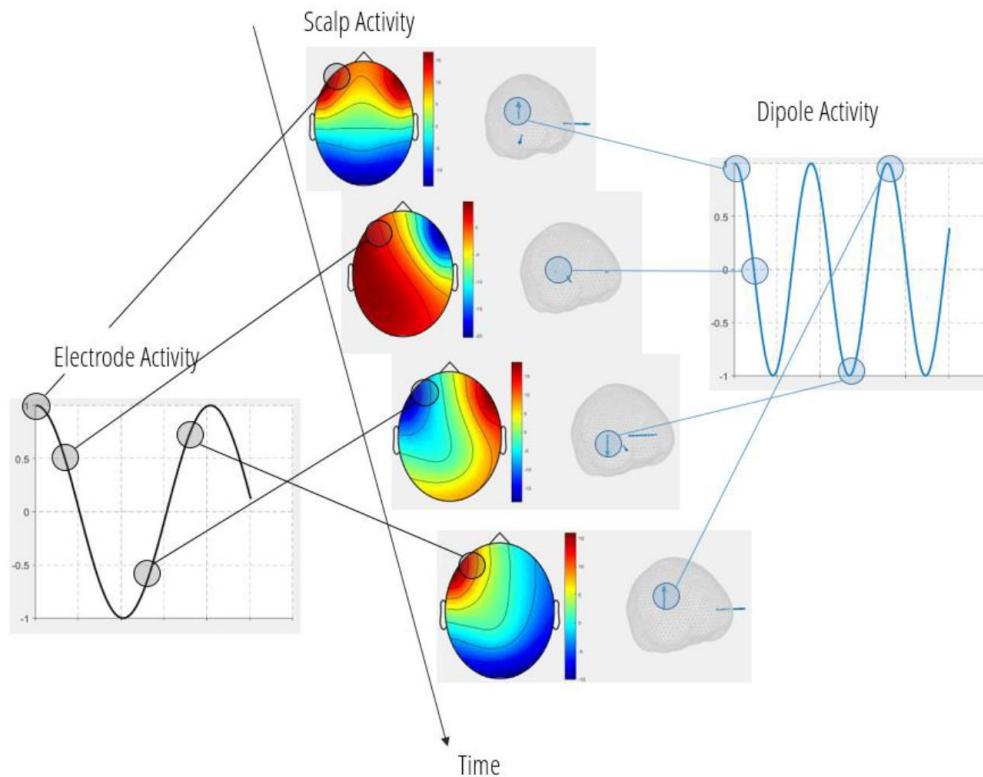
- Dipole generates measurable electrical fields in EEG sensors



# What do we measure with EEG?

## Background

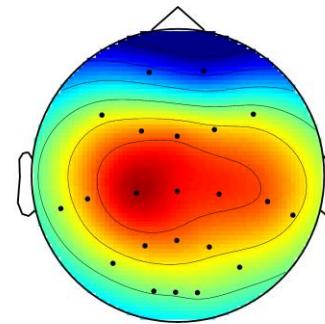
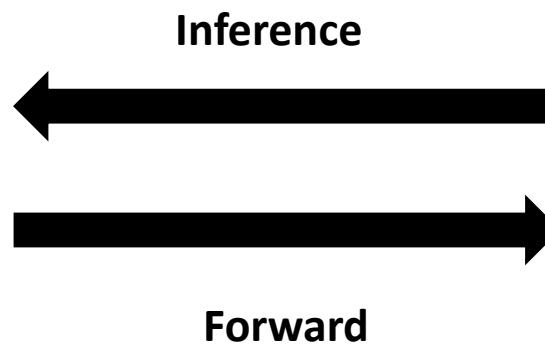
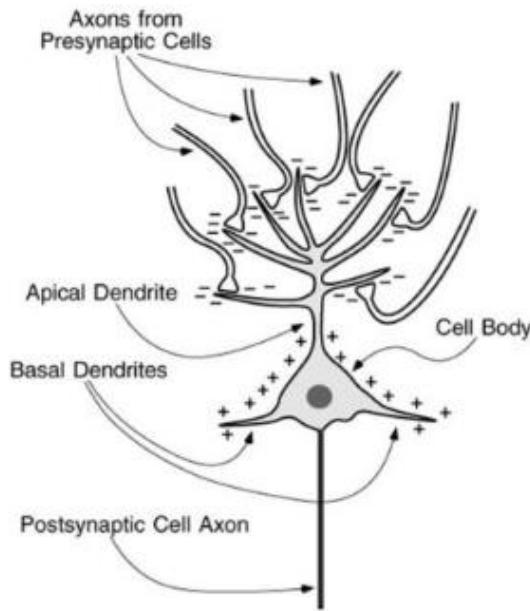
- Dipole moments change over time and lead to changes in measured scalp potentials



# What do we measure with EEG?

## Question

- Can we make inferences about properties of the neuronal sources that generate these signals?

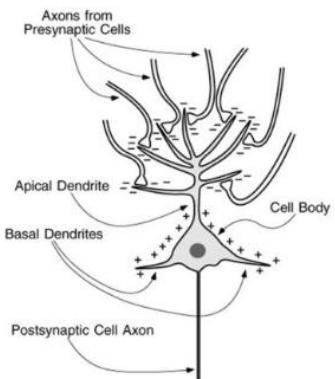


# Scales of analysis



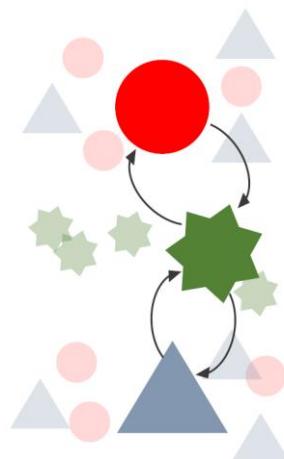
## Single cell

1-10  $\mu\text{m}^2$



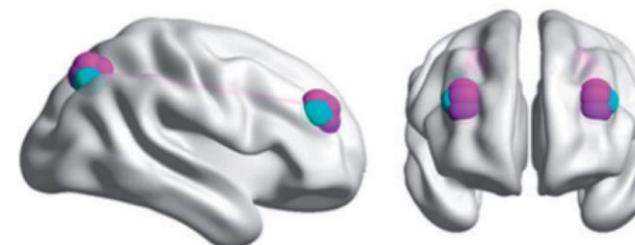
## Cortical Column

1-10  $\text{mm}^2$  to 1-5  $\text{cm}^2$



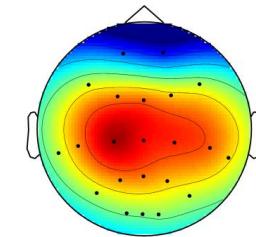
## Brain Networks

5-20  $\text{cm}^2$



## Scalp potentials

30-38  $\text{cm}^2$



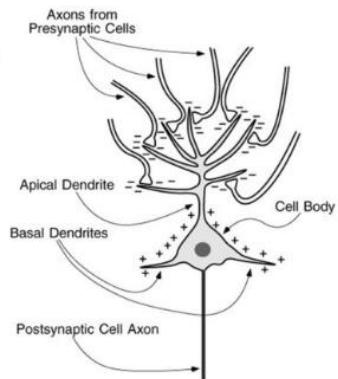
# Scales of analysis

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## Single cell

1-10  $\mu\text{m}^2$



# Source modelling at the microscale



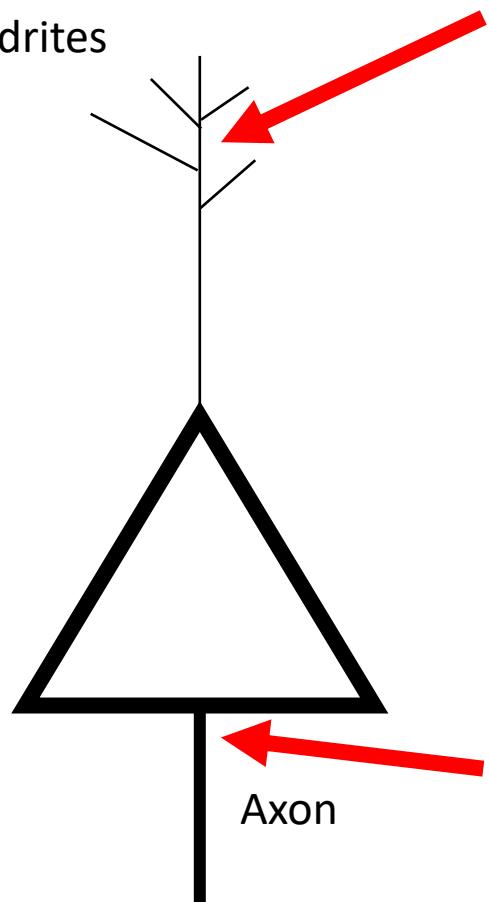
Microscale

Mesoscale

Macroscale

Increasing spatial scale

Dendrites



**Input:** Spikes of a presynaptic cell are converted into postsynaptic potentials (PSPs)

Axon

**Output:** Postsynaptic potentials are converted into a spike

# Source modelling at the microscale

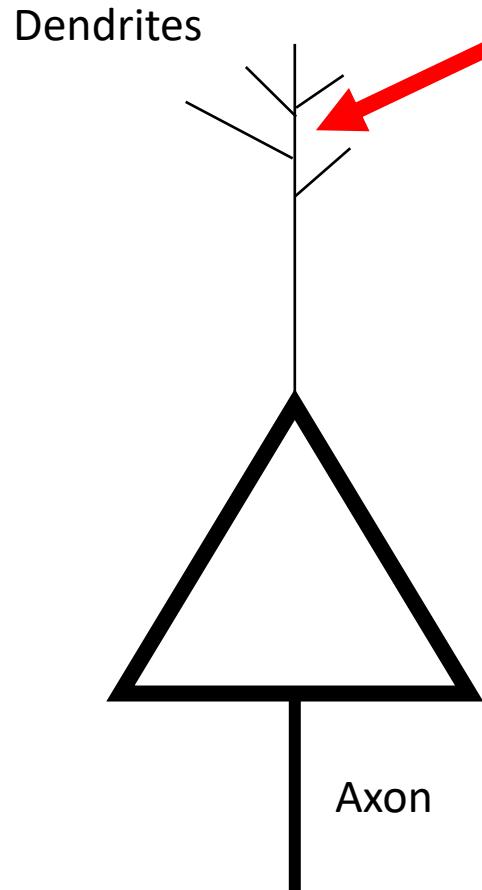


Microscale

Mesoscale

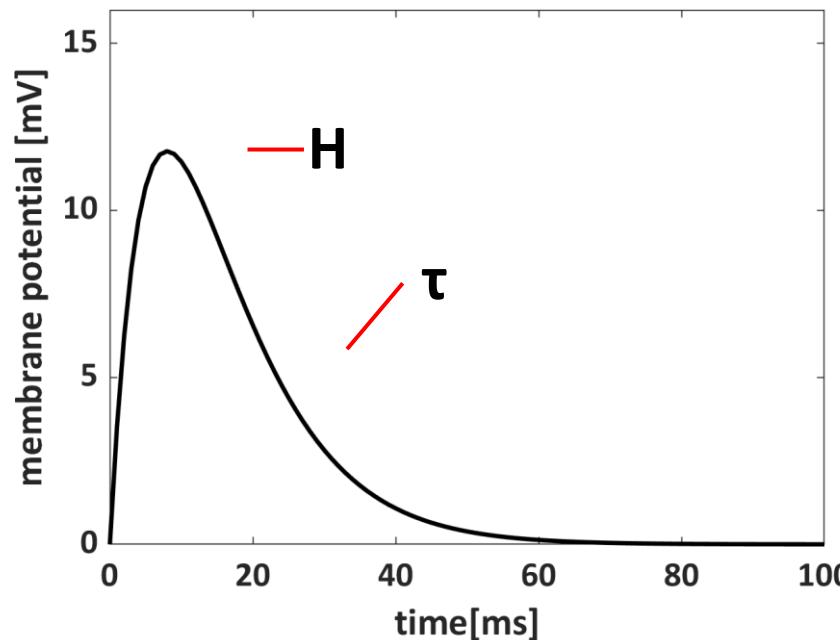
Macroscale

Increasing spatial scale



**Input:** Spikes of a presynaptic cell are converted into postsynaptic potentials (PSPs)

David et al (2005), NeuroImage



$$h(t) = \begin{cases} \frac{Ht \exp(-t/\tau)}{\tau} & t \geq 0 \\ 0 & t < 0 \end{cases}$$

**Convolution** of incoming neural activity with synaptic kernel

# Source modelling at the microscale

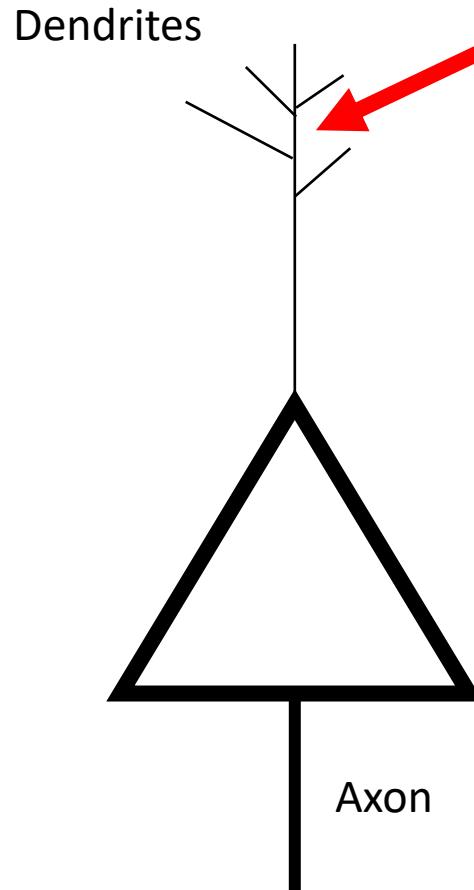


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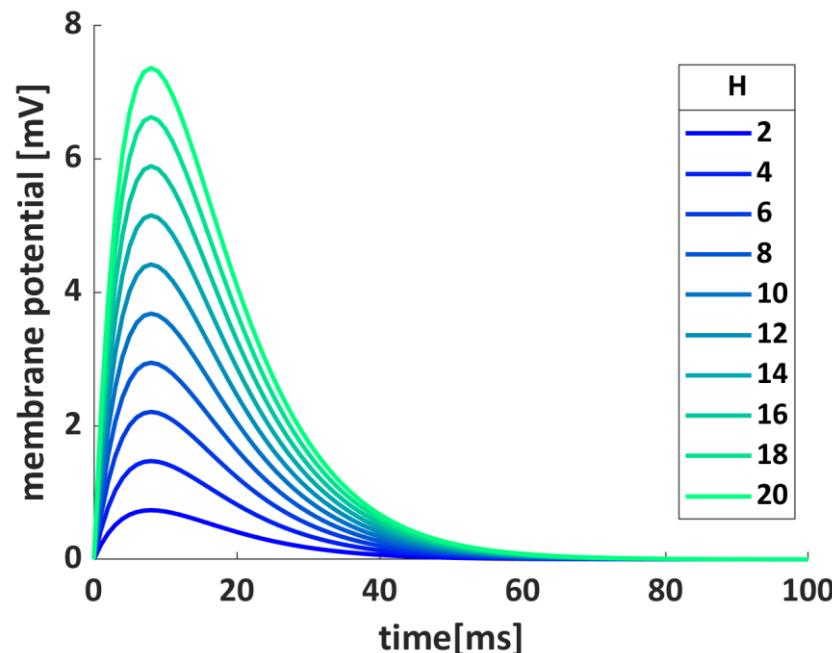
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**Input:** Spikes of a presynaptic cell are converted into postsynaptic potentials (PSPs)



**H:** Maximum postsynaptic potential

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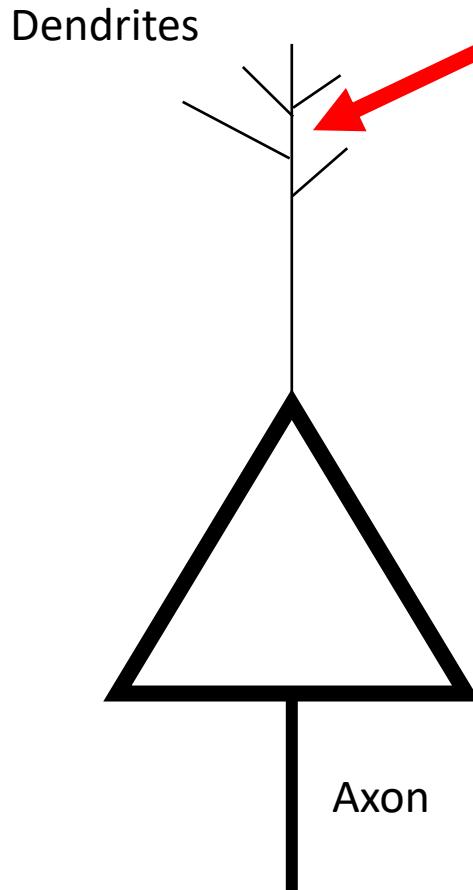


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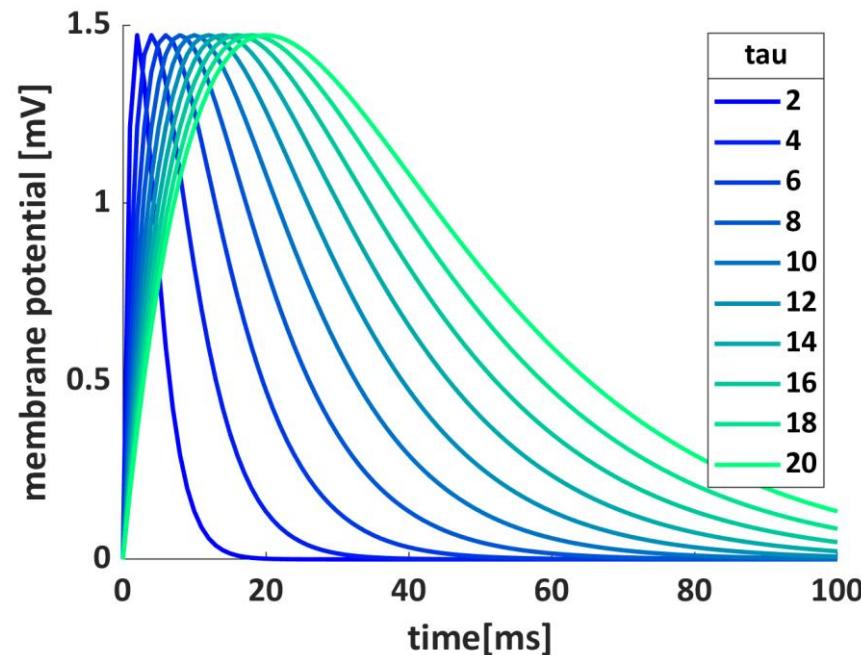
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$\tau$ : Inverse time constant

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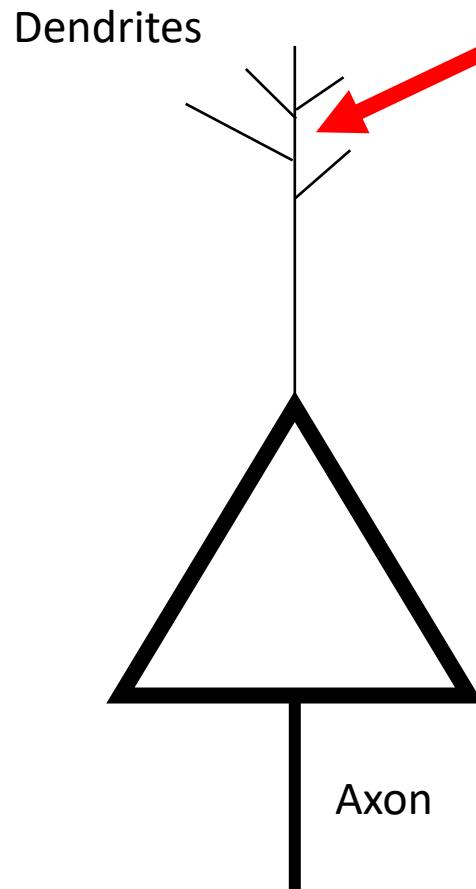


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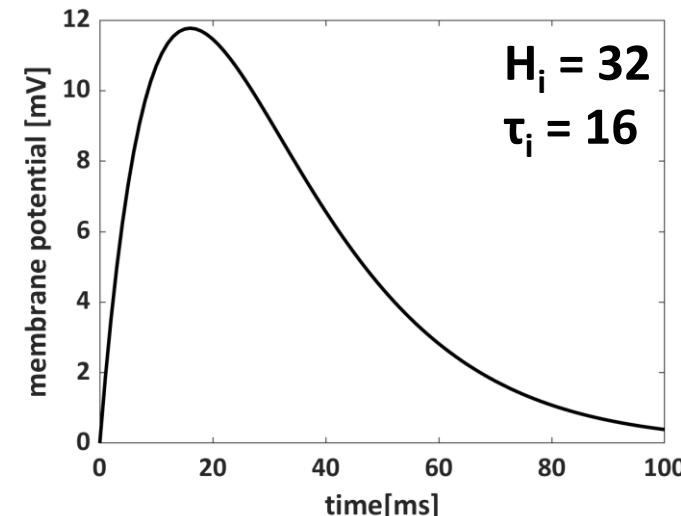
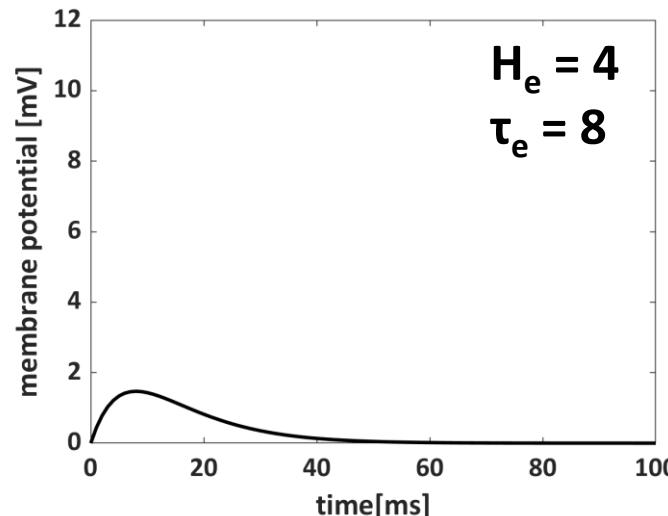
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**Input:** Spikes of a presynaptic cell are converted into postsynaptic potentials (PSPs)



spm\_fx\_erp.m

$H_i > H_e$ , because inhibitory synapse is closer to the soma

Jansen & Ritt (1995), Biol. Cyb.

# Source modelling at the microscale



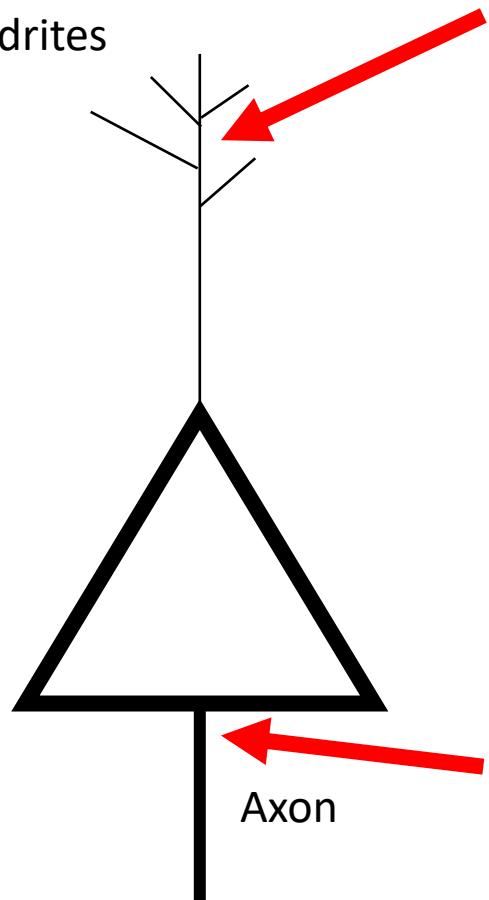
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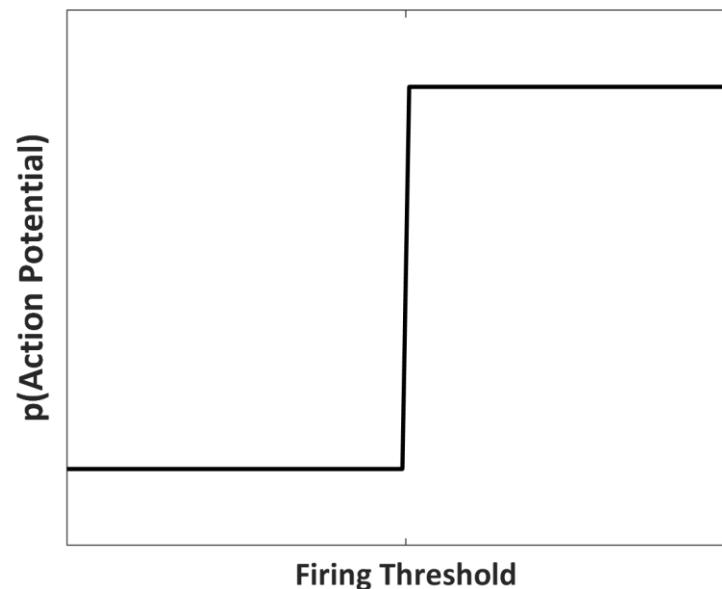
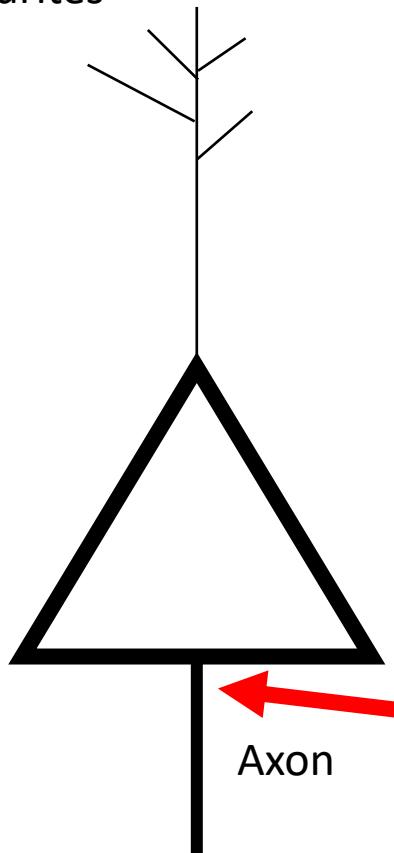
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**Output:** Postsynaptic potentials are converted into a spike

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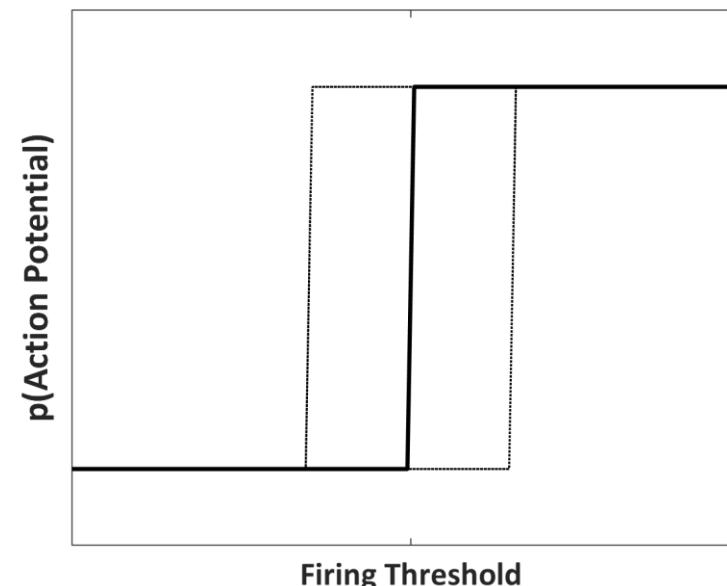
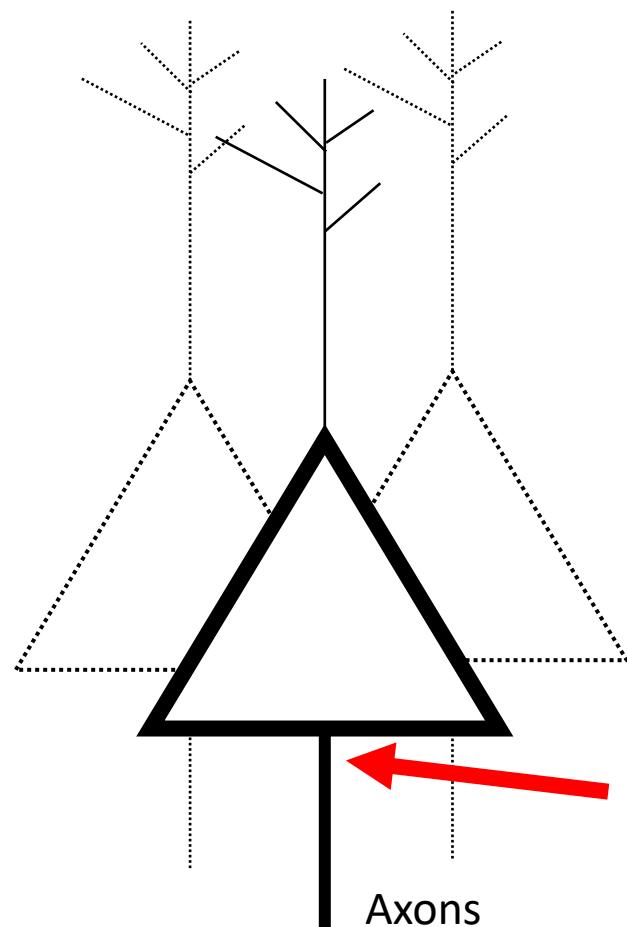
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**Output:** Postsynaptic potentials are converted into a spike

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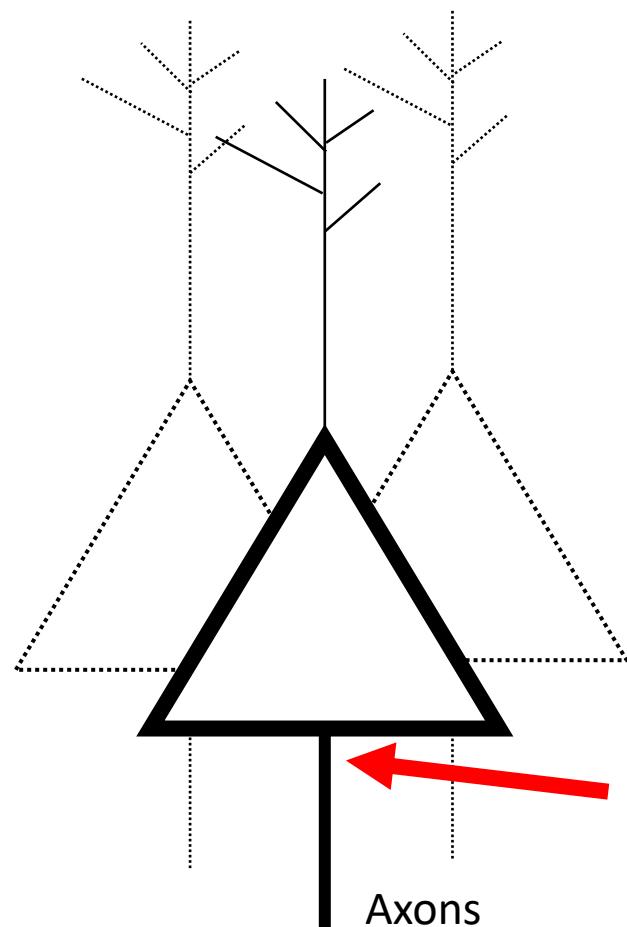
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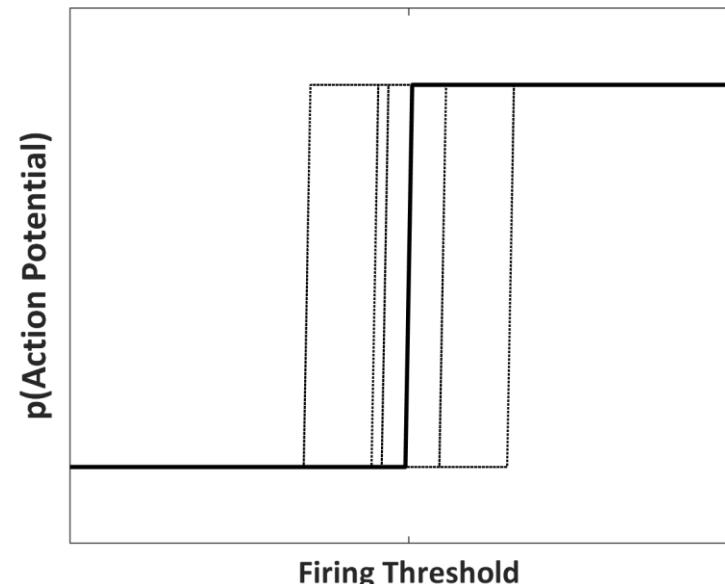
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**Output:** Postsynaptic potentials are converted into a spike



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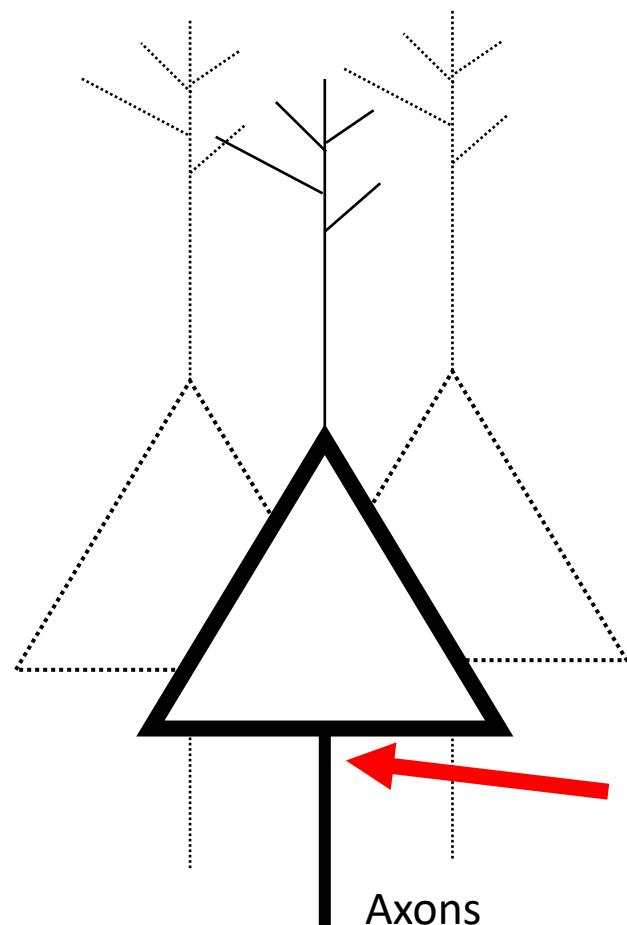
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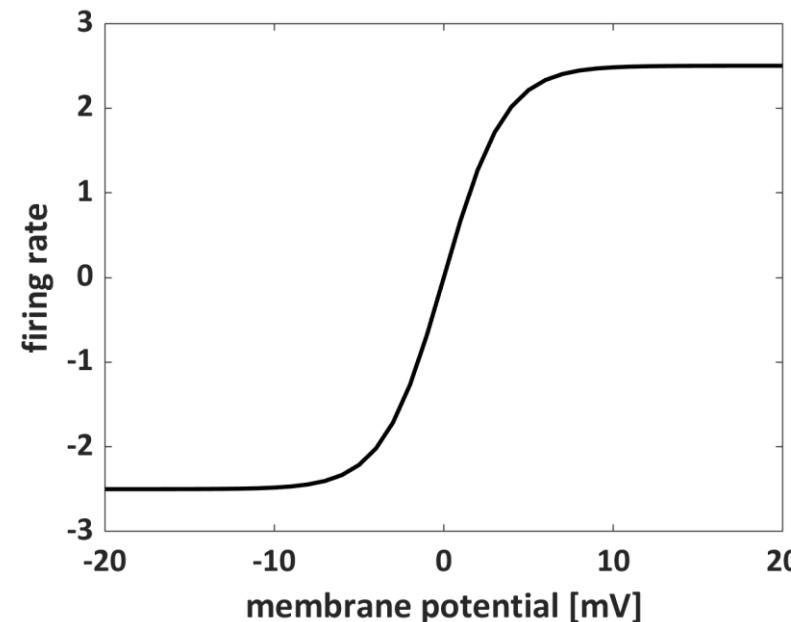
Macroscale

Increasing spatial scale

Dendrites



**Output:** Postsynaptic potentials are converted  
into an average spike rate



Mean-field  
approximation

# Source modelling at the microscale



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**Mean-field  
approximation**

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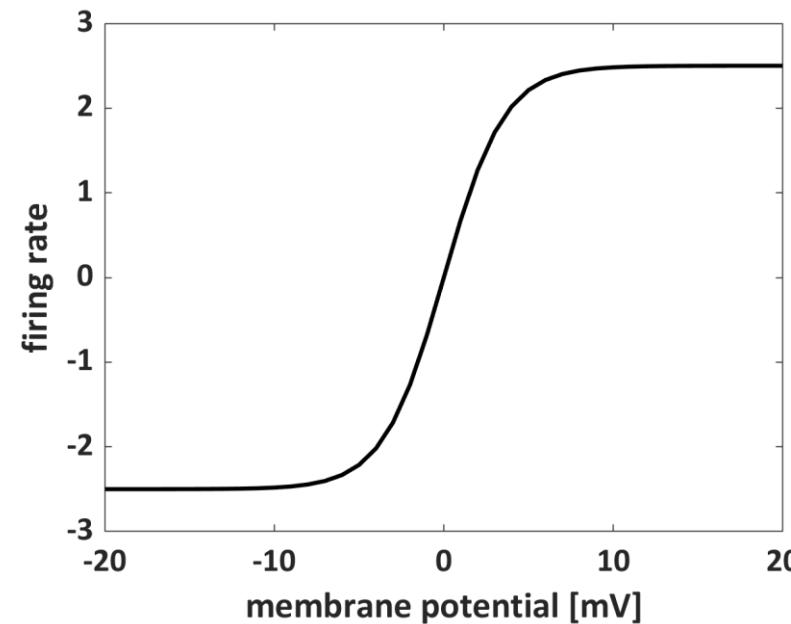
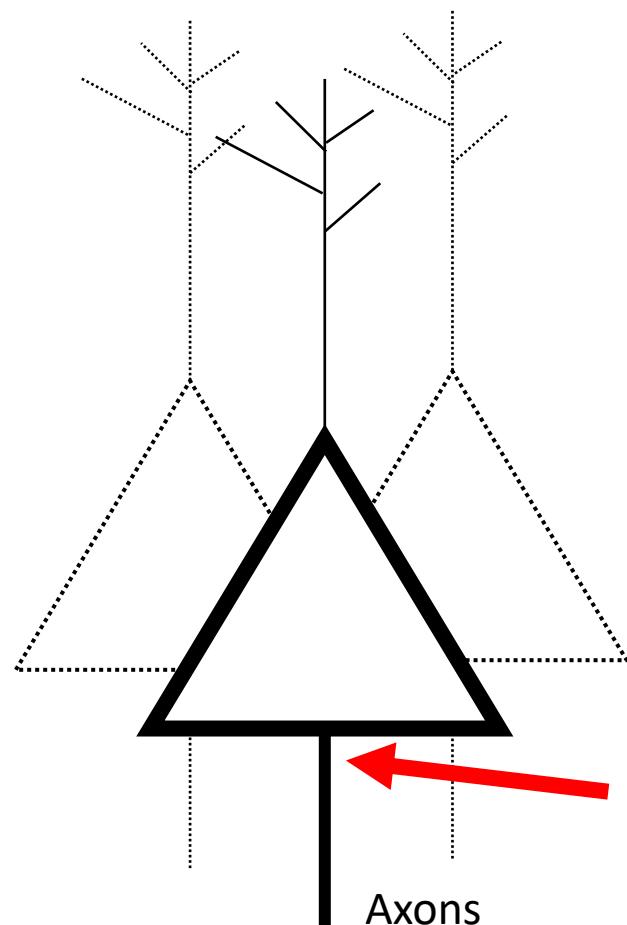
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**Output:** Postsynaptic potentials are converted  
into an average spike rate

$$S(v) = \frac{2e_0}{1 + \exp(-rv)} - e_0$$

$$e_0 = 2.5 \\ r = 0.56$$

David et al (2005), NeuroImage

# Source modelling at the microscale



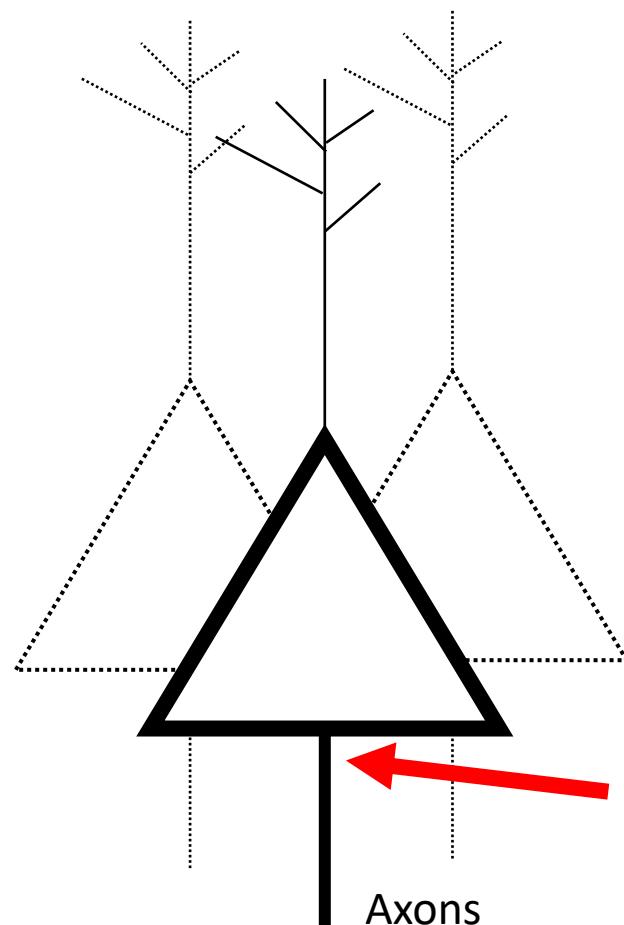
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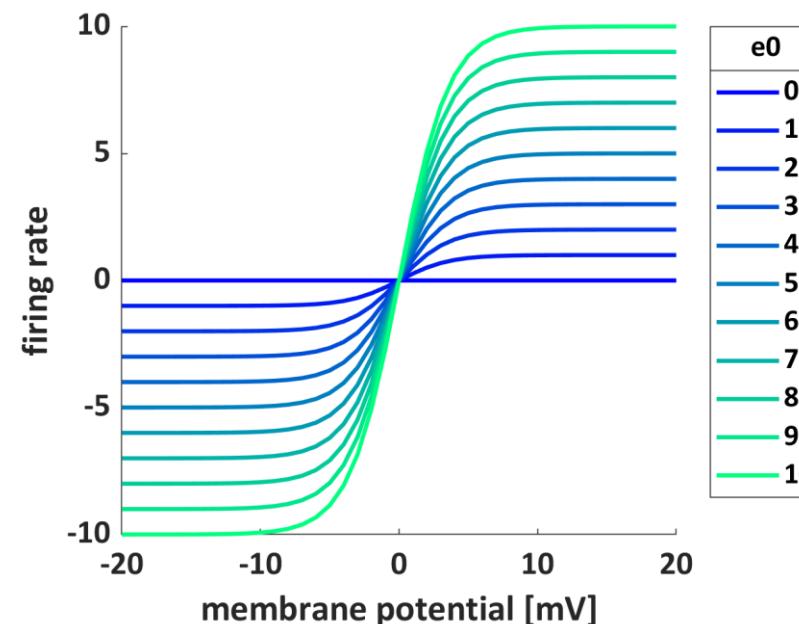
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Dendrites



**Output:** Postsynaptic potentials are converted  
into an average spike rate



$$S(v) = \frac{2e_0}{1 + \exp(-rv)} - e_0$$

**e<sub>0</sub>:** Maximal firing rate

David et al (2005), NeuroImage

# Source modelling at the microscale



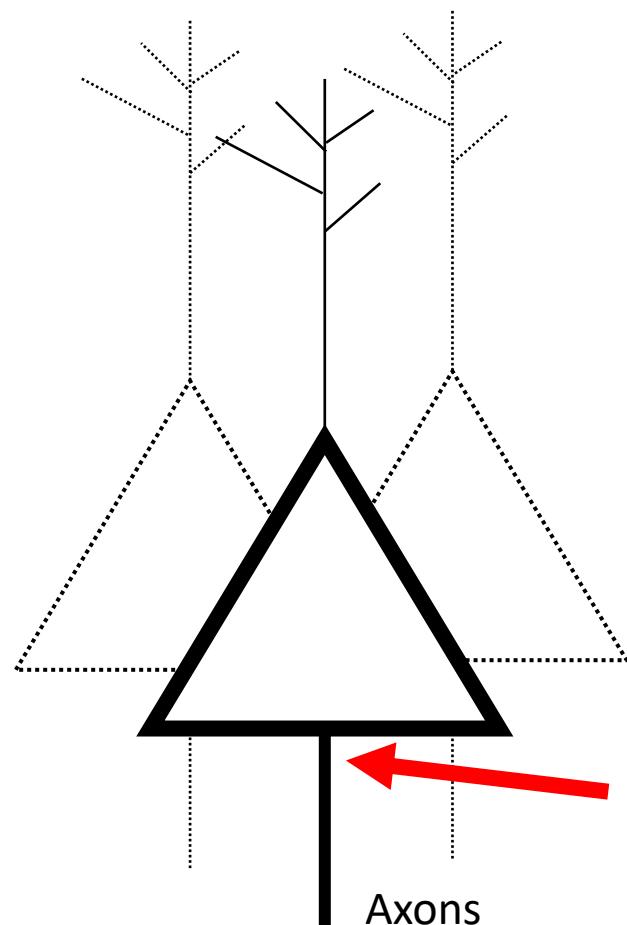
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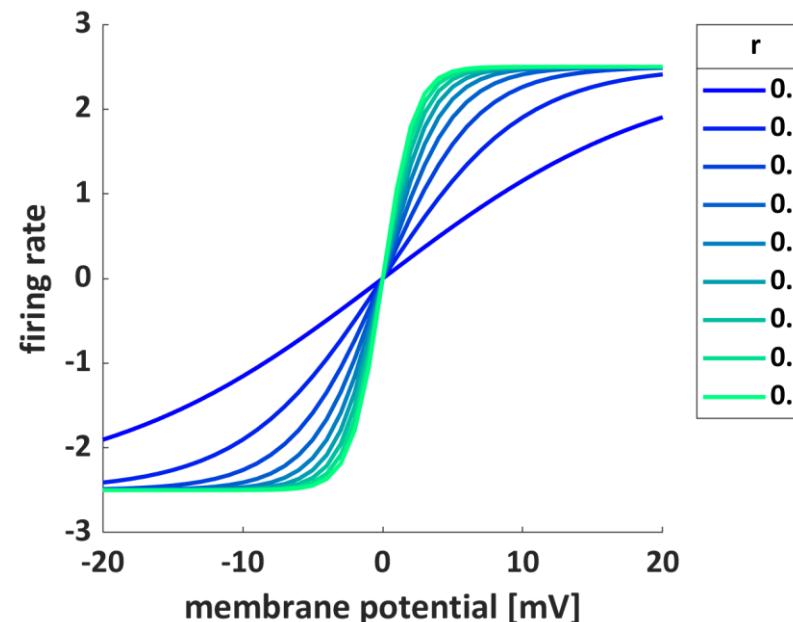
Macroscale

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Dendrites



**Output:** Postsynaptic potentials are converted into an average spike rate



$$S(v) = \frac{2e_0}{1 + \exp(-rv)} - e_0$$

$r$ : Stochasticity

David et al (2005), NeuroImage

# Source modelling at the microscale



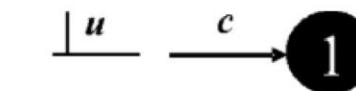
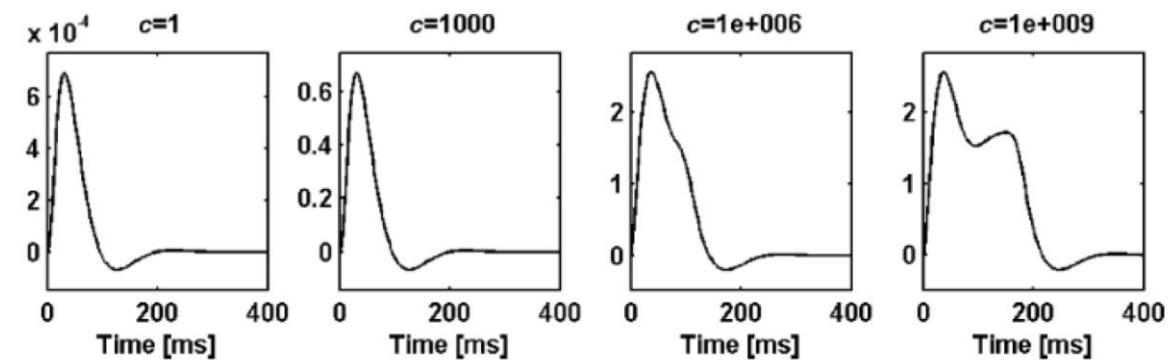
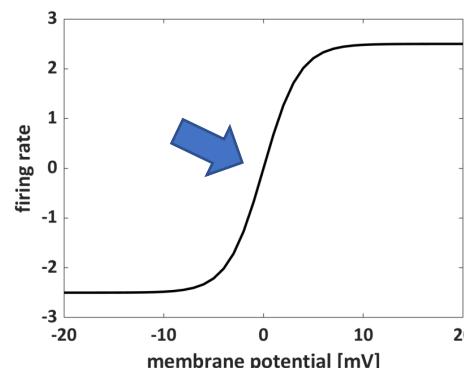
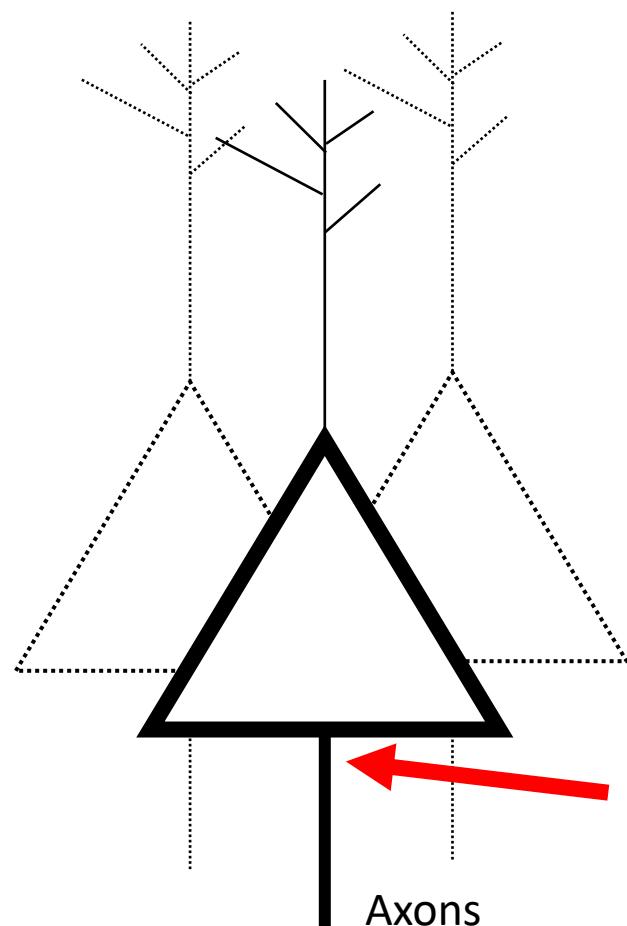
Microscale

Mesoscale

Macroscale

Increasing spatial scale

Dendrites



**Output:** Postsynaptic potentials are converted into an average spike rate

David et al (2005), NeuroImage

# Source modelling at the microscale



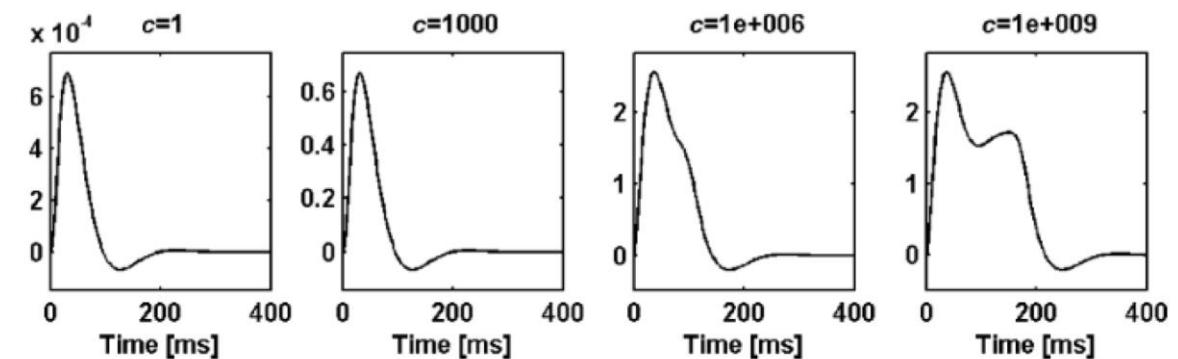
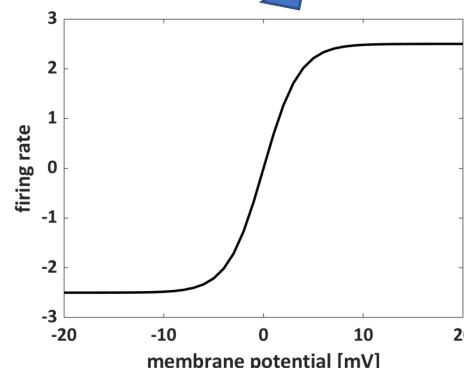
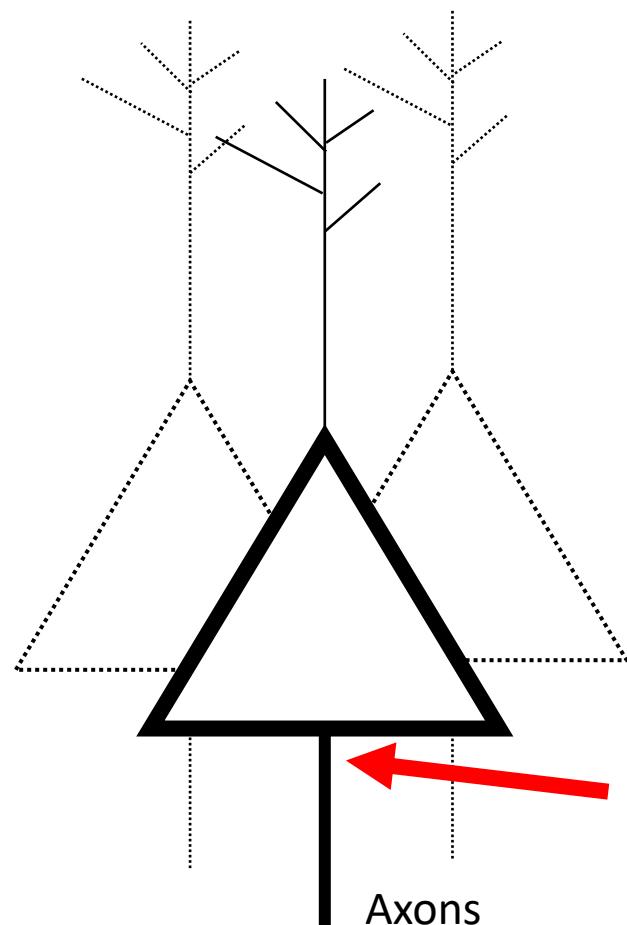
Microscale

Mesoscale

Macroscale

Increasing spatial scale

Dendrites



**Output:** Postsynaptic potentials are converted into an average spike rate

David et al (2005), NeuroImage

# Source modelling at the microscale



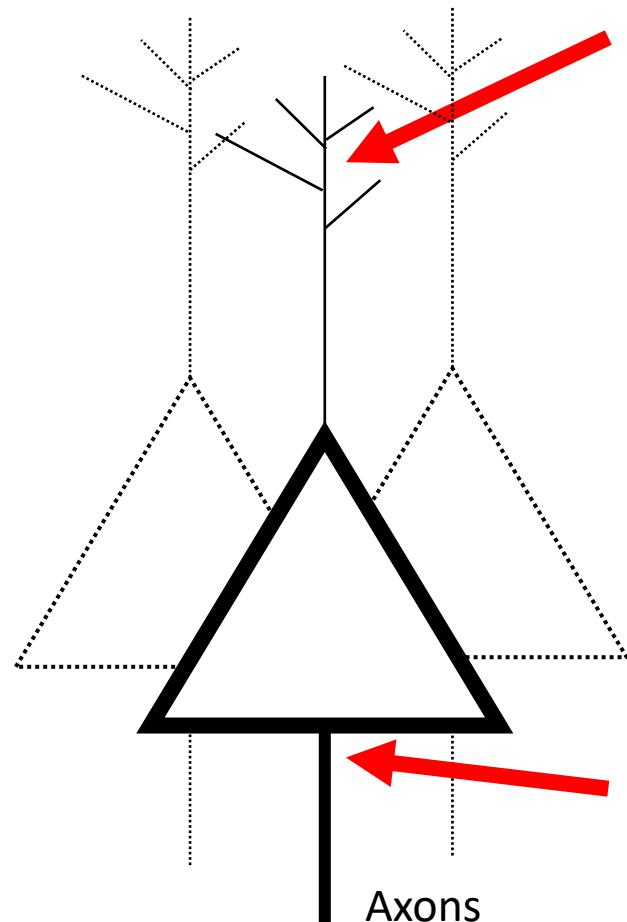
Microscale

Mesoscale

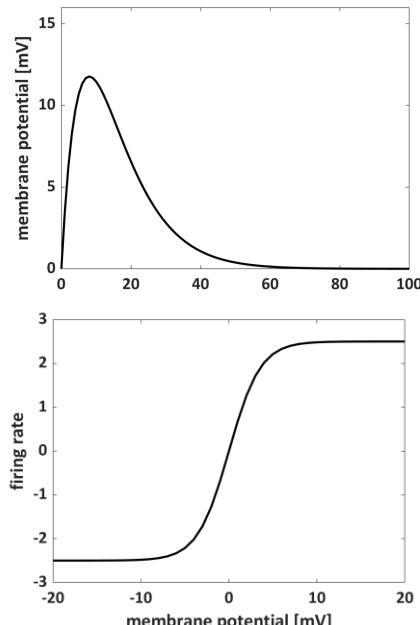
Macroscale

Increasing spatial scale

Dendrites



**Input:** Spike rates of a presynaptic cells are converted into postsynaptic potentials (PSPs)



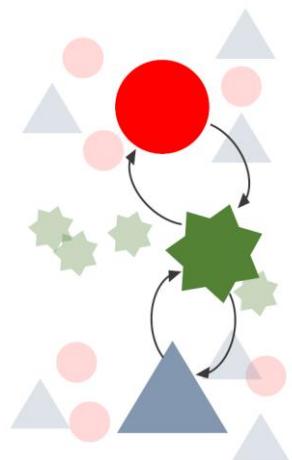
**Output:** Postsynaptic potentials are converted into an average spike rate

# Scales of analysis



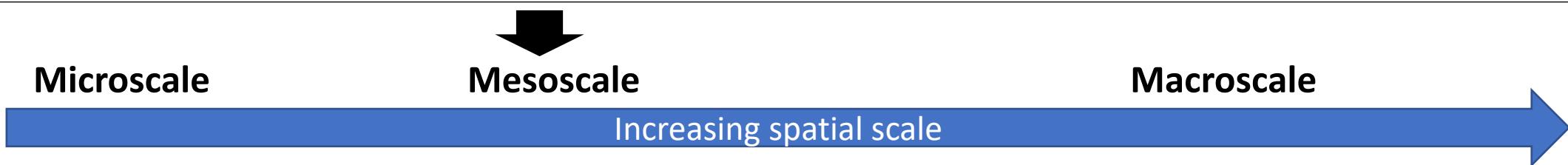
## **Cortical Column**

1-10 mm<sup>2</sup> to 1-5 cm<sup>2</sup>

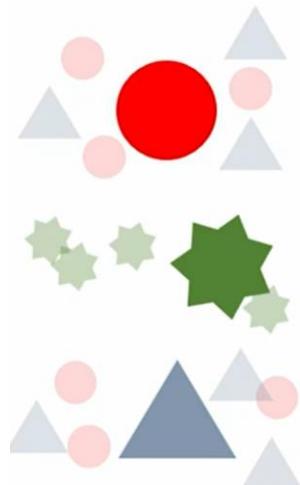


Schöbi (2019), CPC Lecture

# Source modelling at the mesoscale

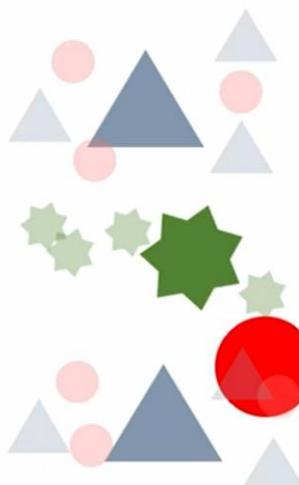


ERP



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

CMC

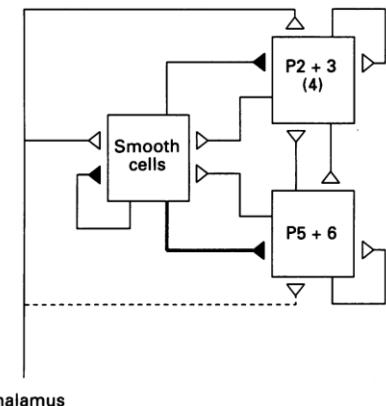


Superficial and deep pyramidal cell populations are modeled individually

# Source modelling at the mesoscale

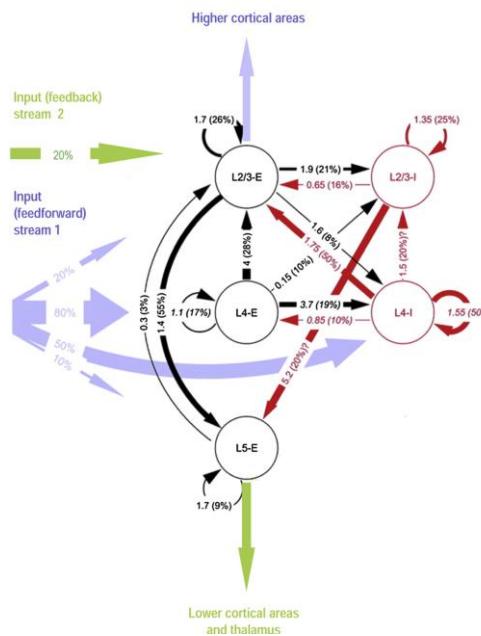


# Original microcircuit



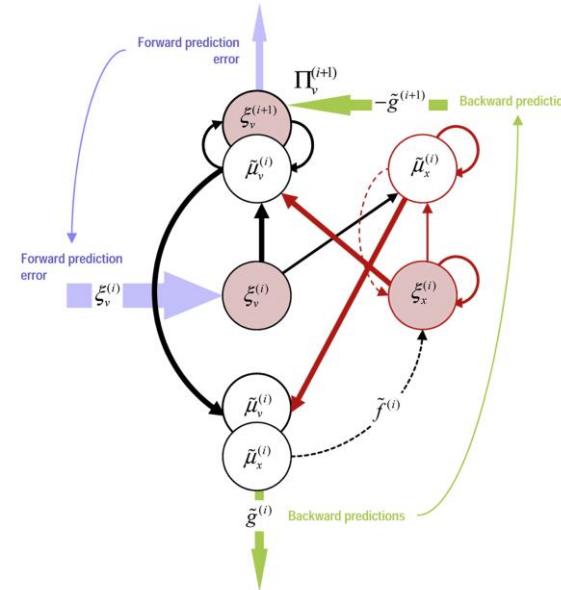
Douglas & Martin (1991),  
J. Physiol.

# Updated microcircuit



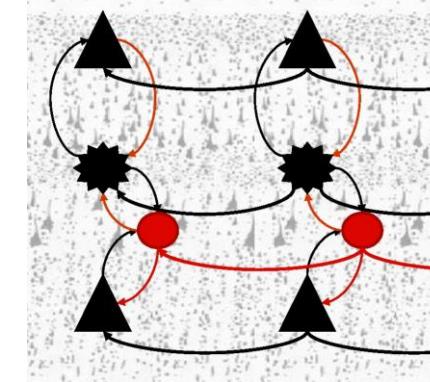
Haeusler & Maass (2006),  
Cerebral Cortex

## Canonical microcircuit



Bastos et al. (2012)  
Neuron

## Reduced microcircuit (DCM)



Pinotsis et al. (2012),  
NeuronImage

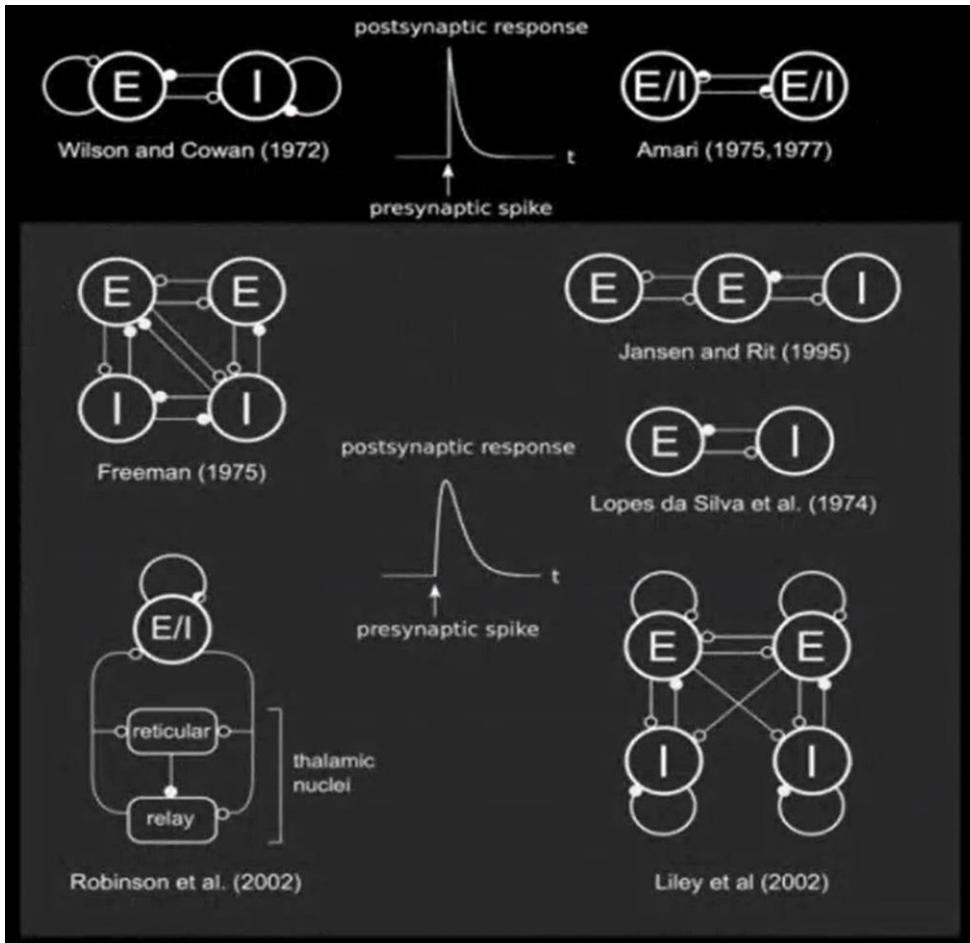
# Source modelling at the mesoscale

# Microscale

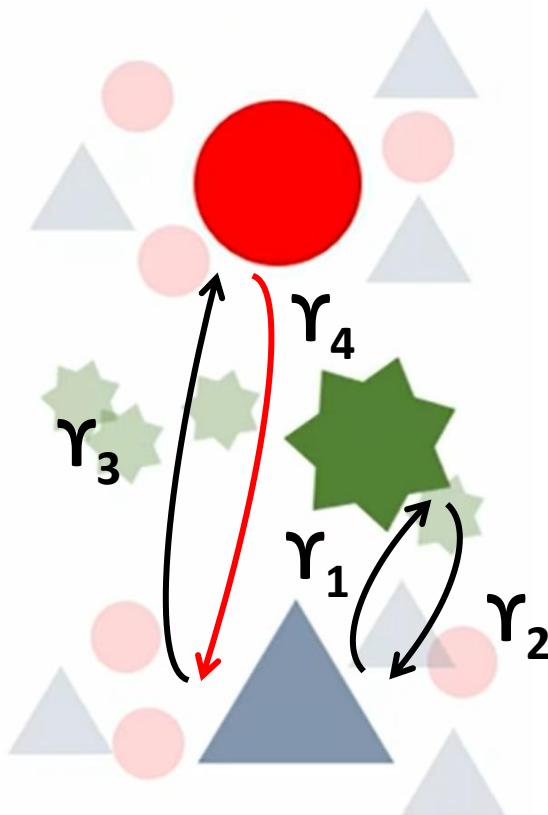
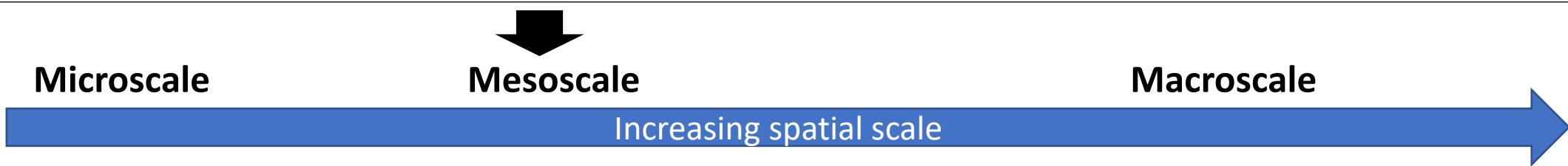
Mesoscale

# Macroscale

## Increasing spatial scale



# Source modelling at the mesoscale



Schöbi (2019), CPC Lecture

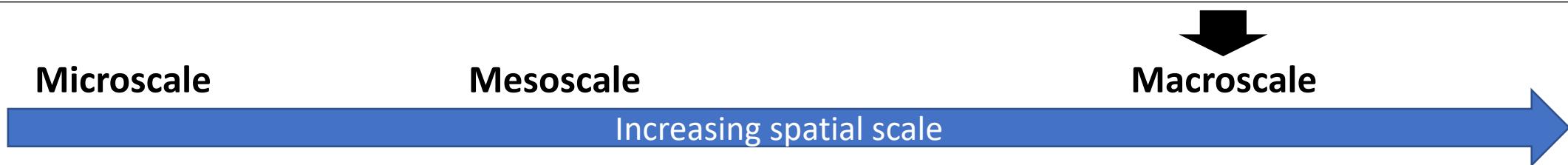
**But what value should  $\gamma$  have?**

$$\gamma_2 = 0.8\gamma_1, \gamma_3 = \gamma_4 = 0.25\gamma_1$$

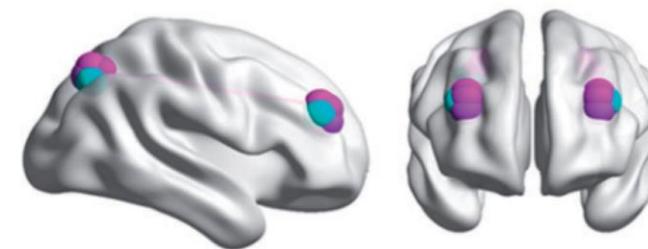
- Based on counts of synapsis
- Animal studies (mouse, cat)
- Visual cortex

Jansen & Ritt (1995), Biol. Cyb.

# Source modelling at the macroscale

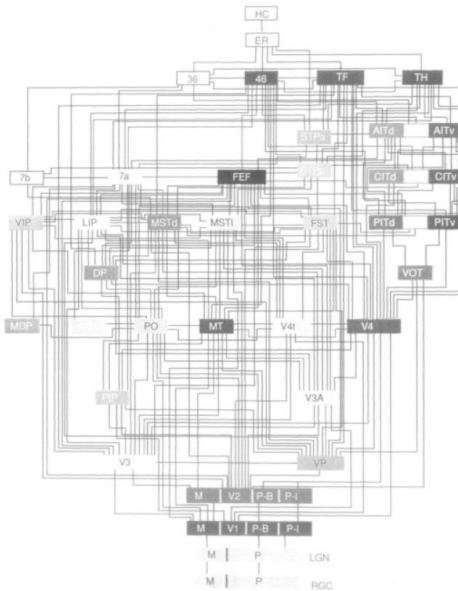
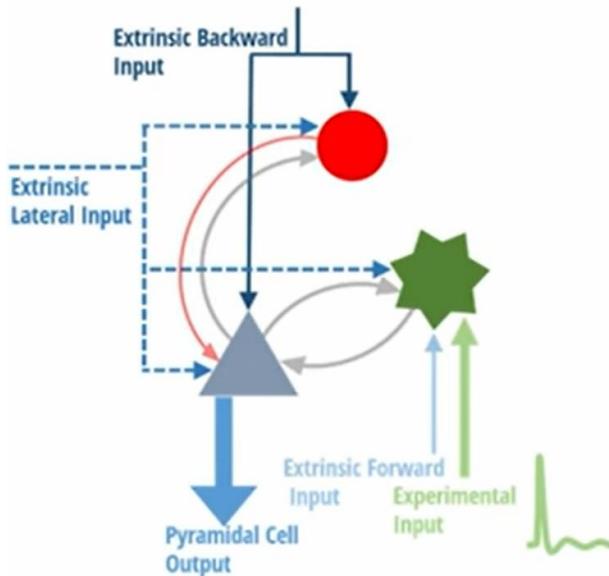
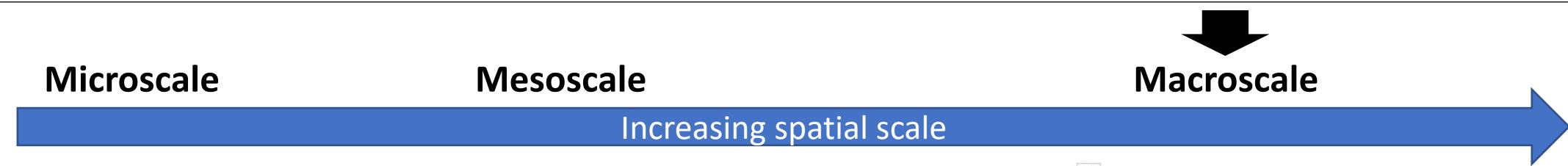


**Brain Networks**  
5-20 cm<sup>2</sup>



Symmonds et al. (2018), Brain

# Source modelling at the macroscale

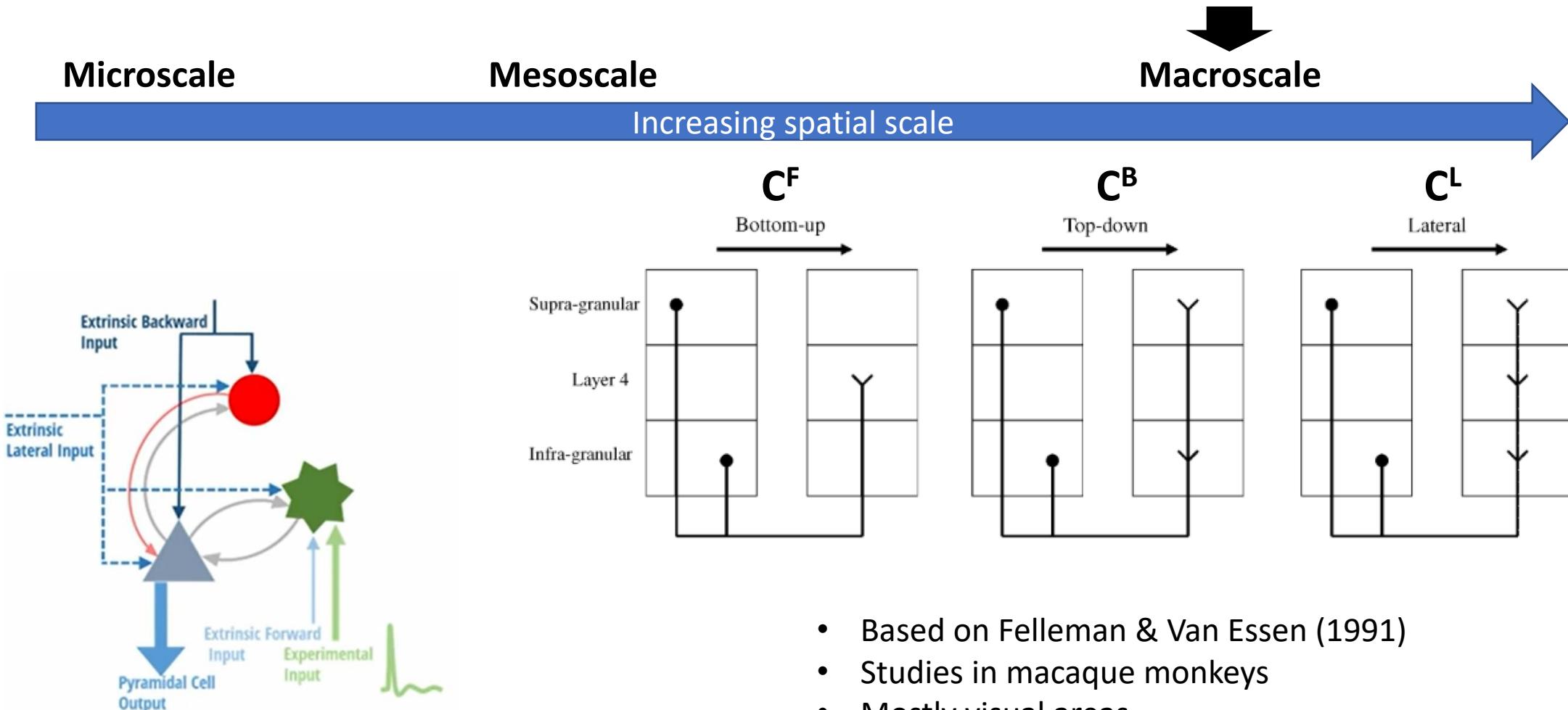


- Based on Felleman & Van Essen (1991)
- Studies in macaque monkeys
- Mostly visual areas
- But see Markov et al (2013)

Schöbi (2019), CPC Lecture

David et al (2005), NeuroImage

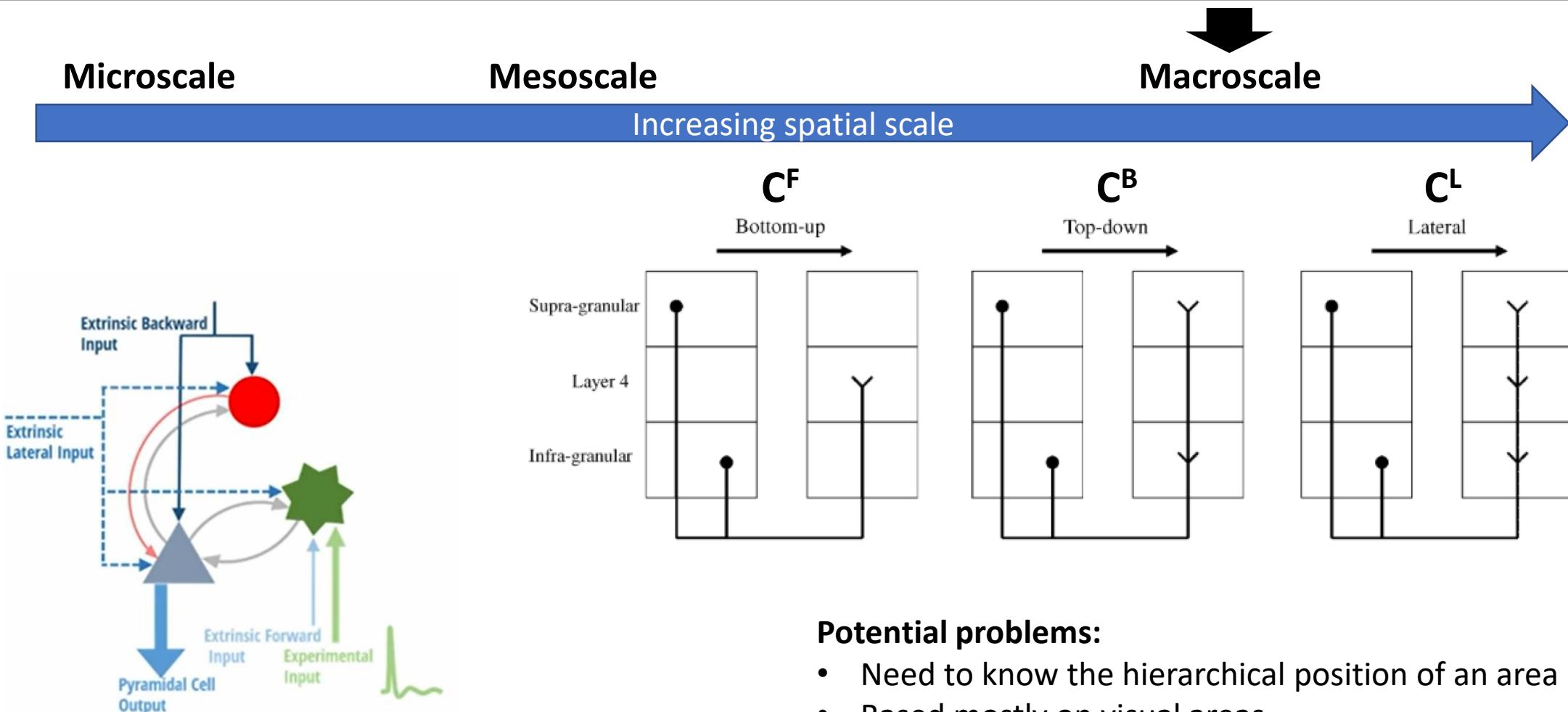
# Source modelling at the macroscale



Schöbi (2019), CPC Lecture

David et al (2005), NeuroImage

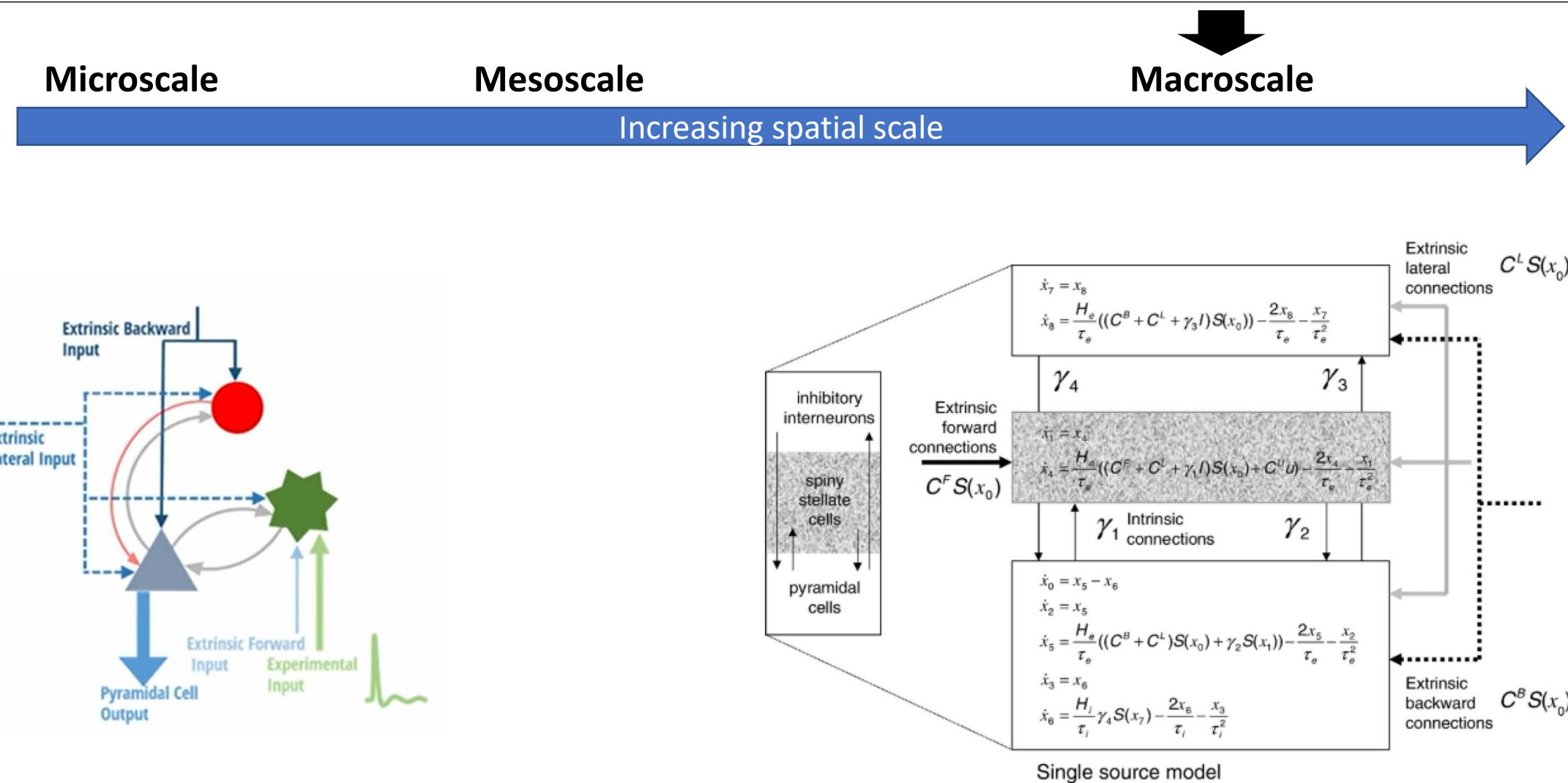
# Source modelling at the macroscale



## Potential problems:

- Need to know the hierarchical position of an area
- Based mostly on visual areas
- Based on areas with a 6-layer structure

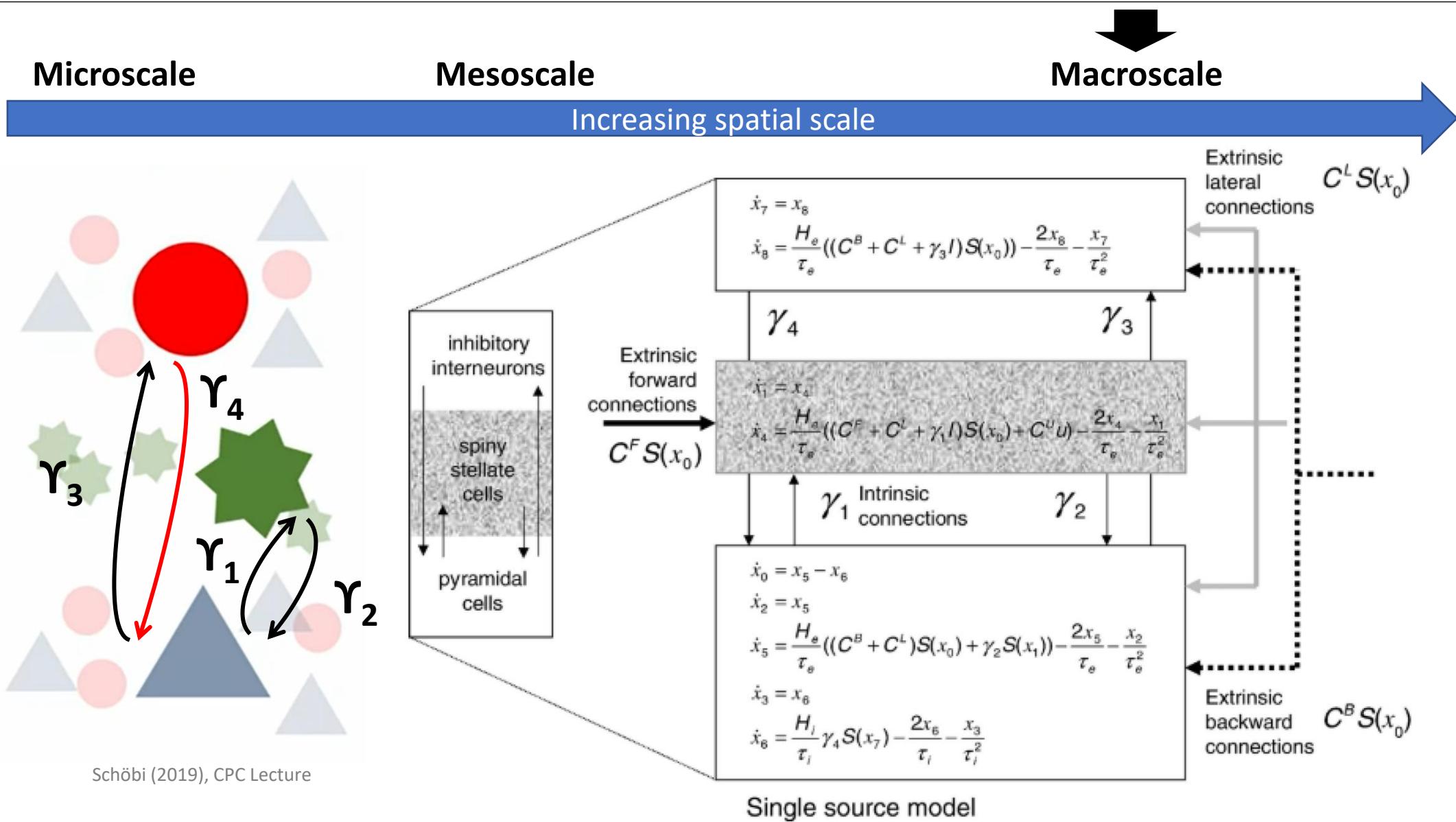
# Source modelling at the macroscale



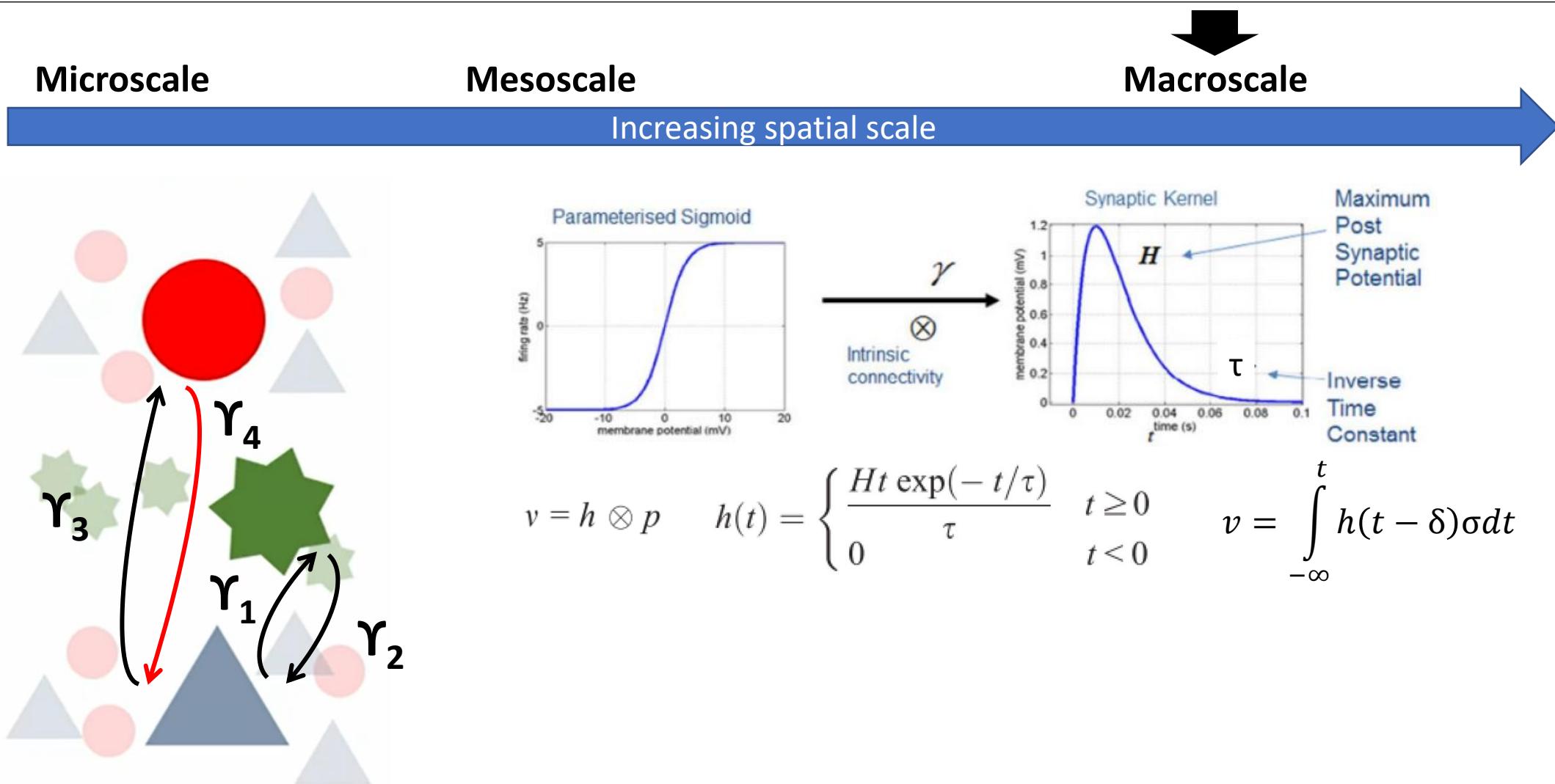
Schöbi (2019), CPC Lecture

David et al (2006), Neuroimage

# Source modelling at the macroscale



# Source modelling at the macroscale

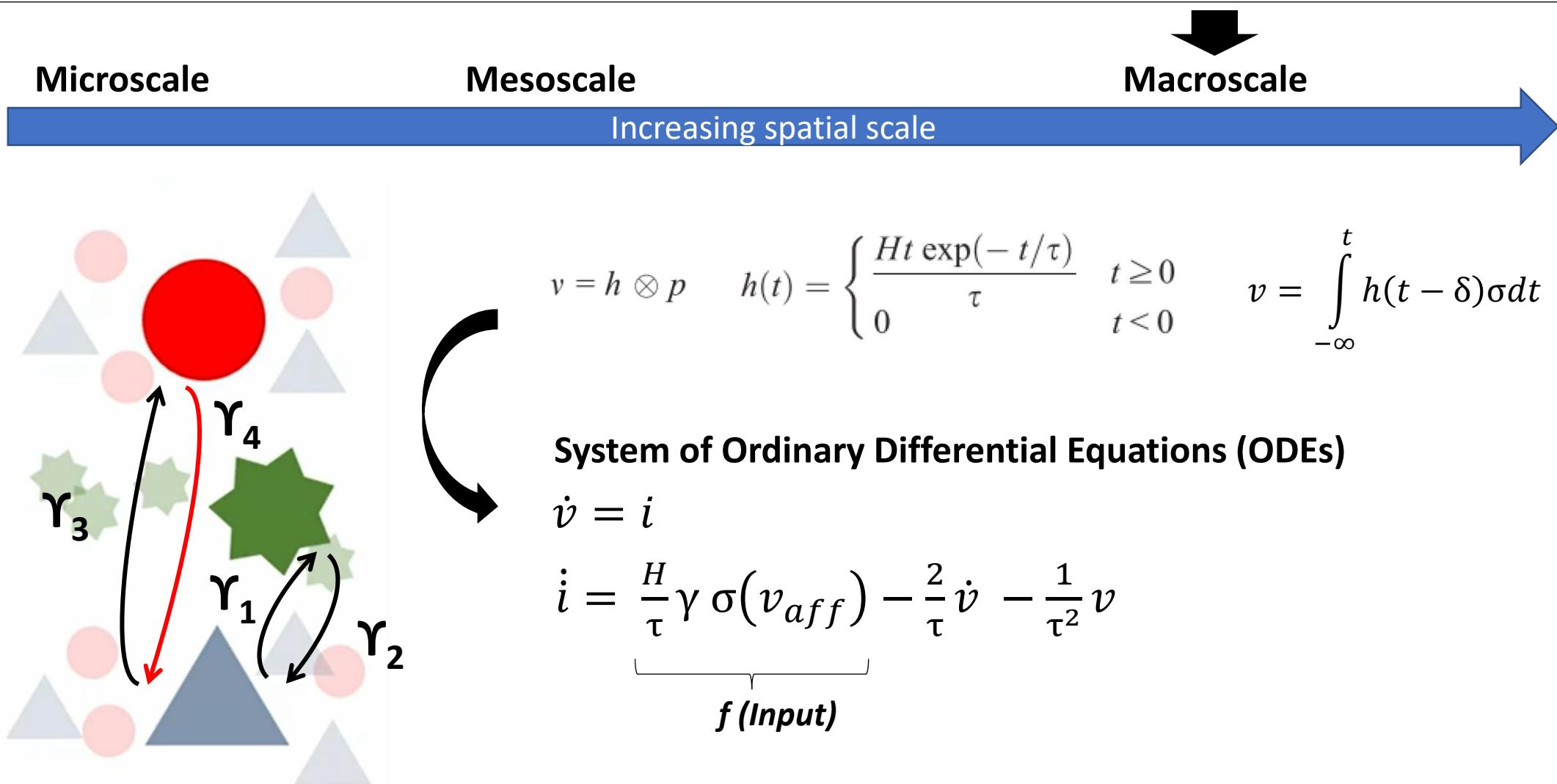


Schöbi (2019), CPC Lecture

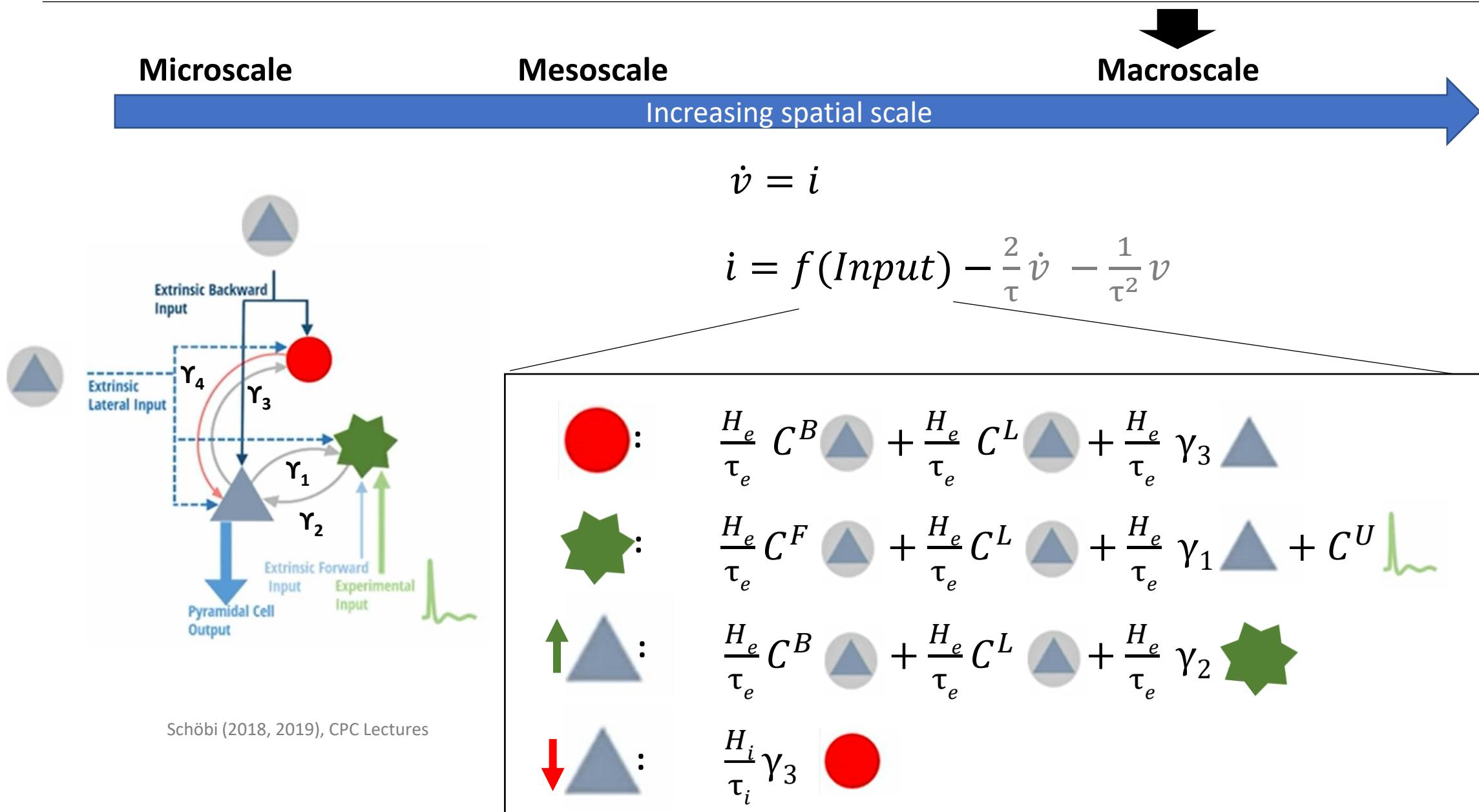
Moran et al (2013), Frontiers in Computational Neuroscience

55

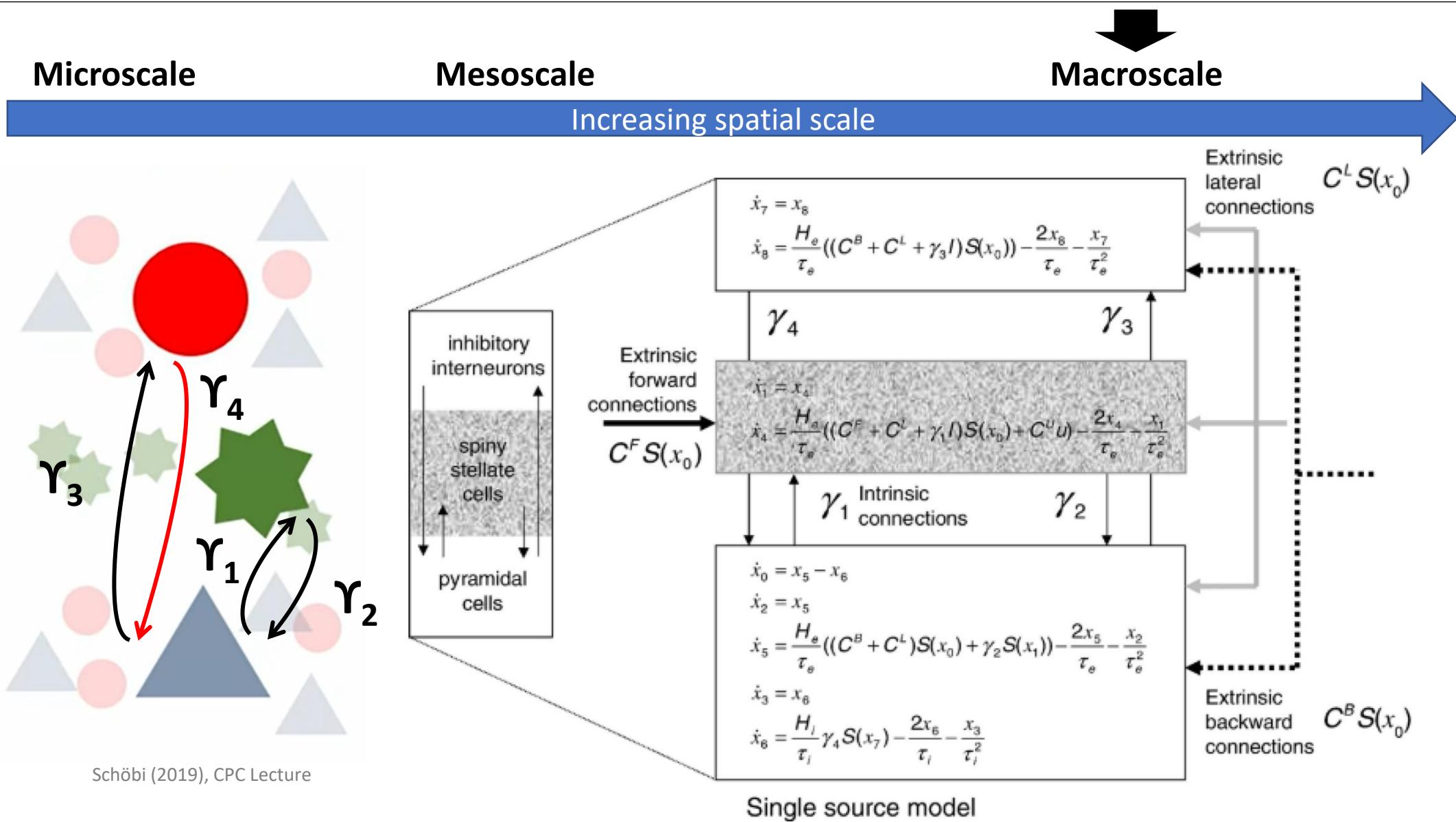
# Source modelling at the macroscale



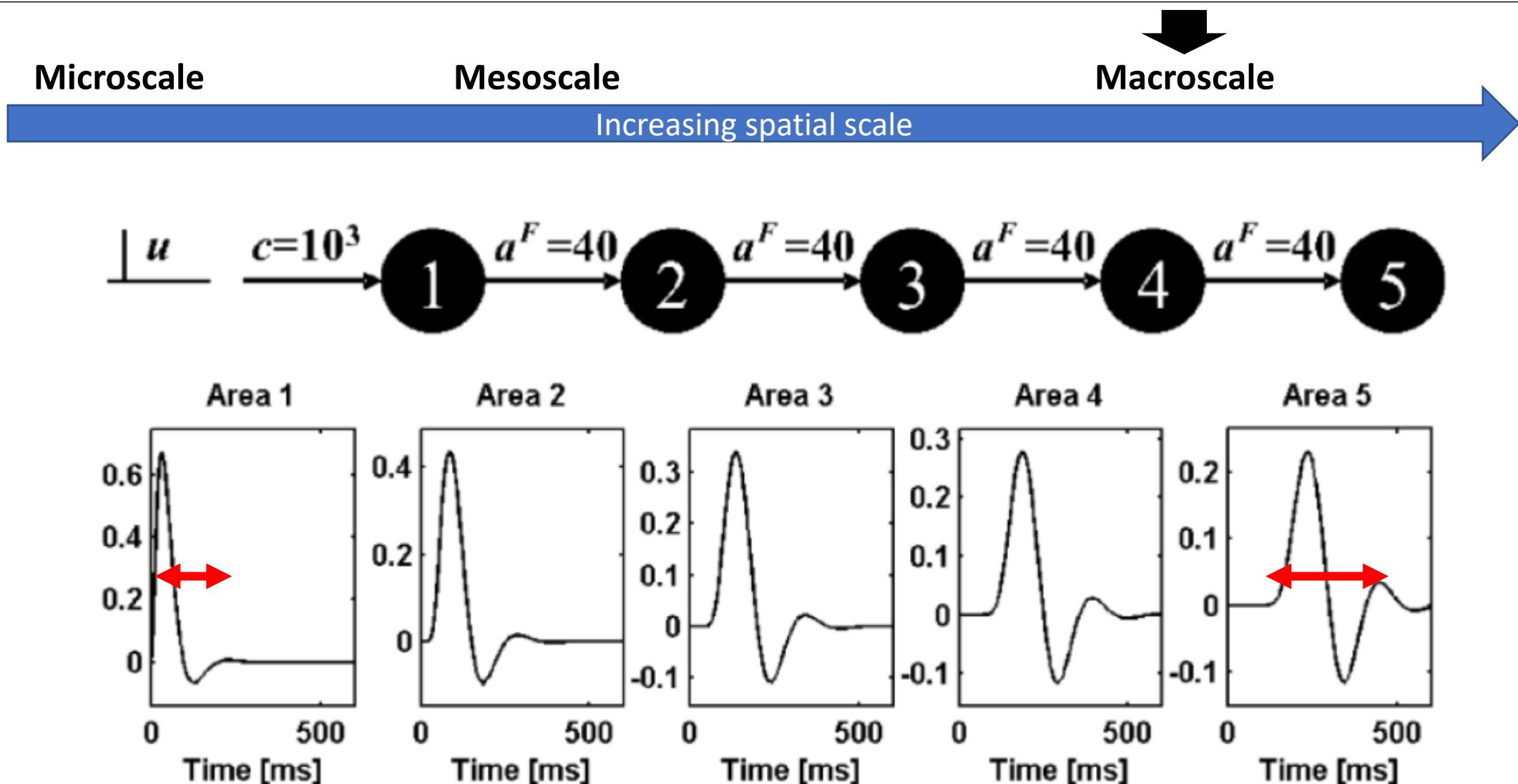
# Source modelling at the macroscale



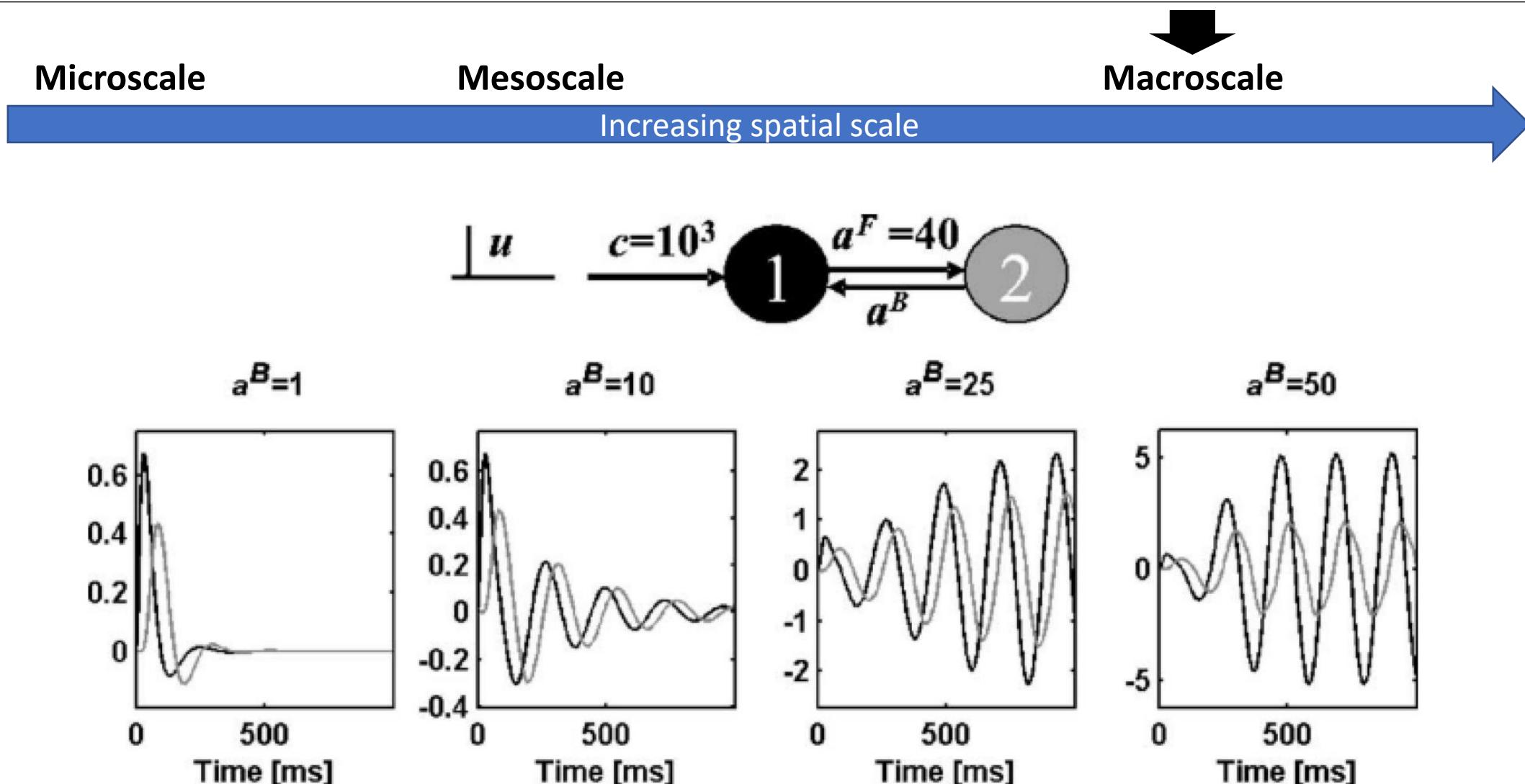
# Source modelling at the macroscale



# Source modelling at the macroscale

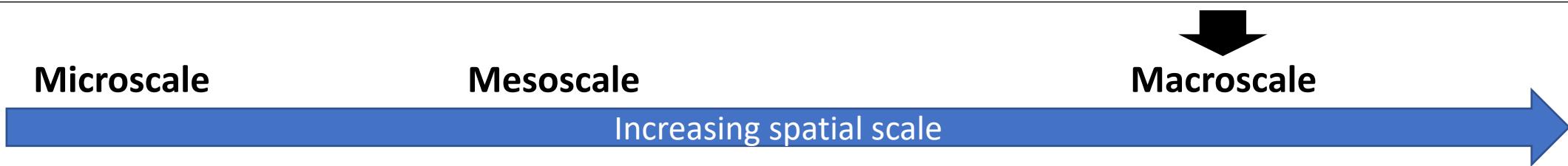


# Source modelling at the macroscale

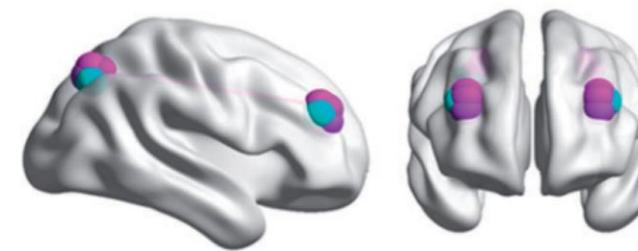


# Scales of analysis

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**Brain Networks**  
5-20 cm<sup>2</sup>



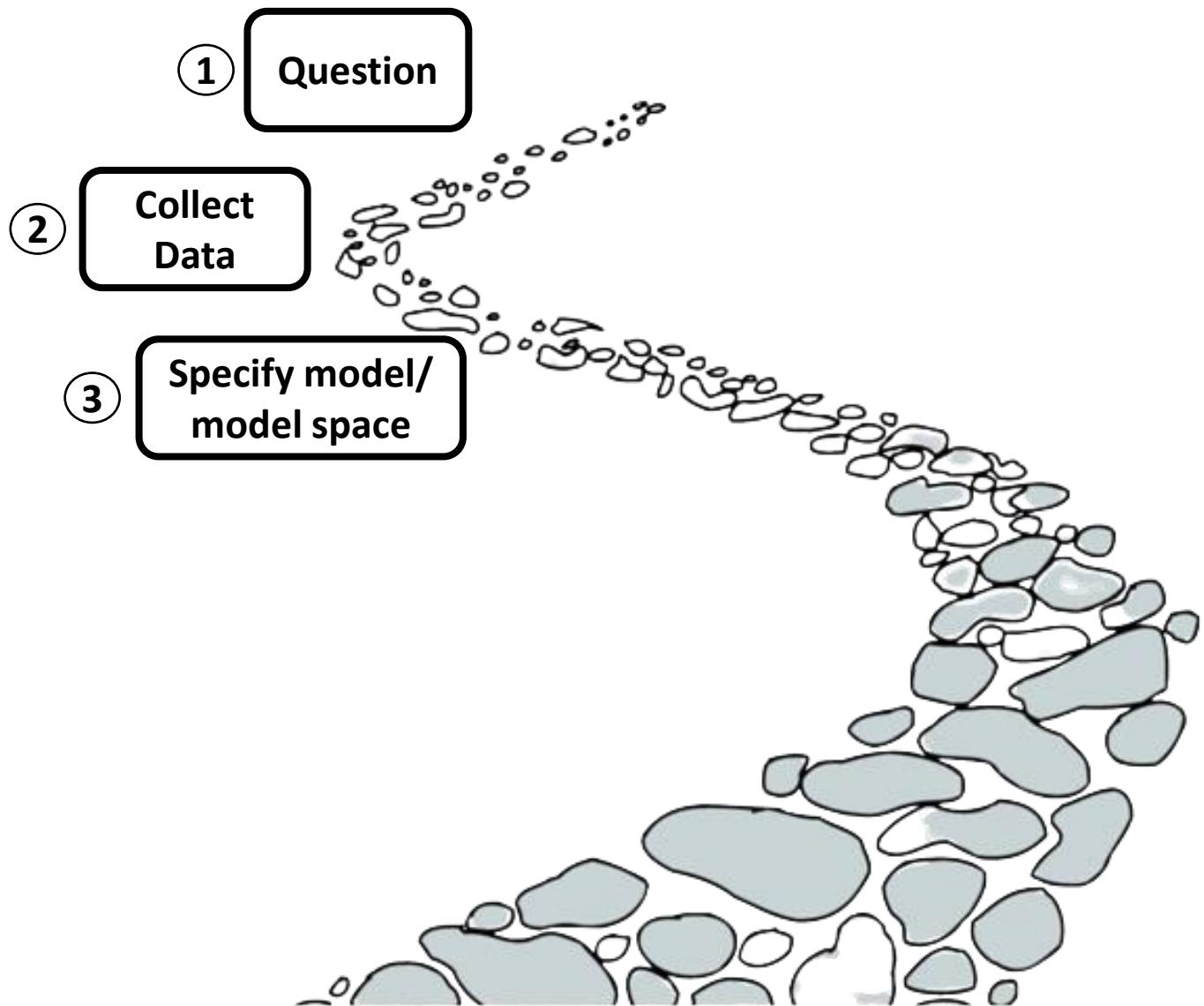
Symmonds et al. (2018), Brain

# Specify model/model space

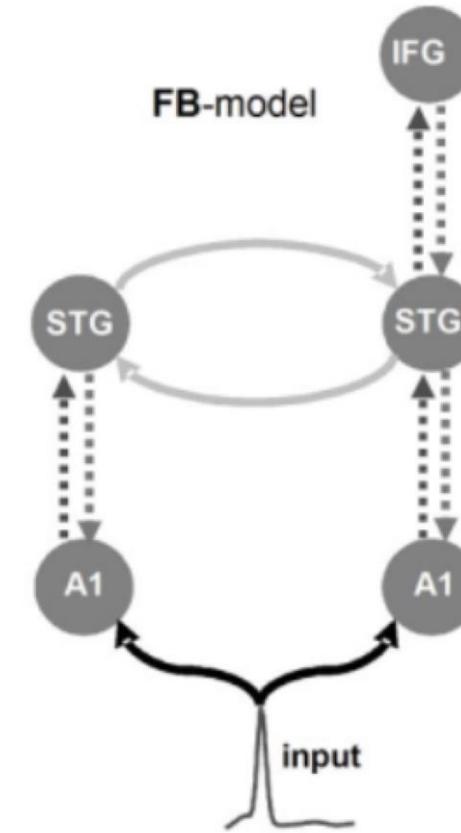
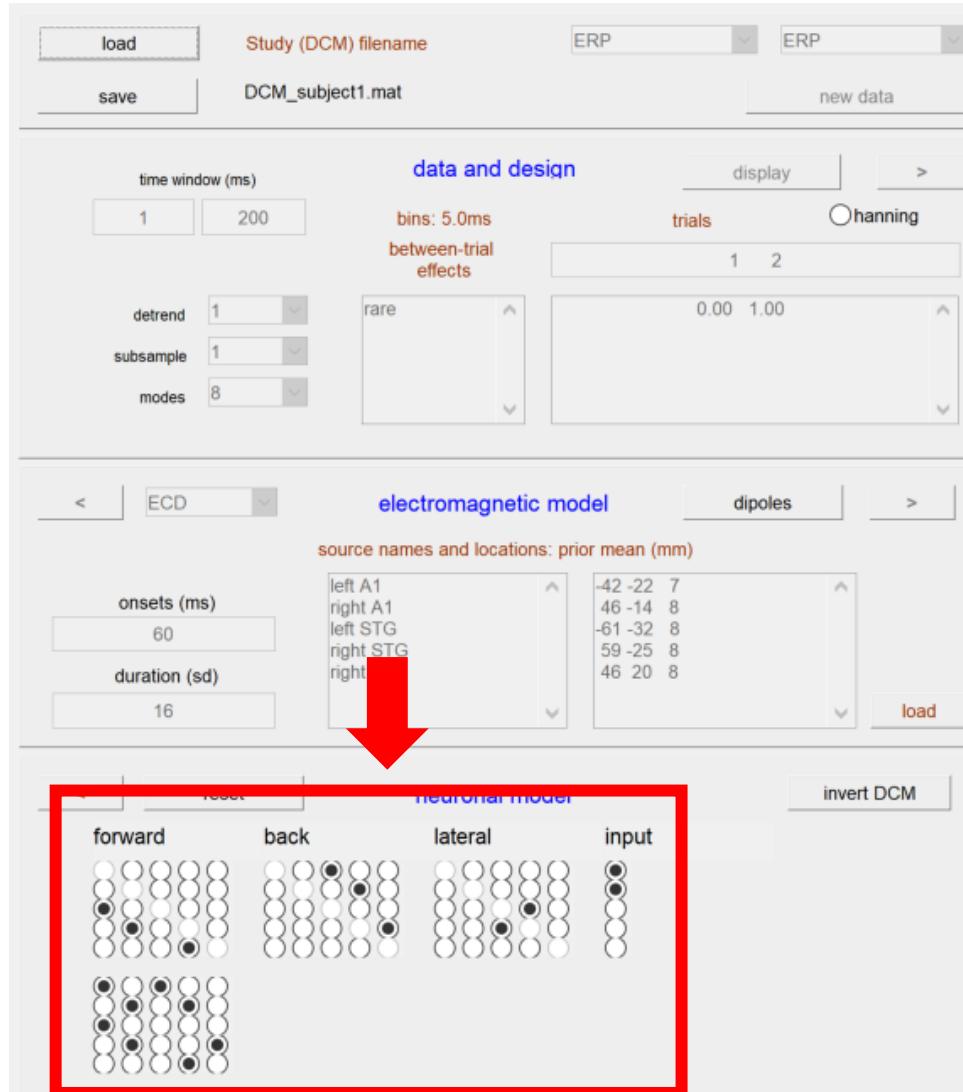
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## Steps

- Translate your question into a model comparison or a parameter inference problem
- Select regions
- Select a variant of DCM
- Example: The “ERP” model
- **Specify connectivity architecture**

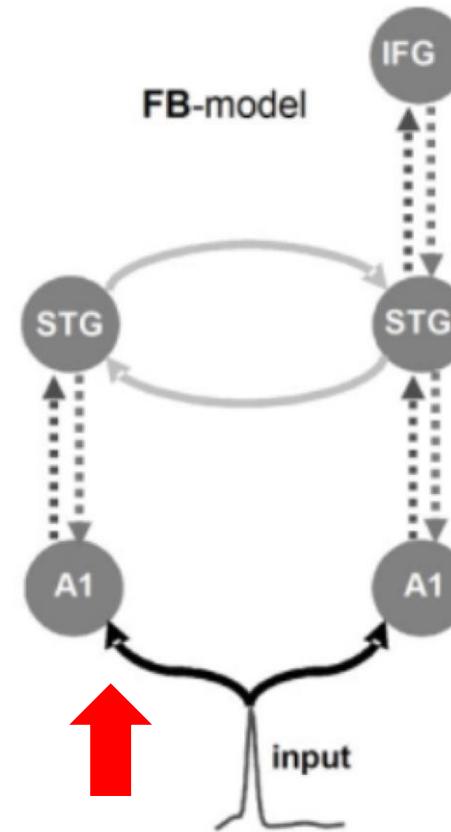
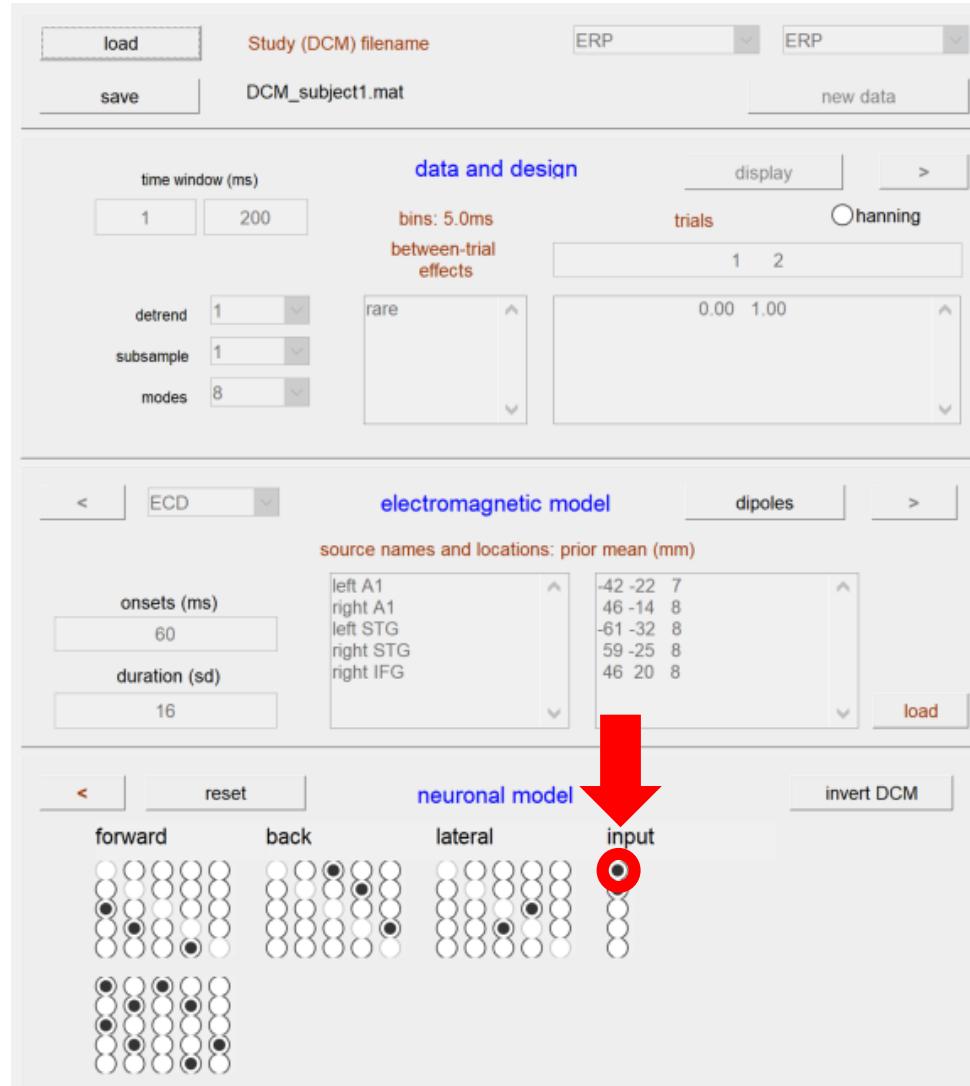


# The DCM analysis path



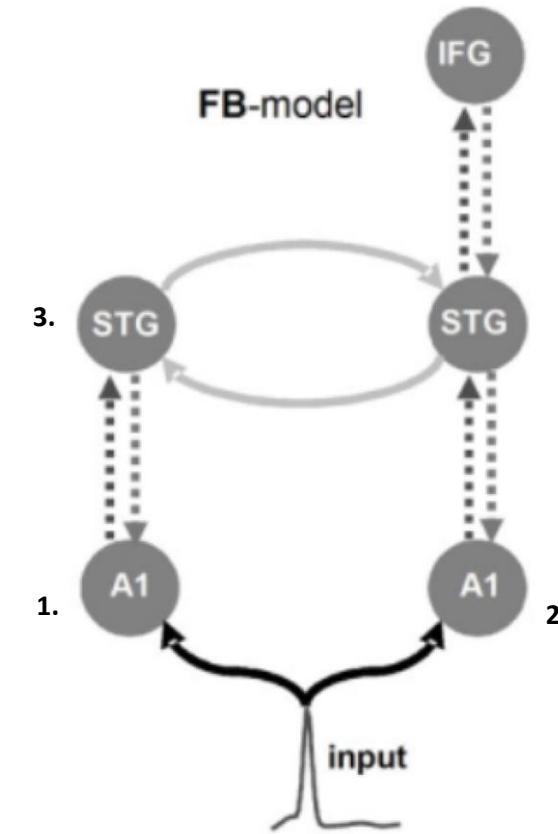
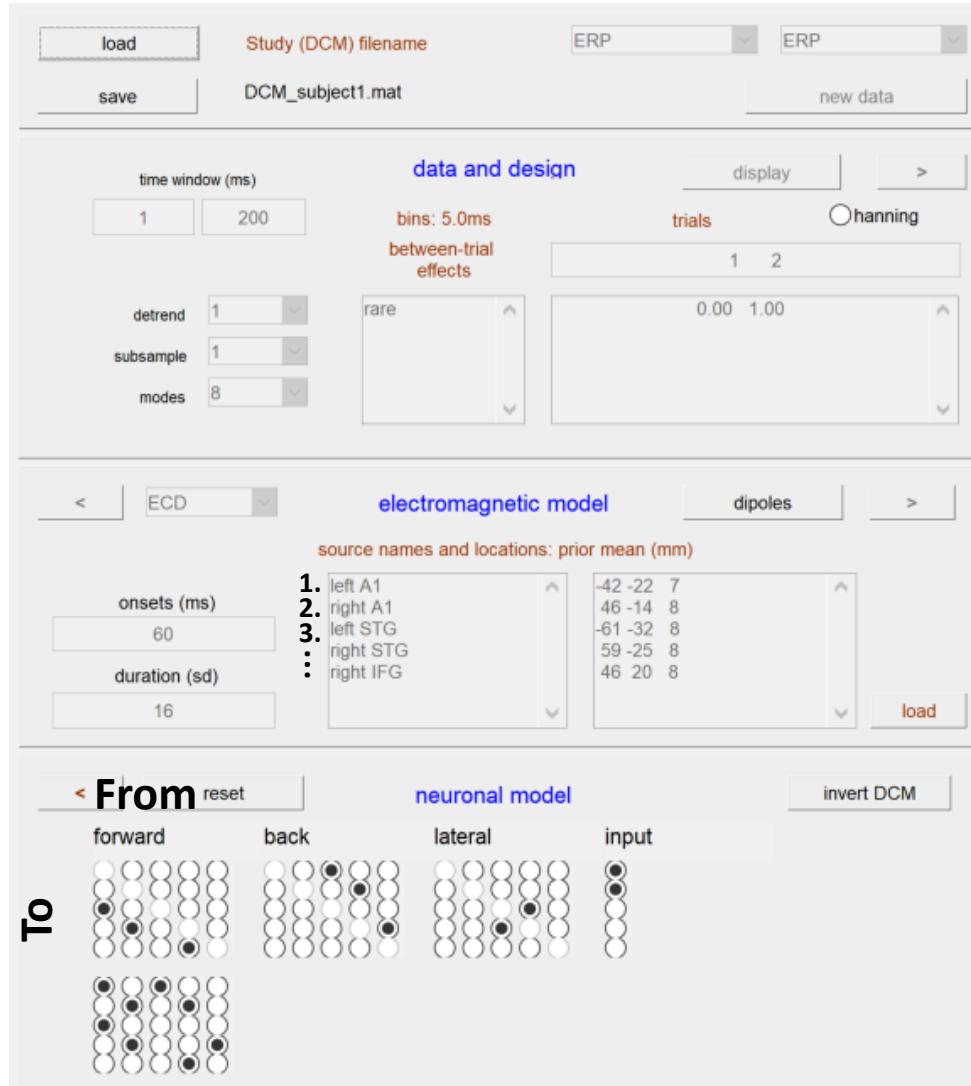
Garrido et al. (2009), HBM

# The DCM analysis path



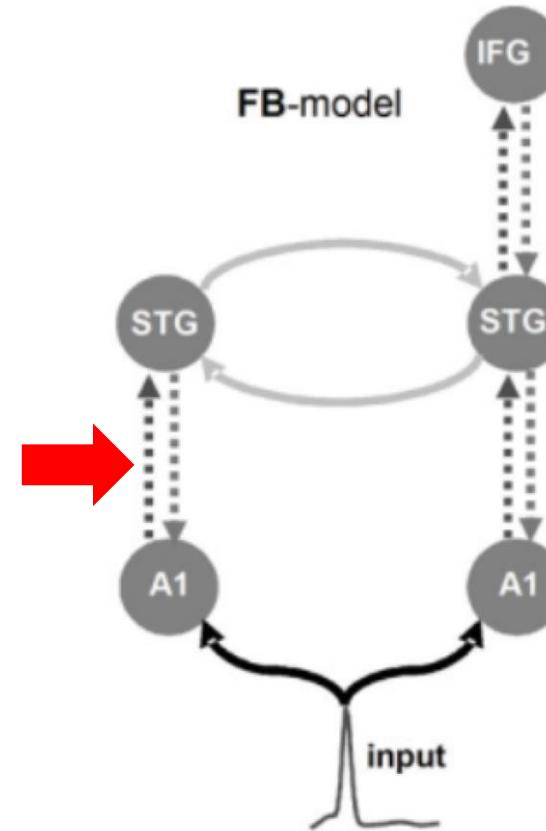
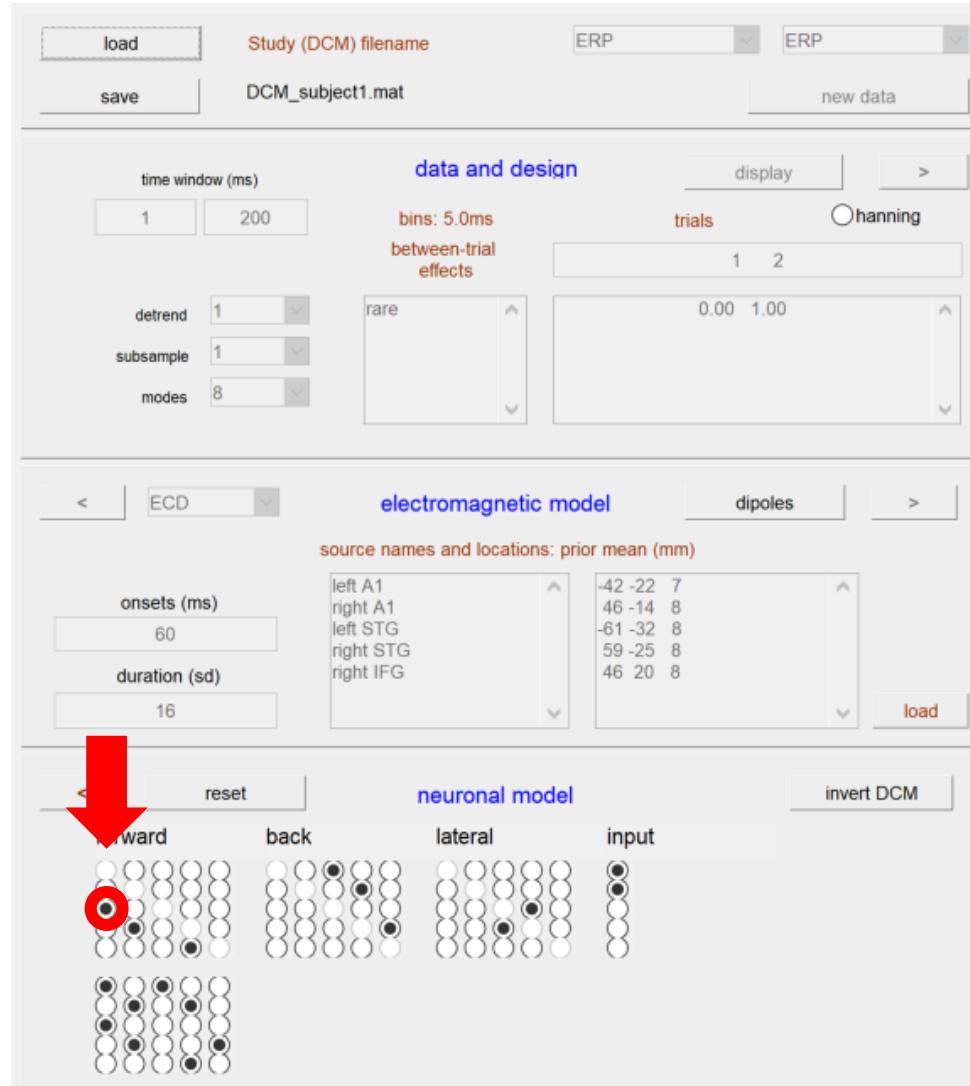
Garrido et al. (2009), HBM

# The DCM analysis path



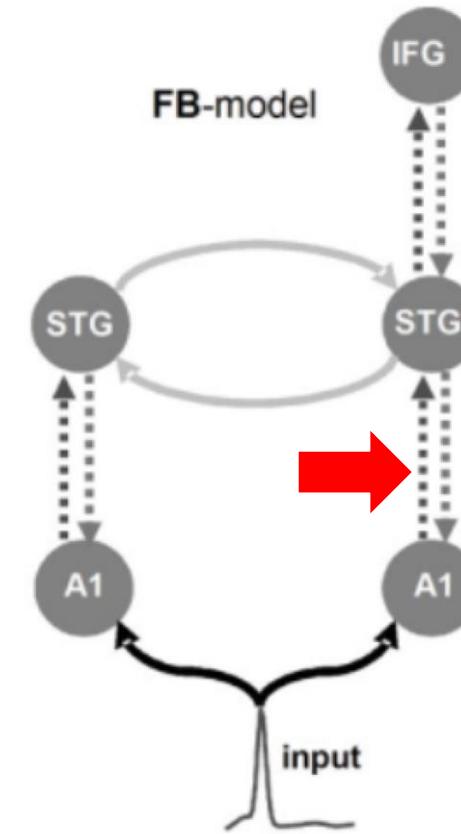
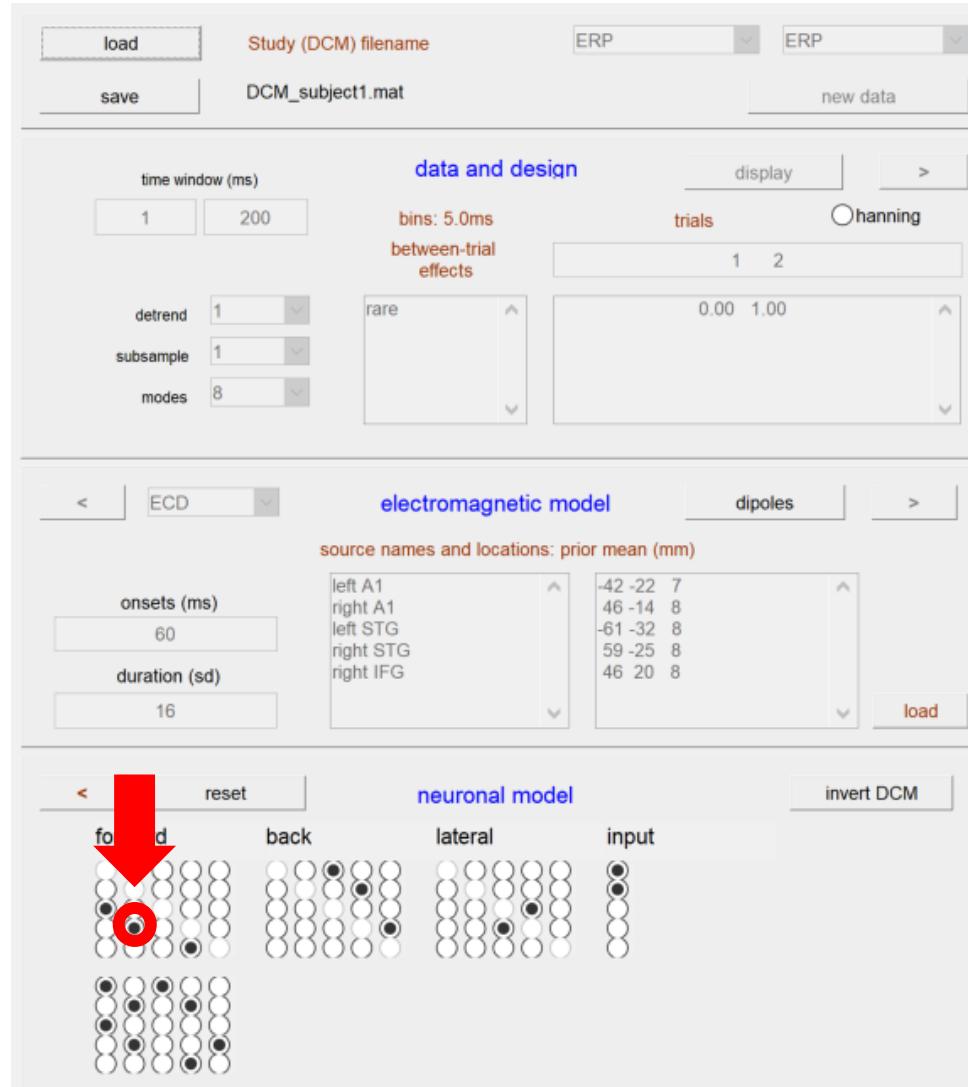
Garrido et al. (2009), HBM

# The DCM analysis path



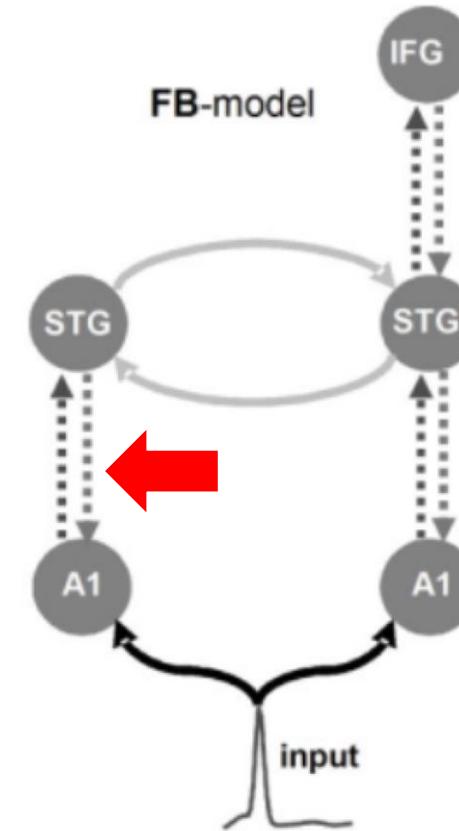
Garrido et al. (2009), HBM

# The DCM analysis path



Garrido et al. (2009), HBM

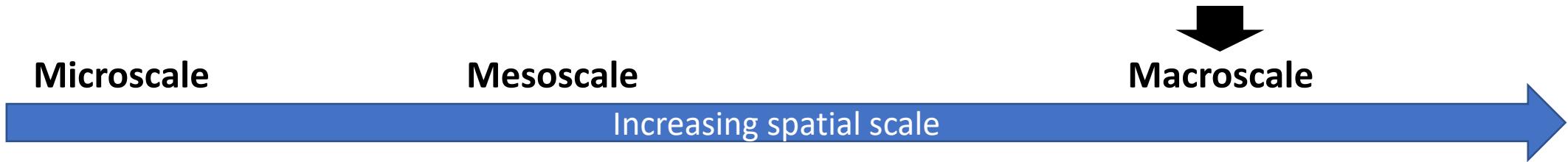
# The DCM analysis path



Garrido et al. (2009), HBM

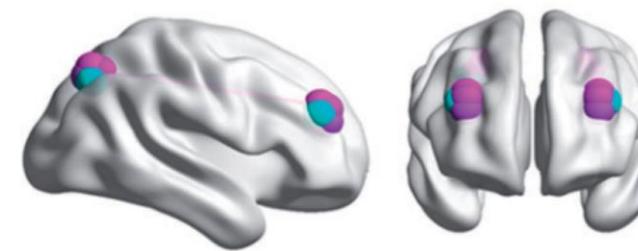
# Scales of analysis

---



**Brain Networks**

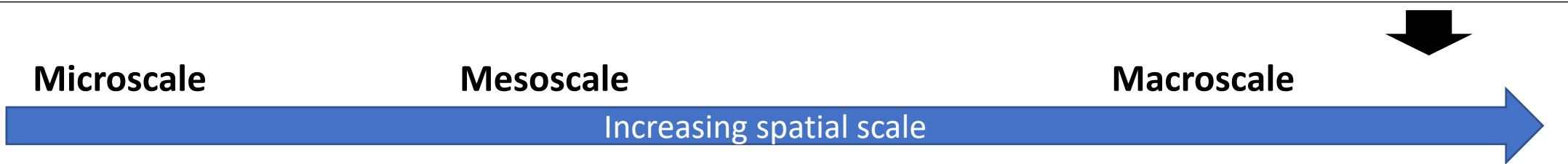
5-20 cm<sup>2</sup>



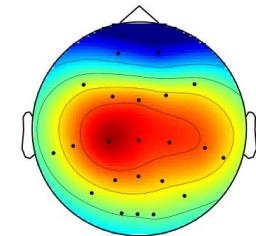
Symmonds et al. (2018), Brain

# Scales of analysis

---



**Scalp potentials**  
30-38 cm<sup>2</sup>



# Scales of analysis



**Scalp potentials**  
30-38 cm<sup>2</sup>

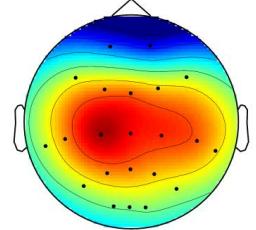
Scalp responses

Leadfield

$$y = Lx + \epsilon$$

↑

Neuronal source activity

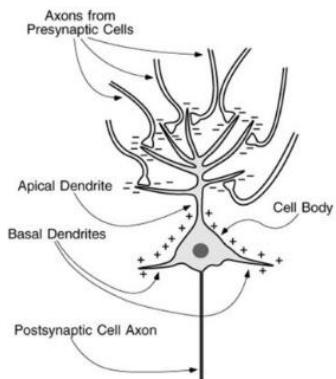


# Scales of analysis



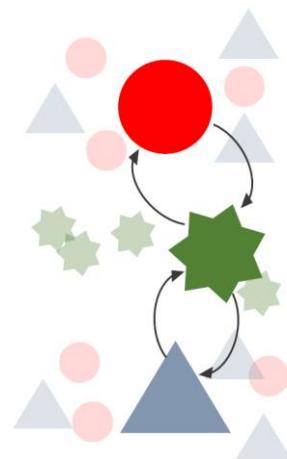
## Single cell

1-10  $\mu\text{m}^2$



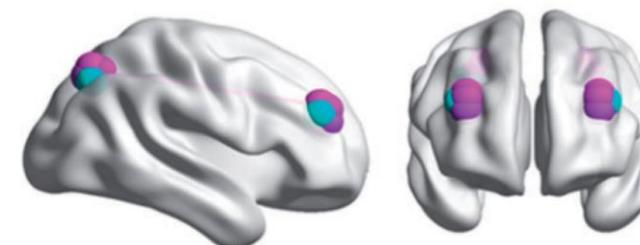
## Cortical Column

1-10  $\text{mm}^2$  to 1-5  $\text{cm}^2$



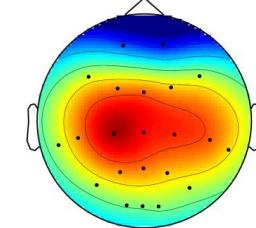
## Brain Networks

5-20  $\text{cm}^2$



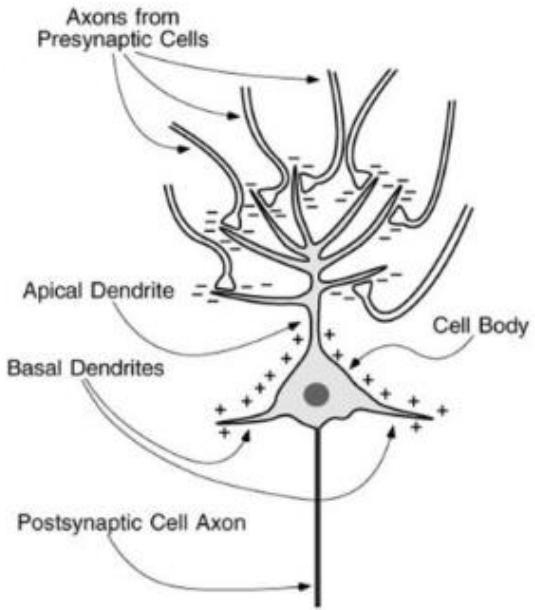
## Scalp potentials

30-38  $\text{cm}^2$

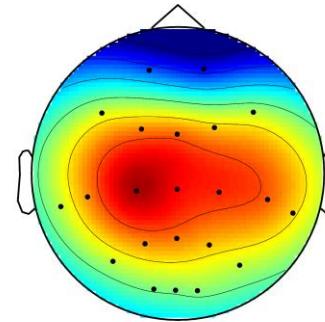


# What do we measure with EEG?

---



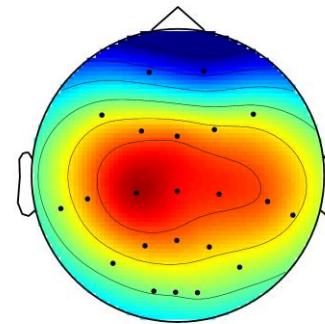
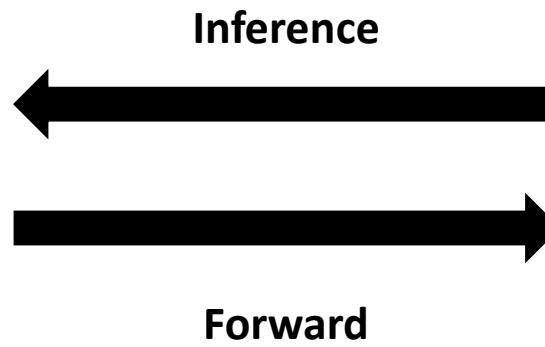
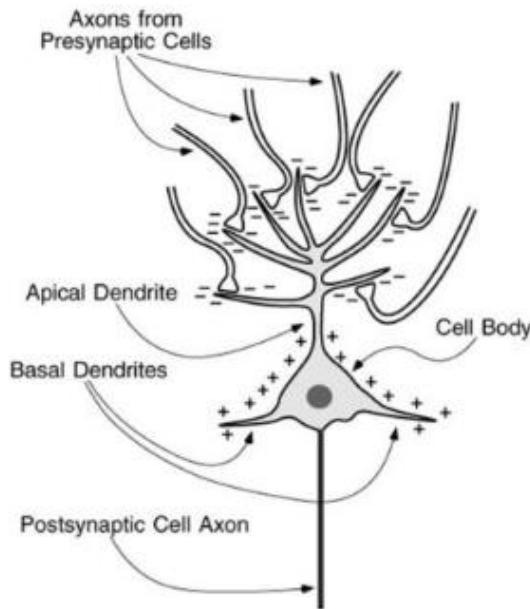
**Forward**



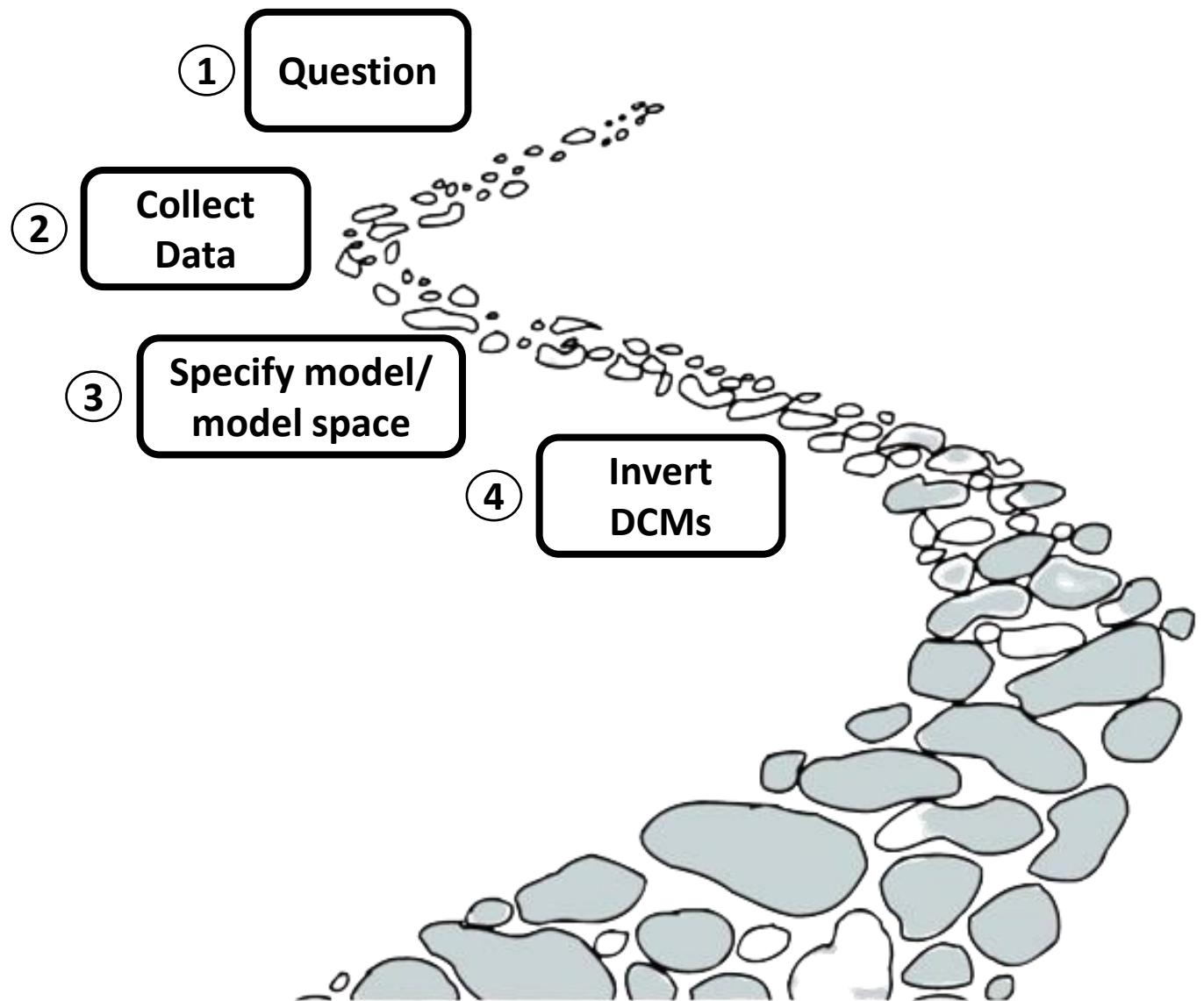
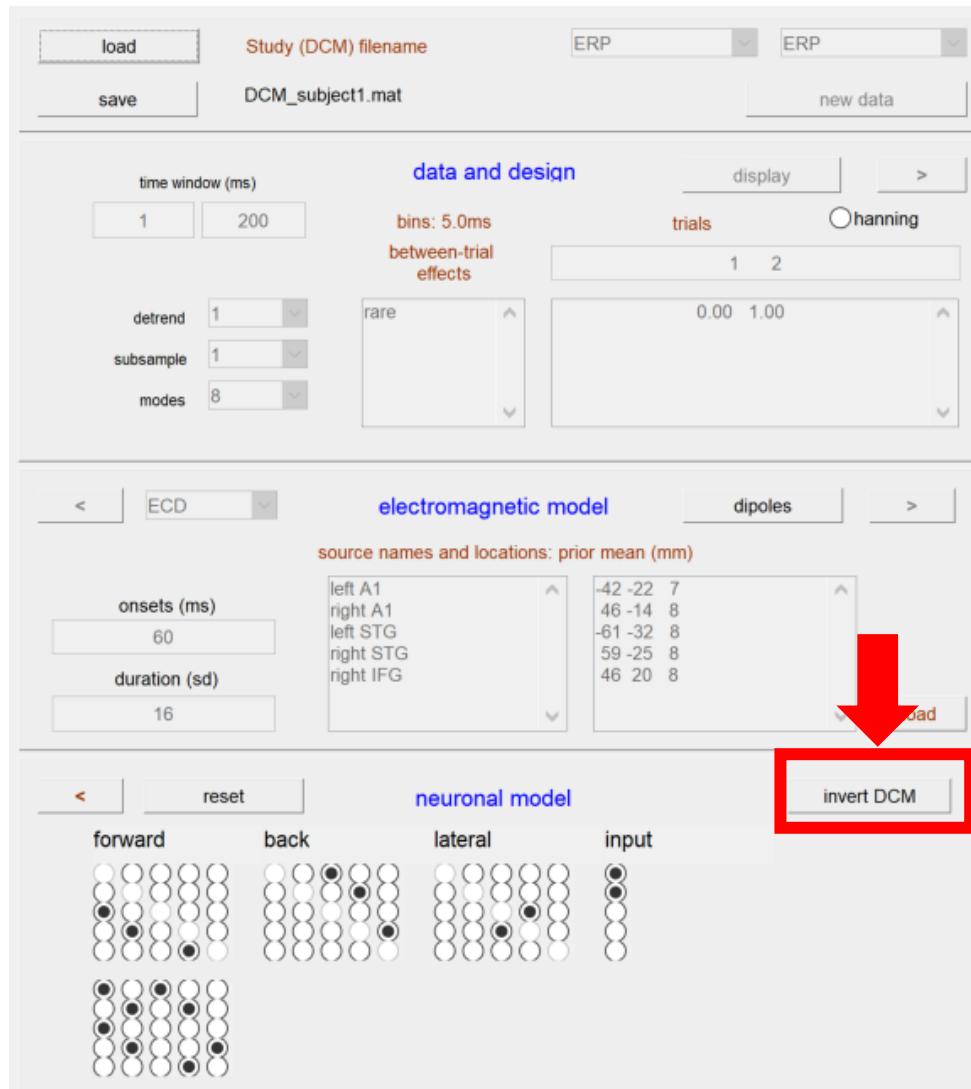
# What do we measure with EEG?

## Question

- Can we make inferences about properties of the neuronal sources that generate these signals?

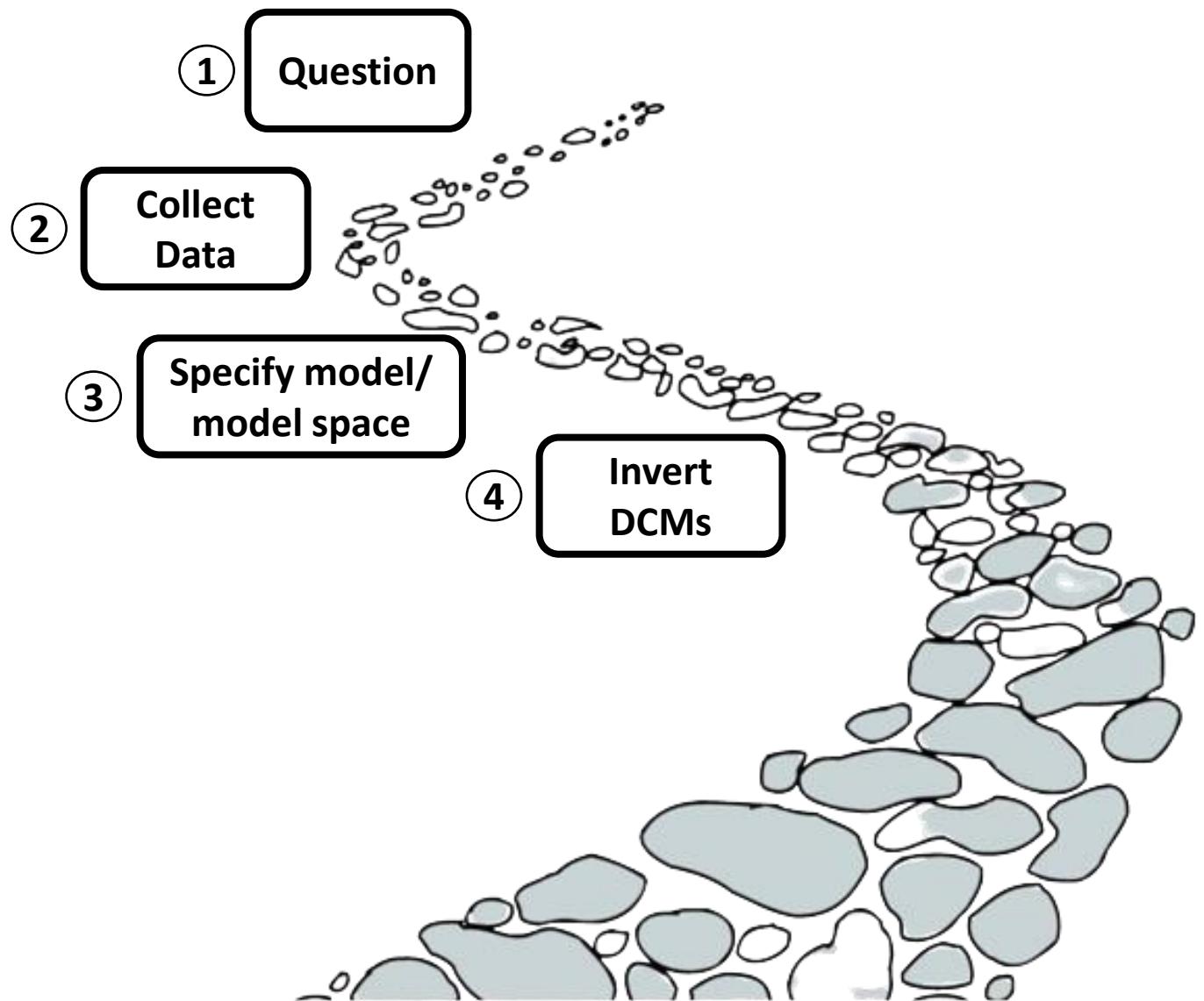


# The DCM analysis path

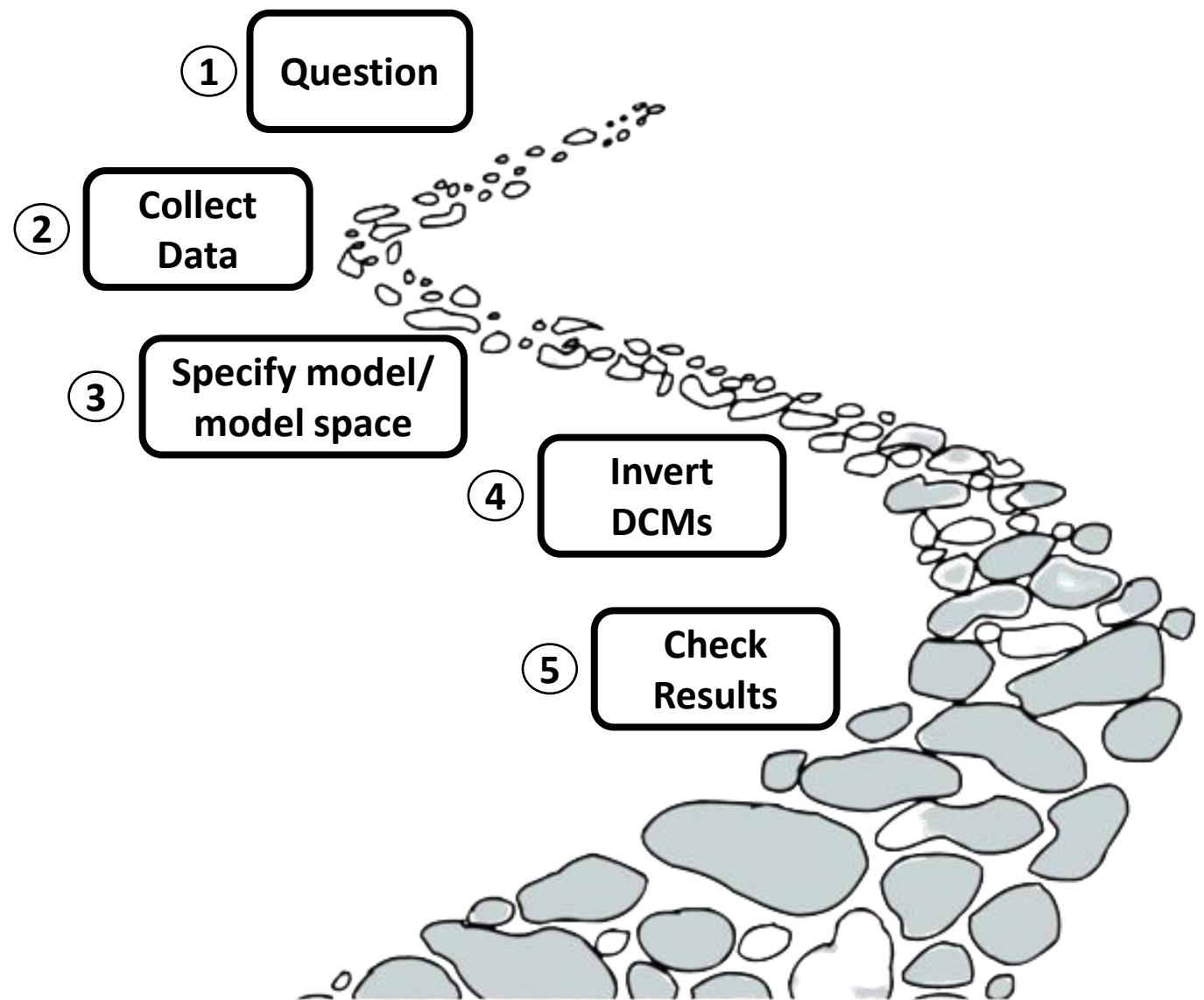
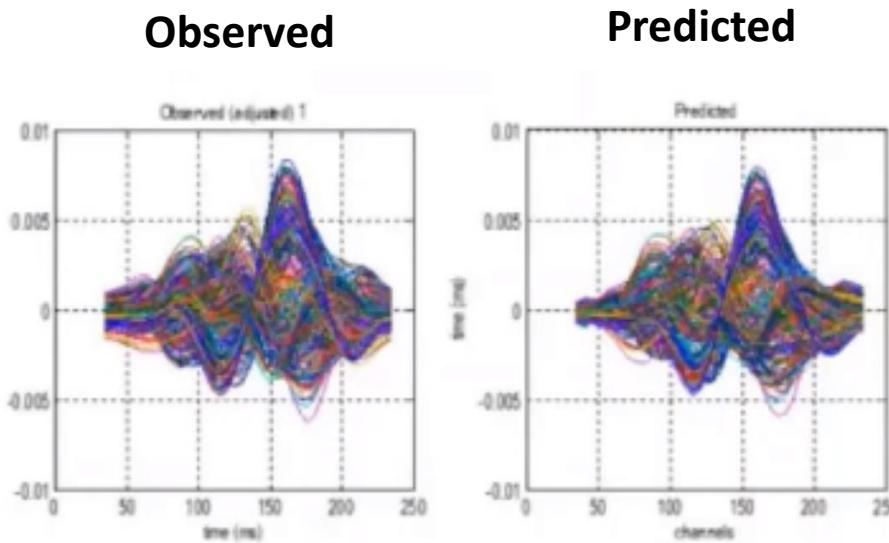


# The DCM analysis path

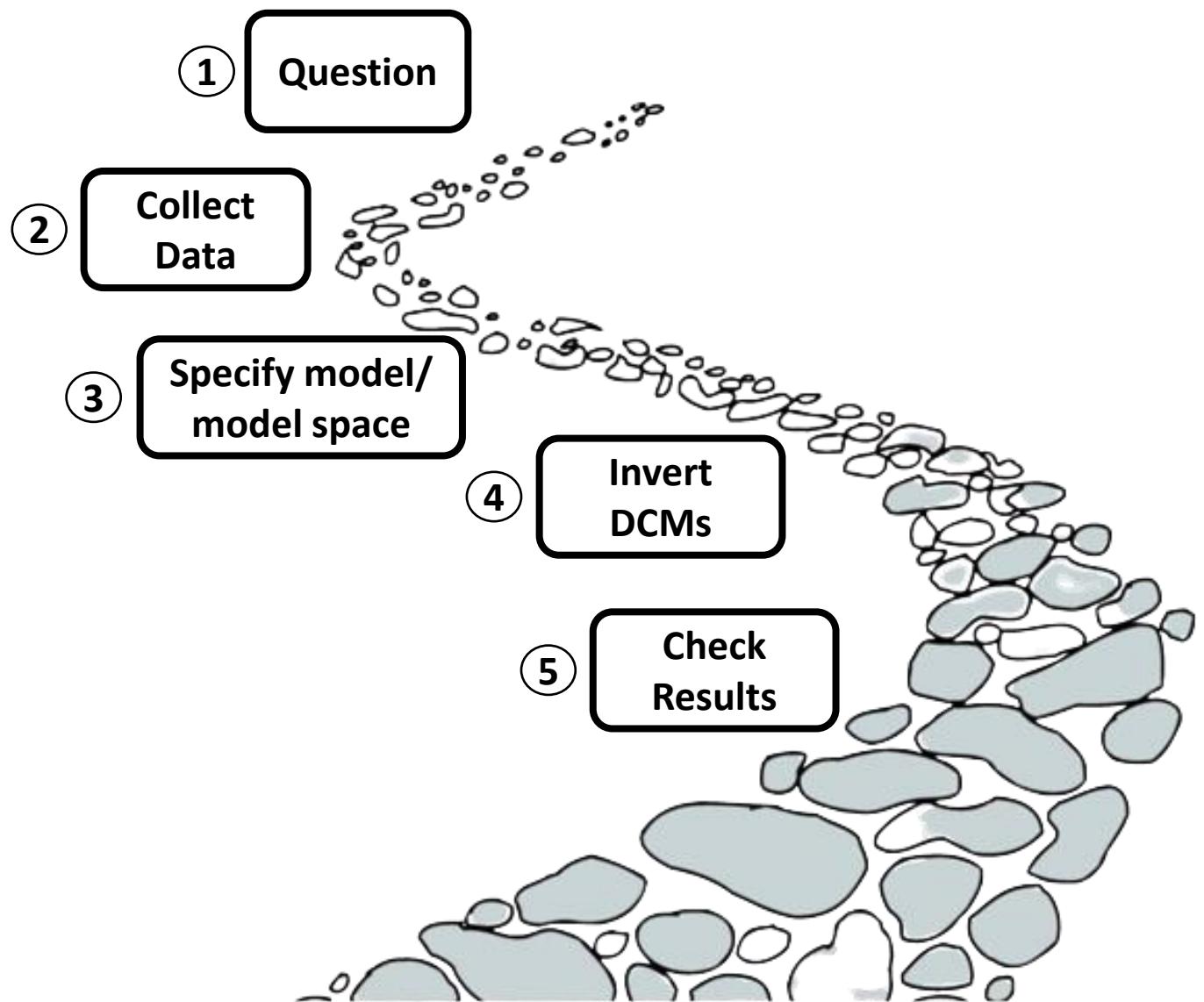
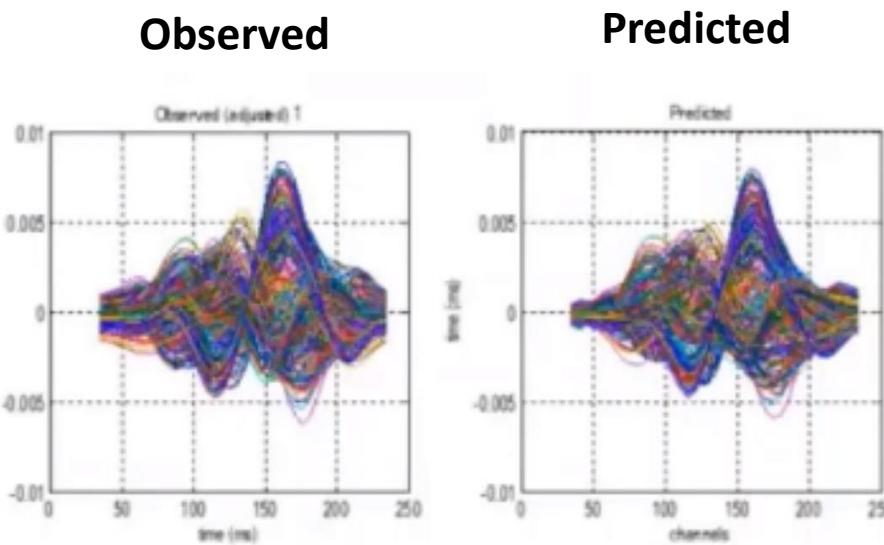
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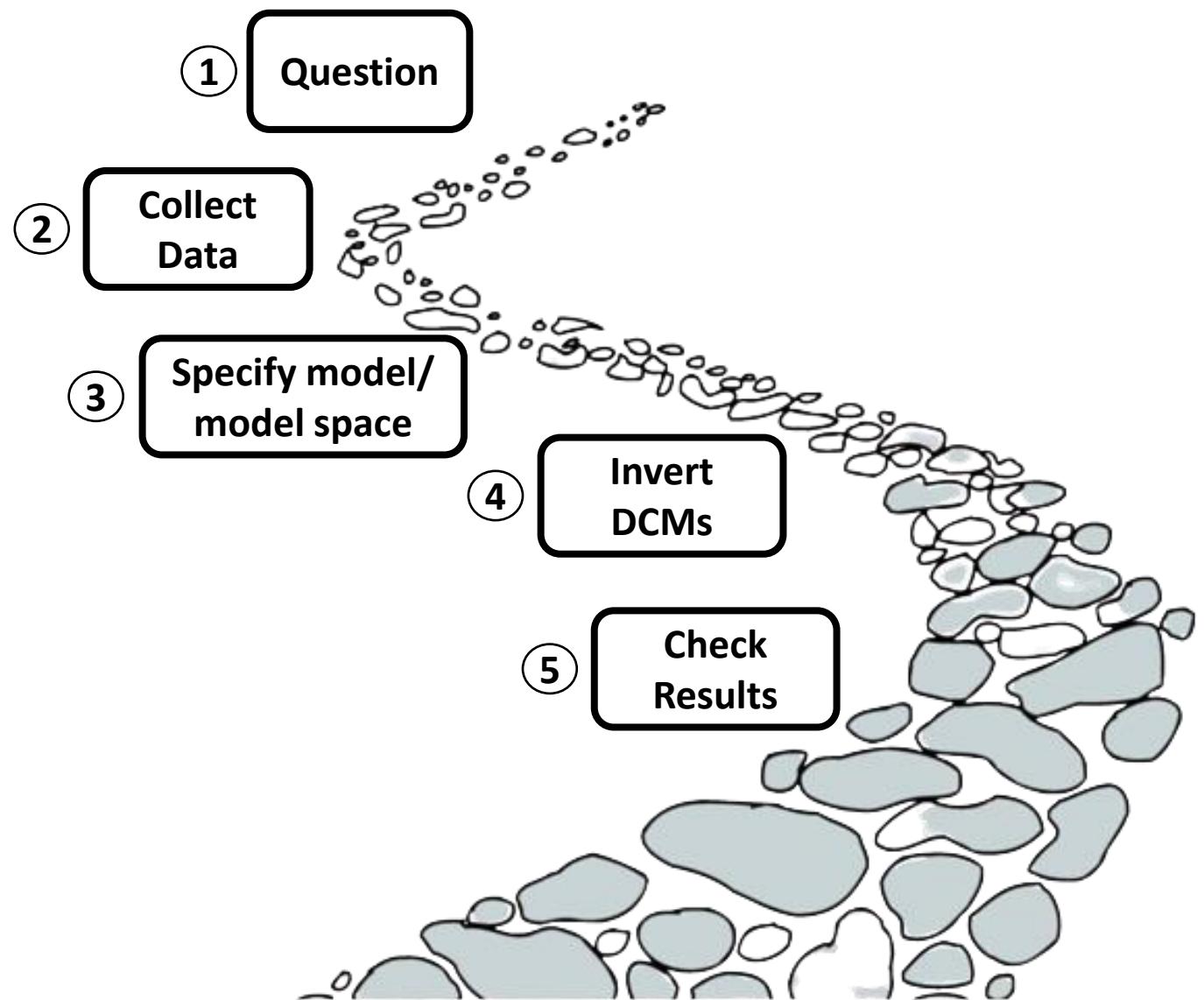
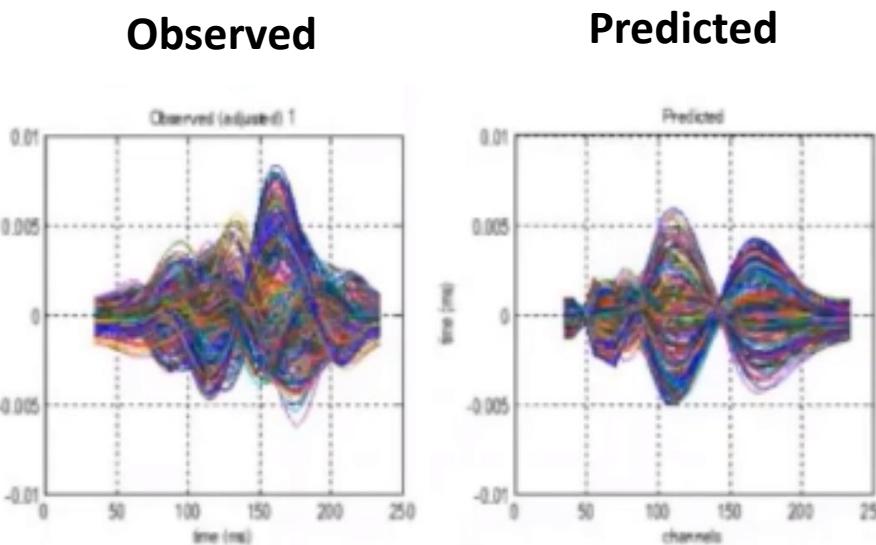
# The DCM analysis path



# The DCM analysis path



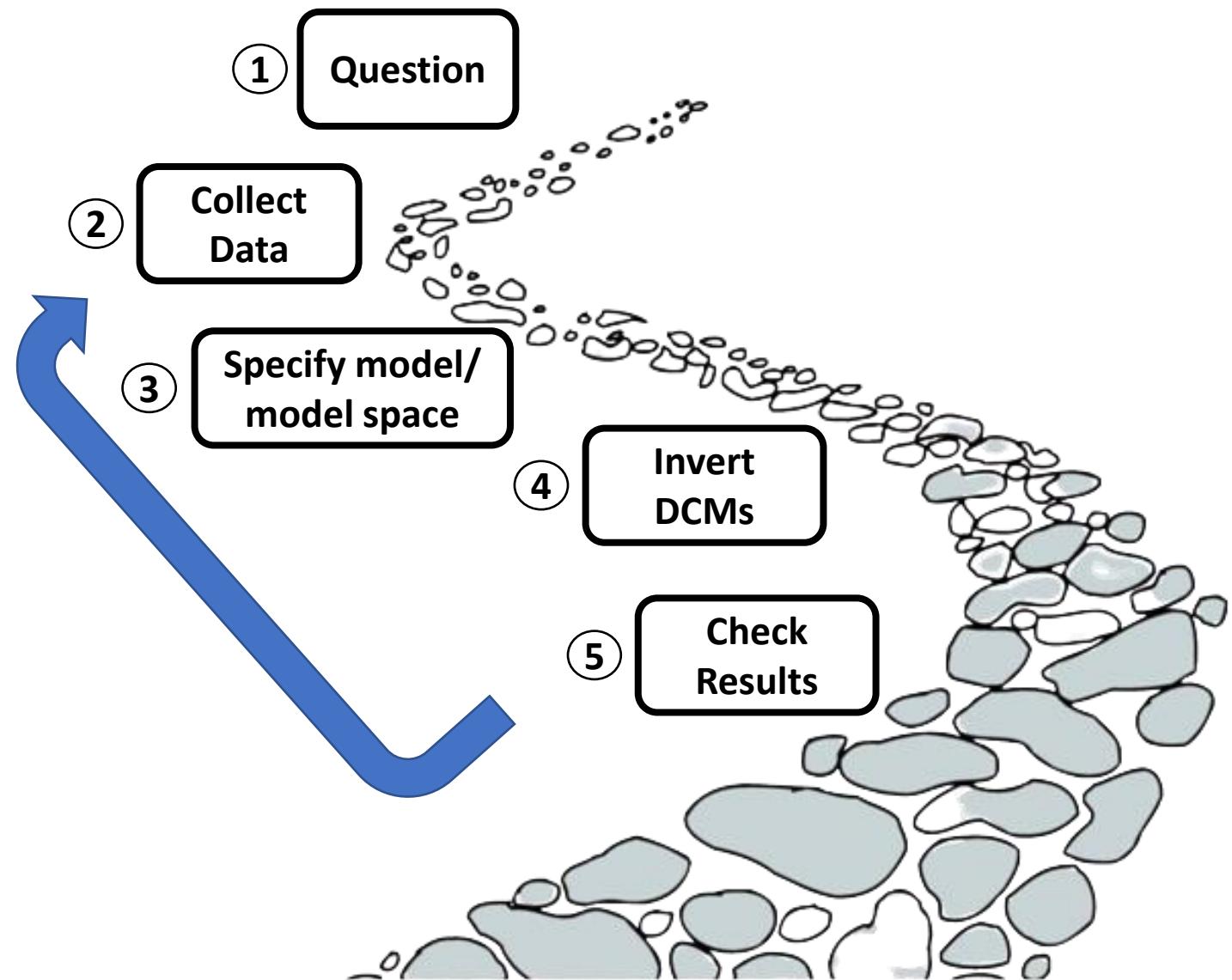
# The DCM analysis path



# The DCM analysis path

## Check your data

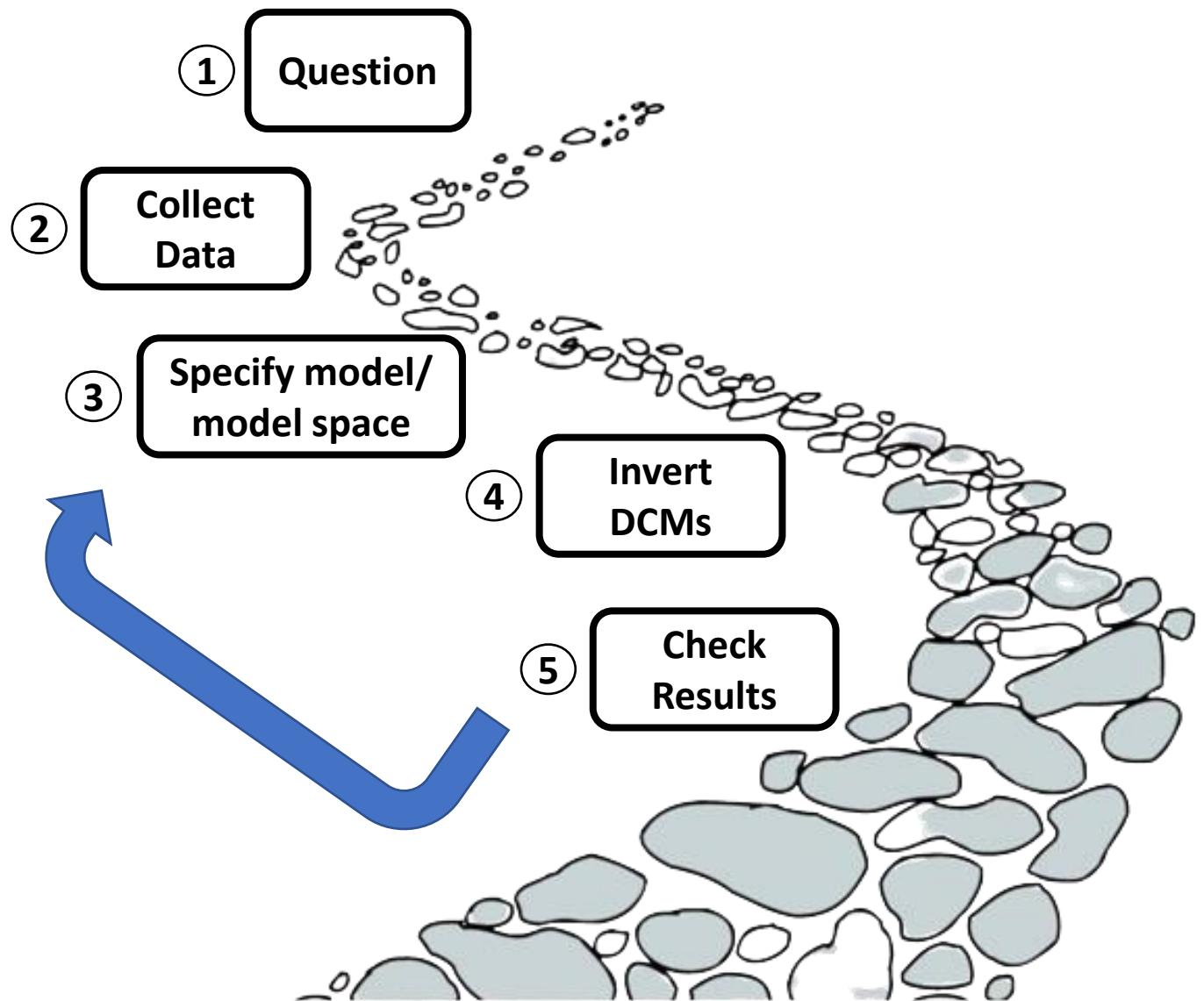
- Did something go wrong with preprocessing?
- Are there artefact?
- Is there high-frequency noise?
- ...



# The DCM analysis path

## Check your sources

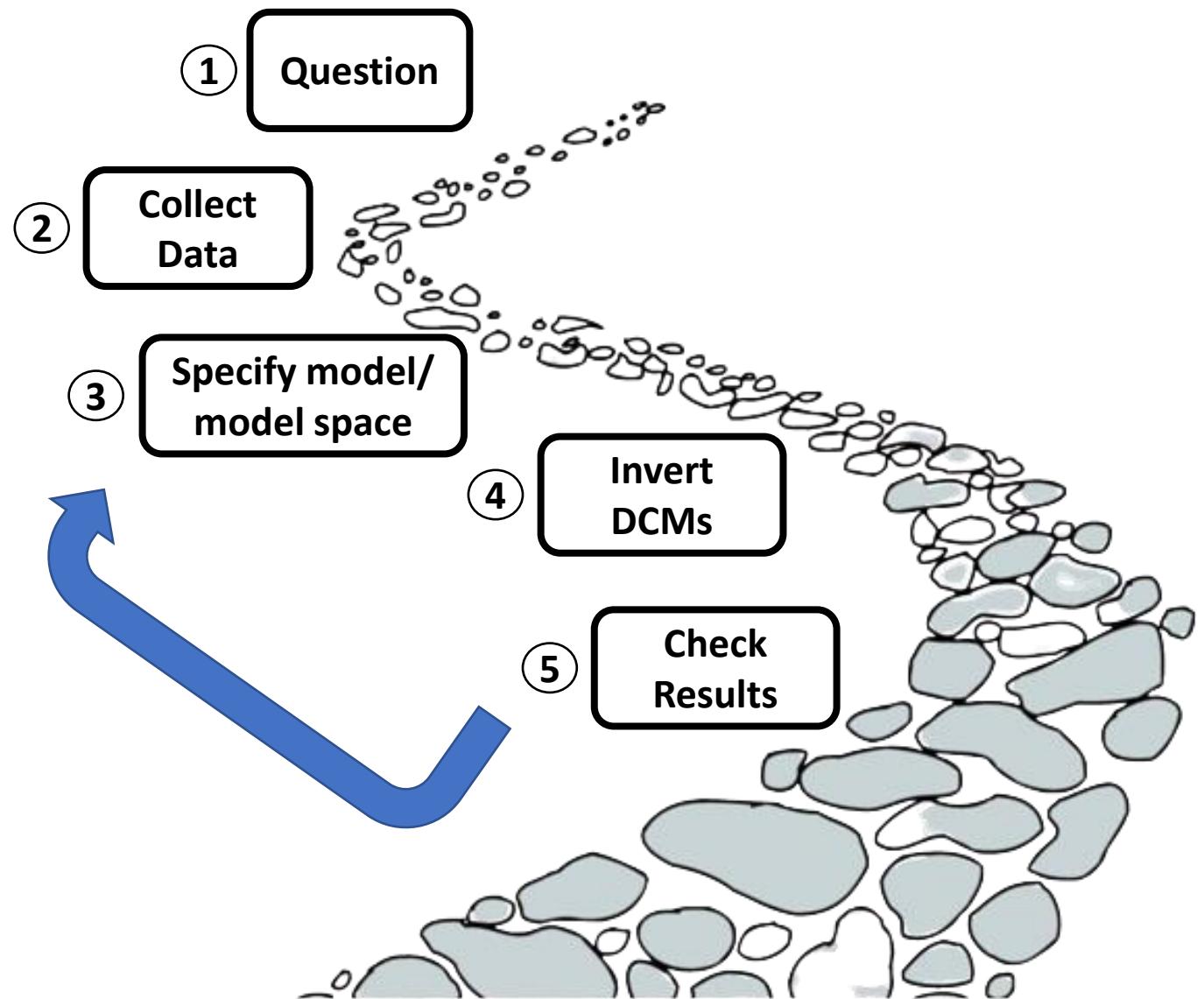
- Is there a relevant source that you have not considered in your network?



# The DCM analysis path

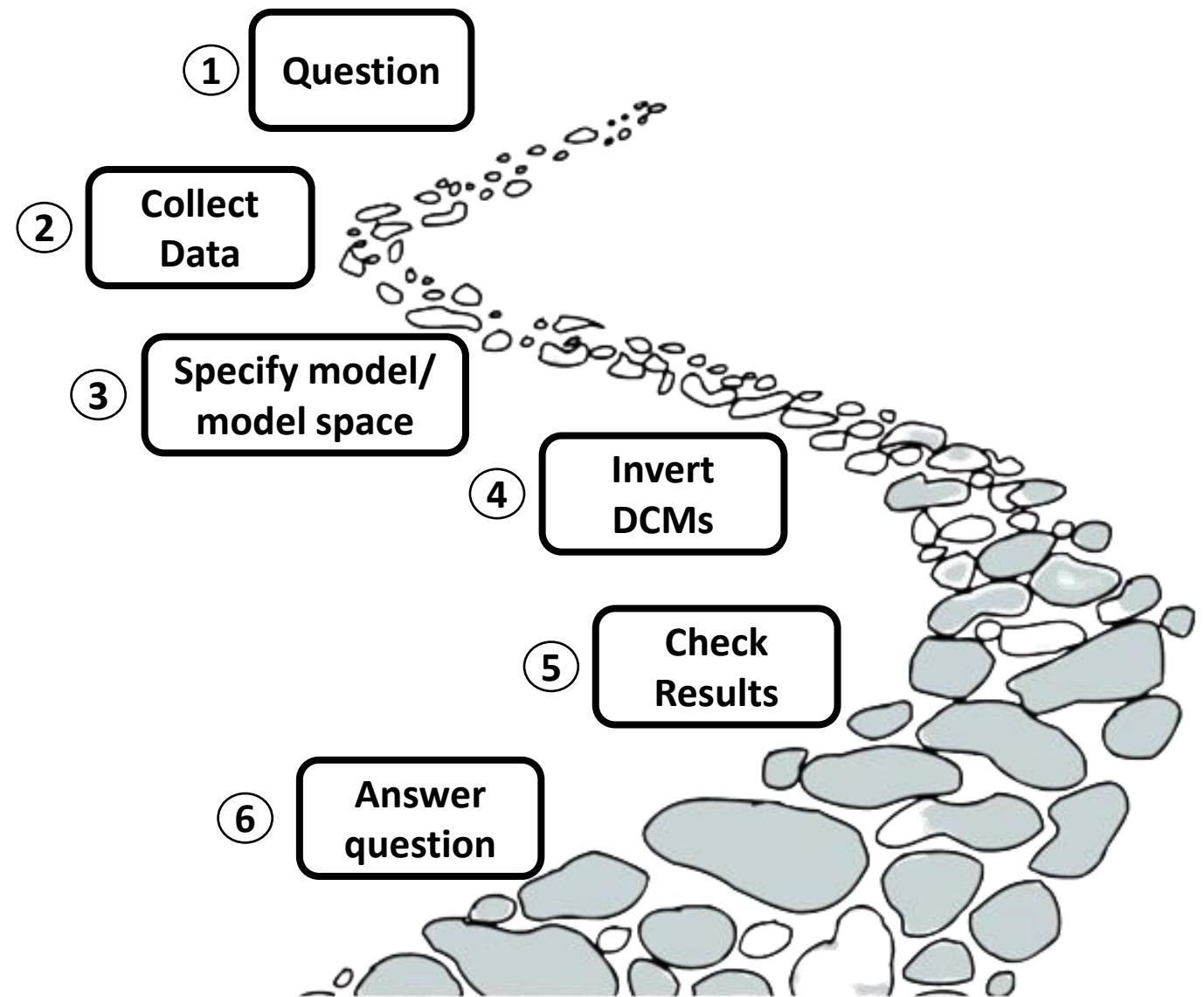
## Check your model

- If you simulate from your model can you produce the effects in simulations?
- Do you need to estimate additional parameters?
- Pick a more complex model?



# The DCM analysis path

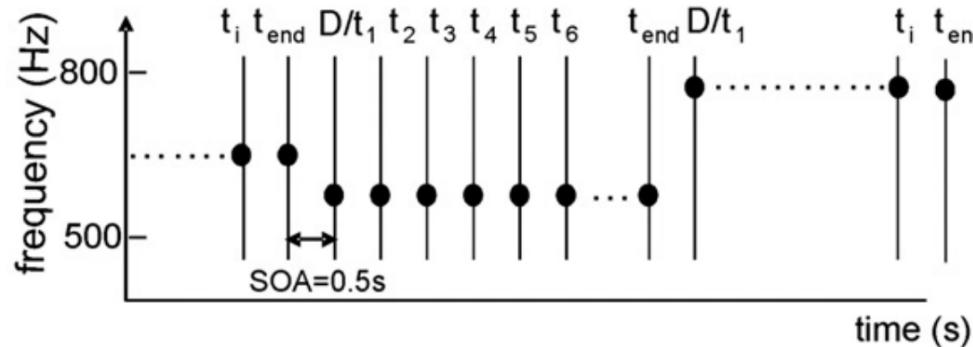
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# Example 1

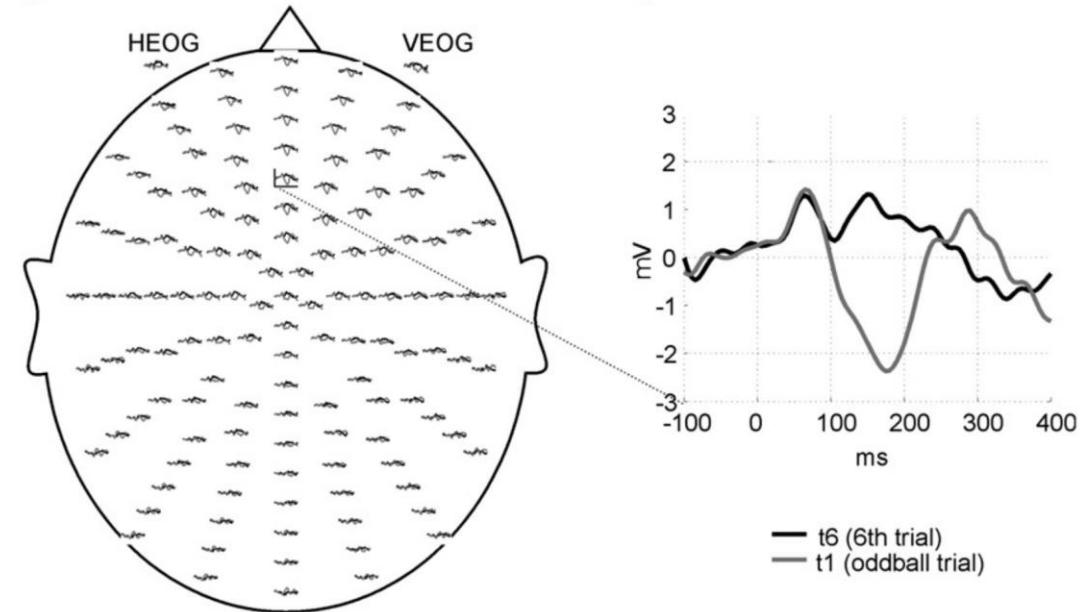
What network explains the mismatch negativity best?

- Roving paradigm to elicit MMN



$D/t_1$  = deviant

$t_i$  = trial  $i$ ,  $1 \leq i \leq 11$

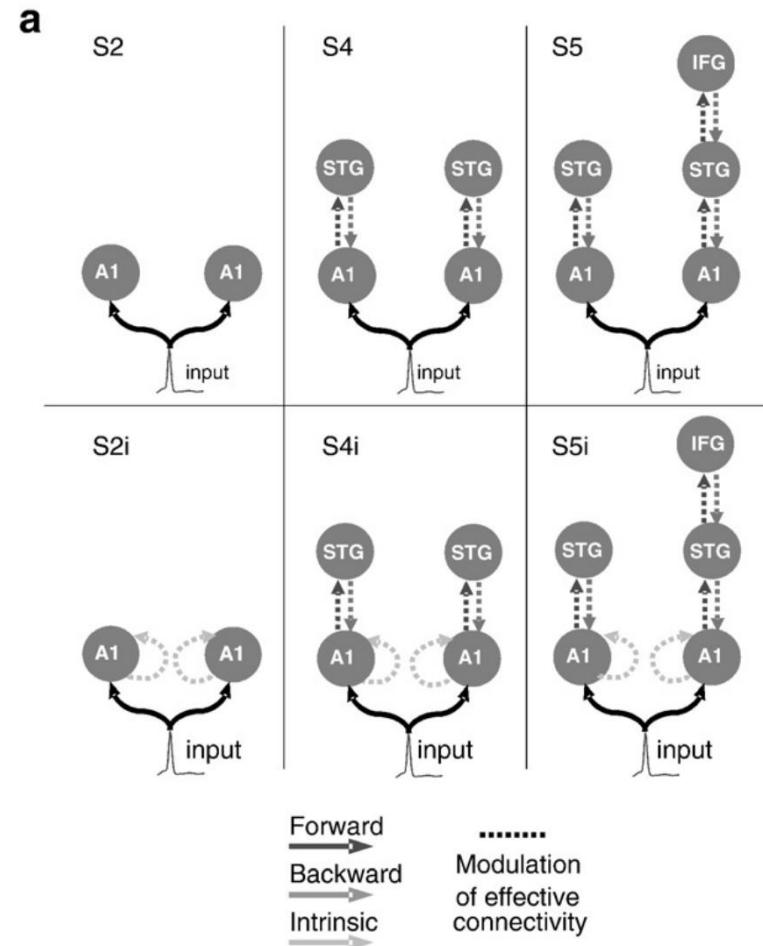
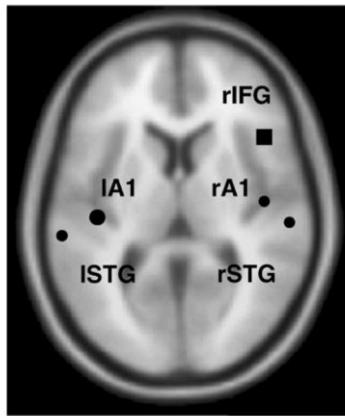


Garrido et al. (2006), NeuroImage

# Example 1

What network explains the mismatch negativity best?

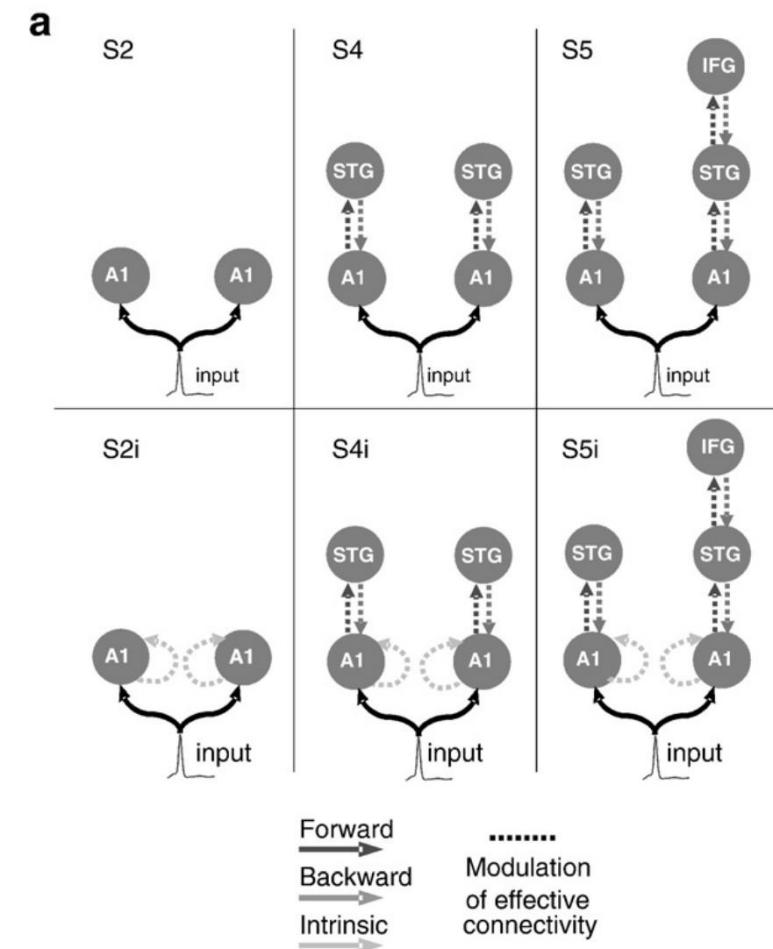
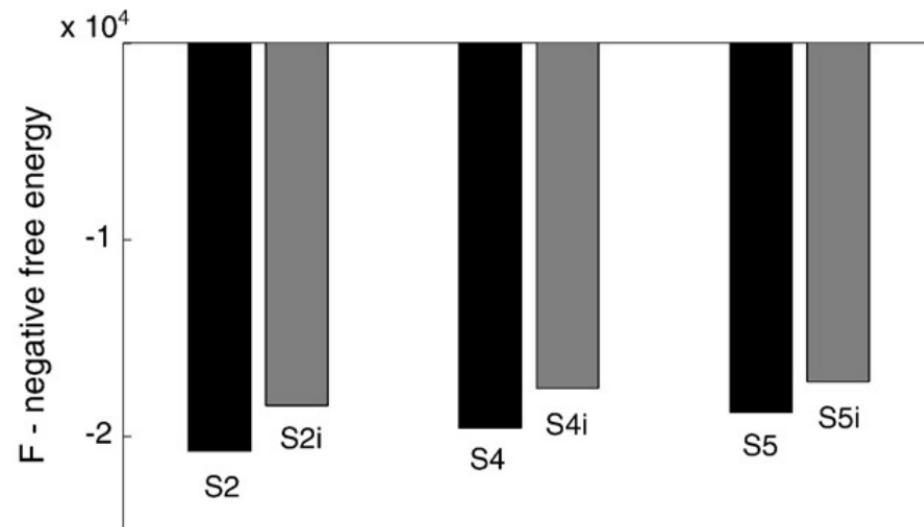
- Model space: Networks with different regions



# Example 1

## What network explains the mismatch negativity best?

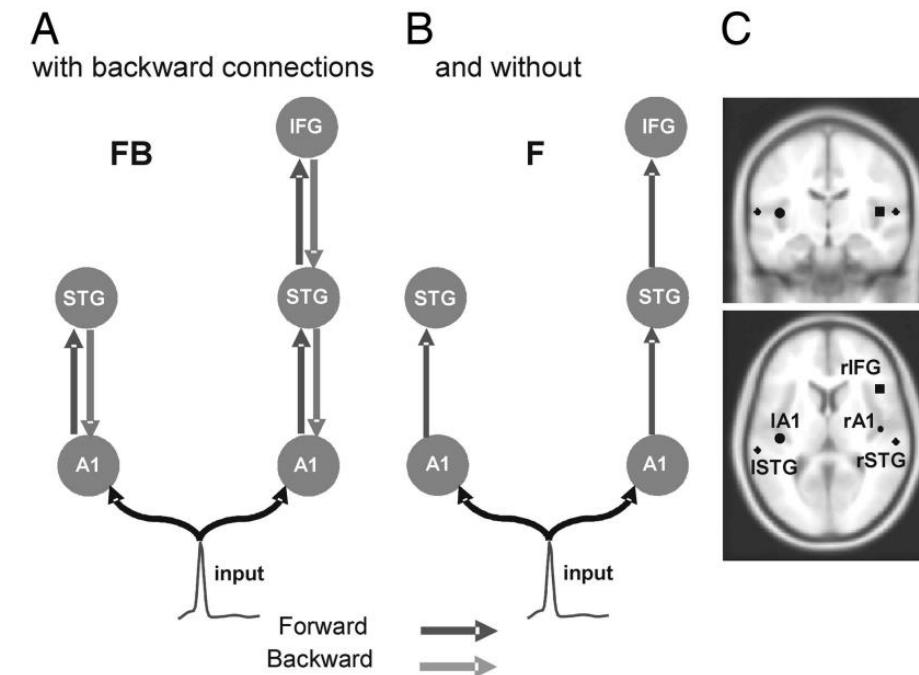
- Model space: Networks with different regions



# Example 2

Are backward connections required to explain the mismatch negativity?

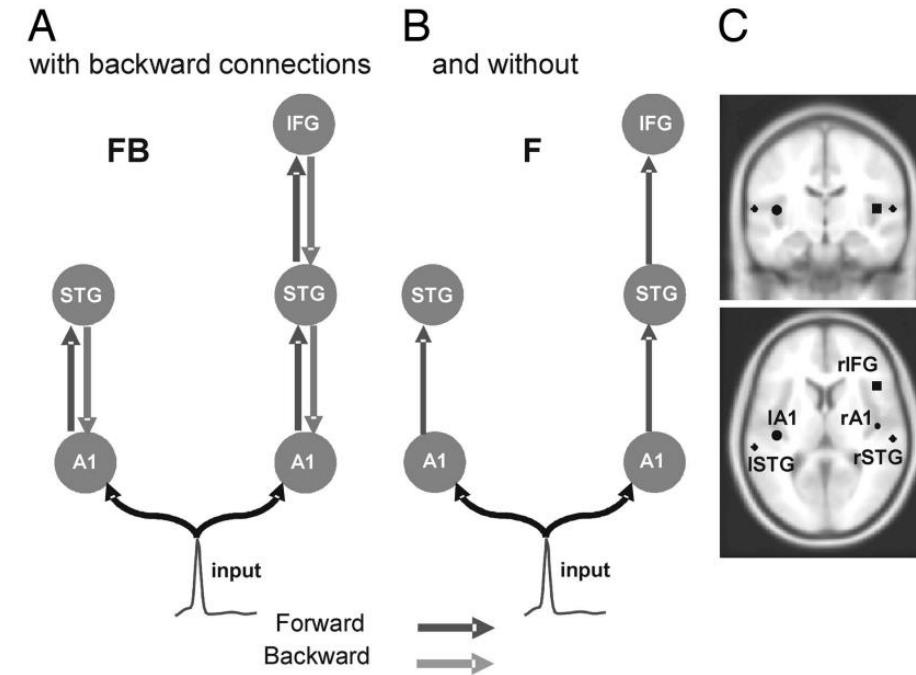
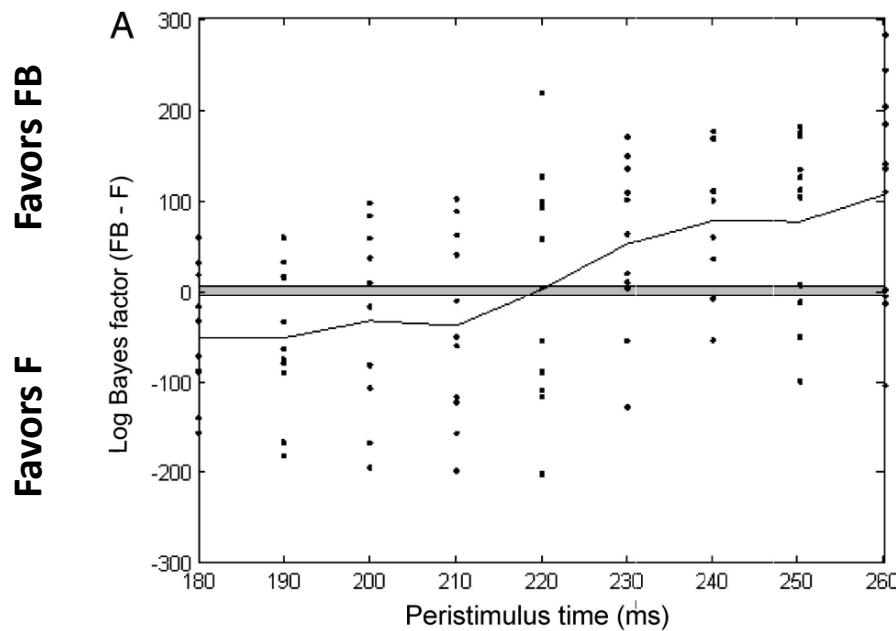
- Model space: Models with and without backward connections



# Example 2

Are backward connections required to explain the mismatch negativity?

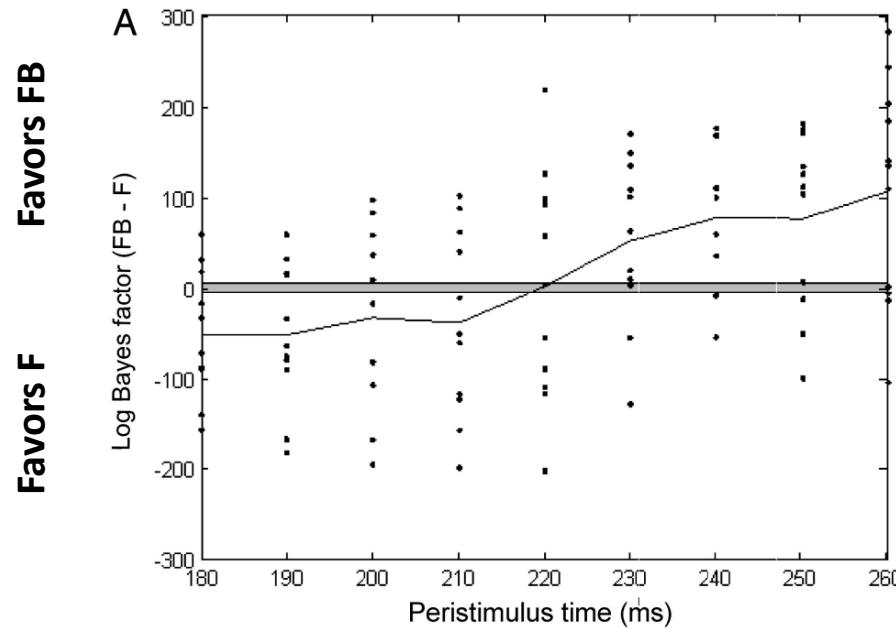
- Model space: Models with and without backward connections



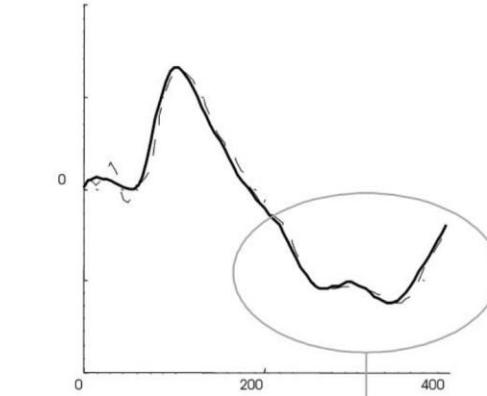
# Example 2

Are backward connections required to explain the mismatch negativity?

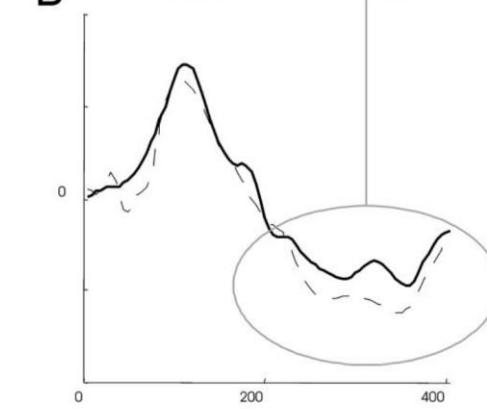
- Model space: Models with and without backward connections



A Predicted and observed response with backward connections



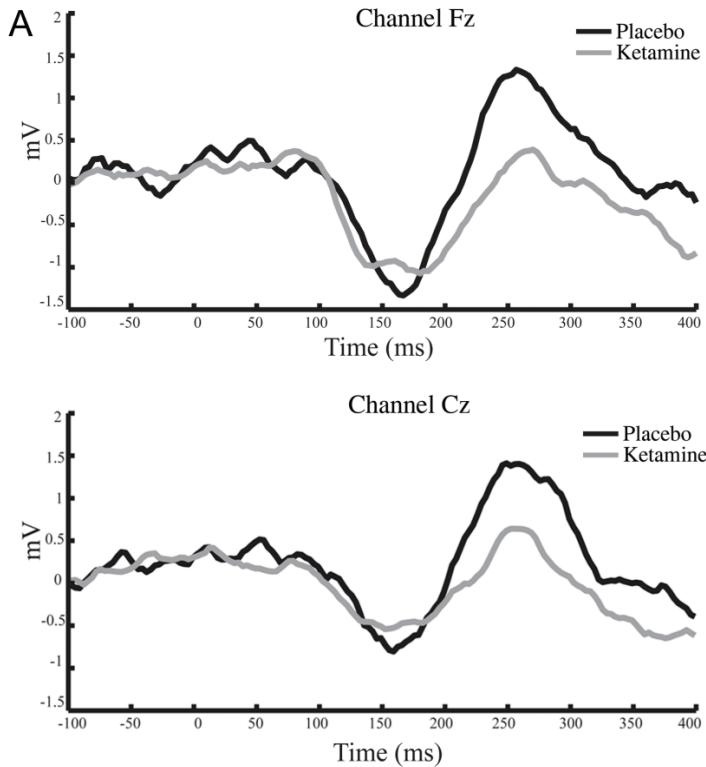
B and without backward connections



# Example 3

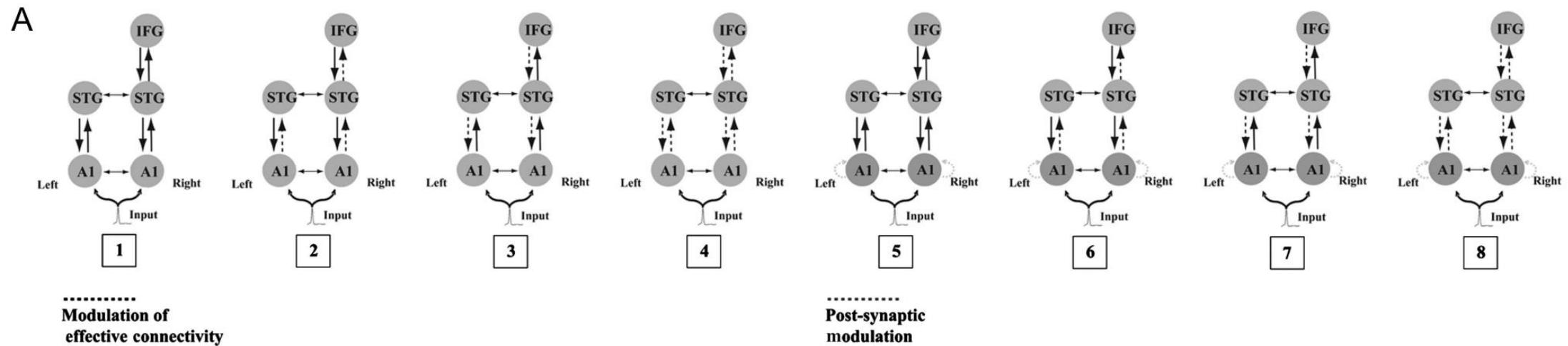
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How does ketamine affect connectivity during the mismatch negativity?



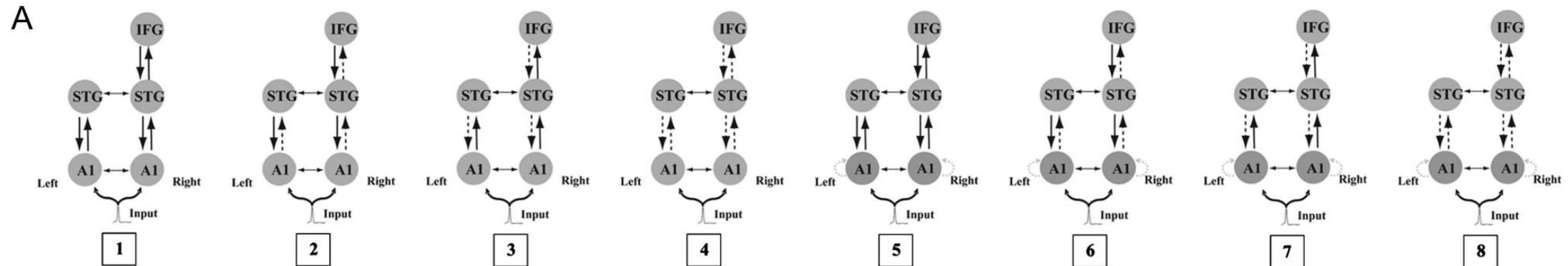
# Example 3

How does ketamine affect connectivity during the mismatch negativity?



# Example 3

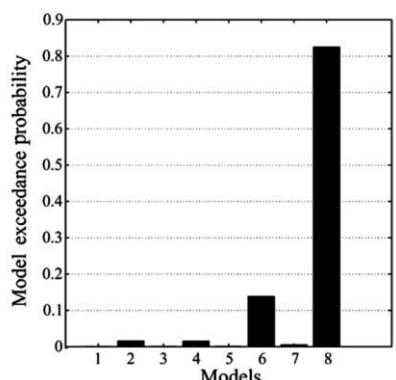
How does ketamine affect connectivity during the mismatch negativity?



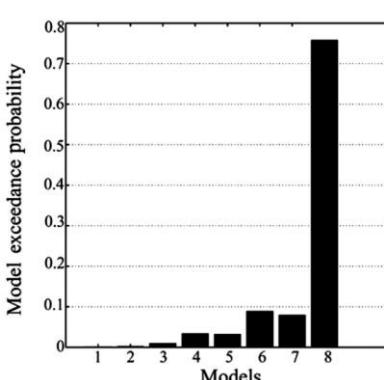
Modulation of effective connectivity

Post-synaptic modulation

B Placebo: population-level best model

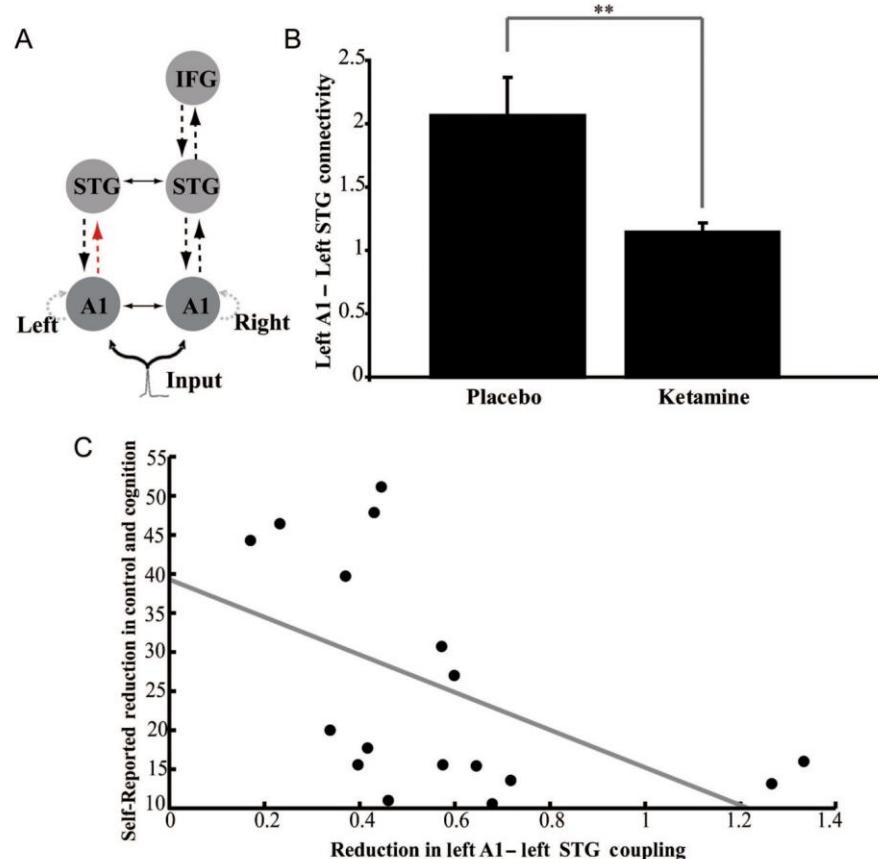


C Ketamine: population-level best model



# Example 3

How does ketamine affect connectivity during the mismatch negativity?



# Example 4

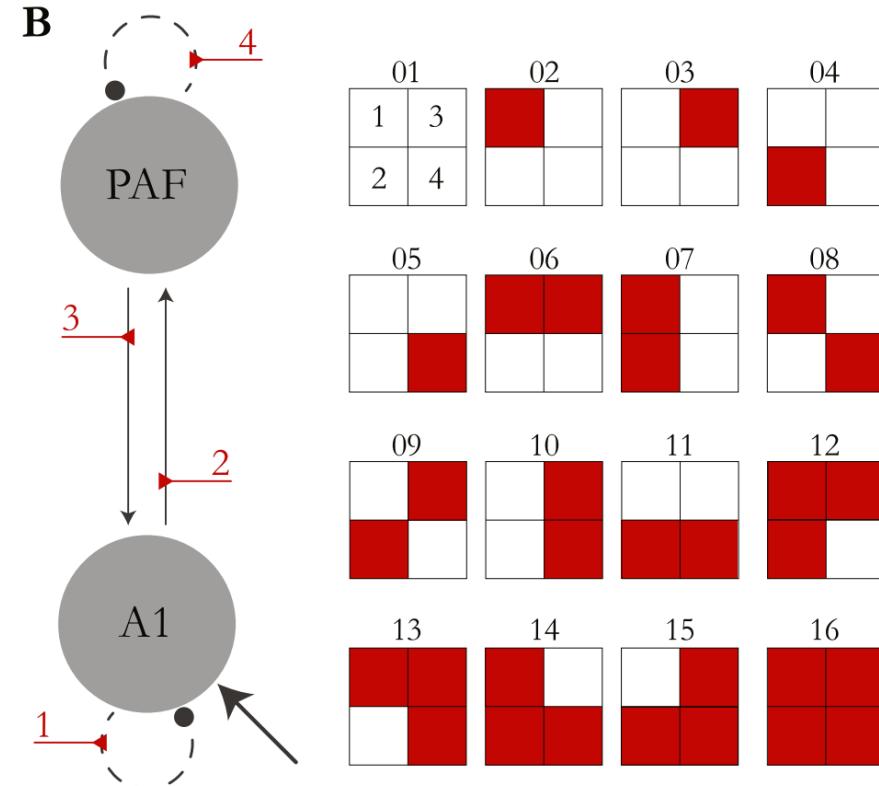
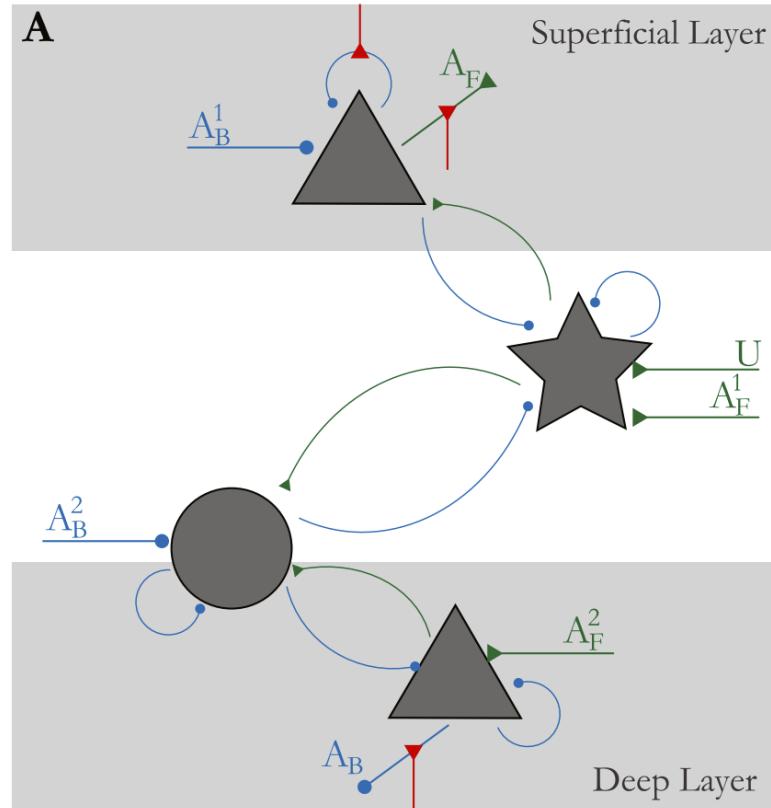
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**What is the role of muscarinic receptor function for MMN generation?**

- Oddball task in rats
- Pharmacological intervention: muscarine receptor agonist scopolamine and antagonist pilocarpine
- Intracranial recordings
- DCM: Canonical microcircuit/LFP

# Example 4

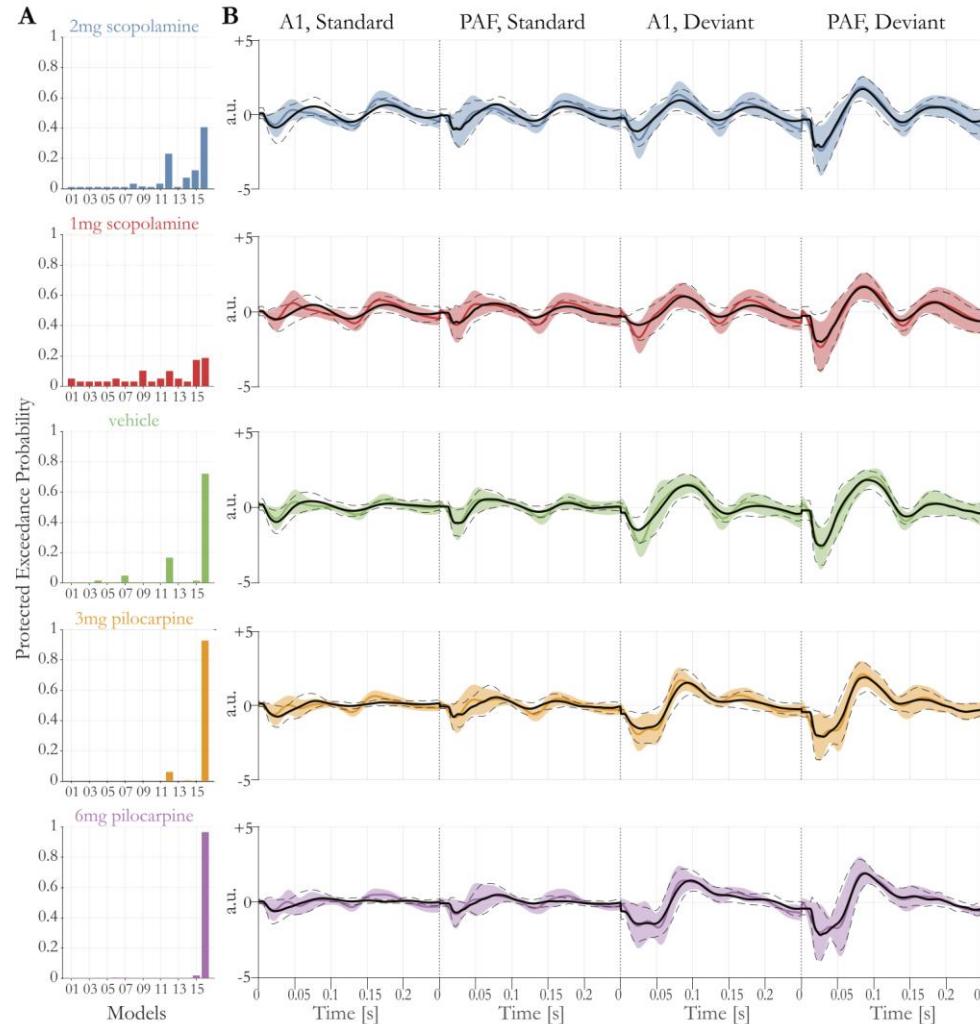
What is the role of muscarinic receptor function for MMN generation?



# Example 4

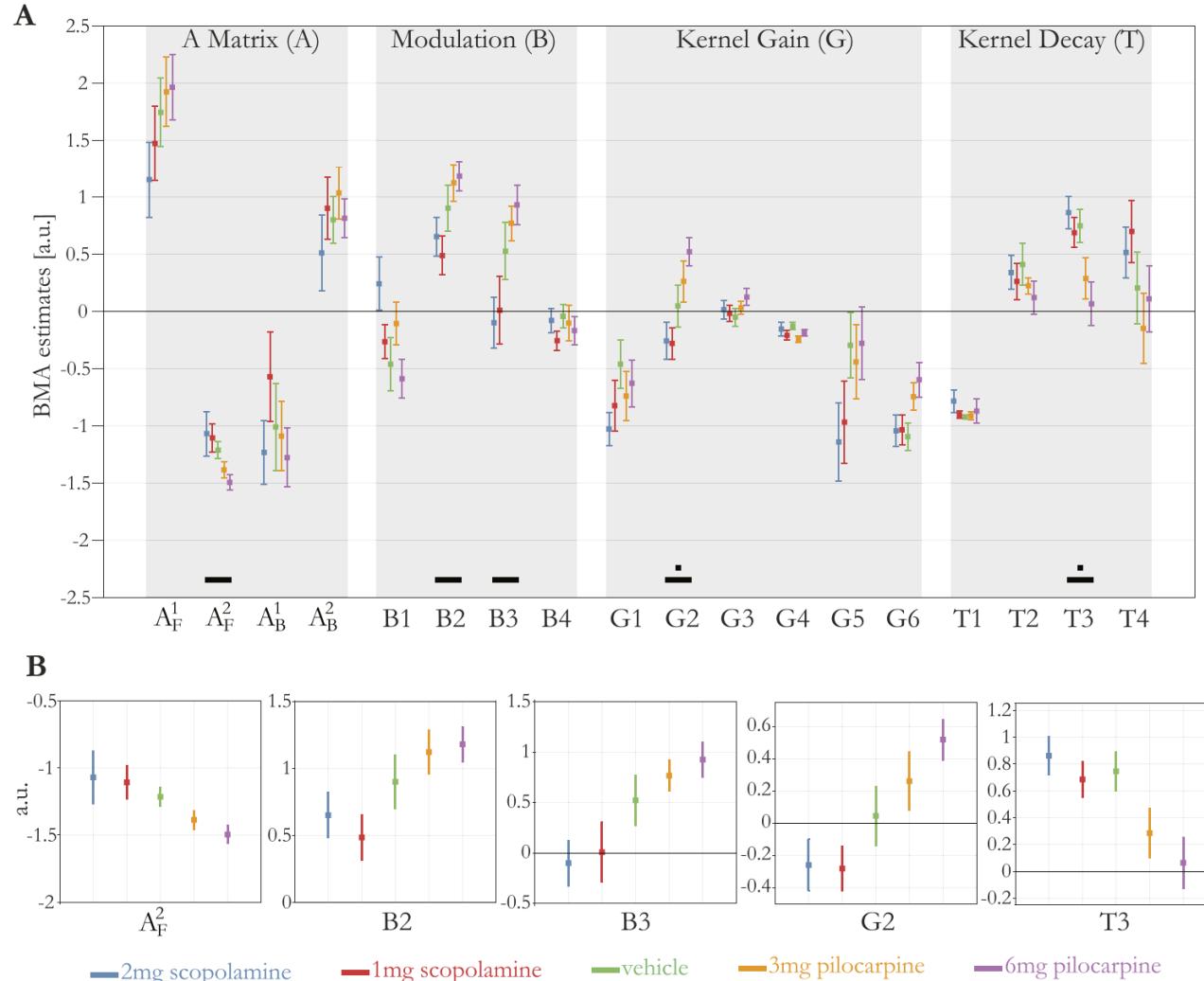
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What is the role of muscarinic receptor function for MMN generation?



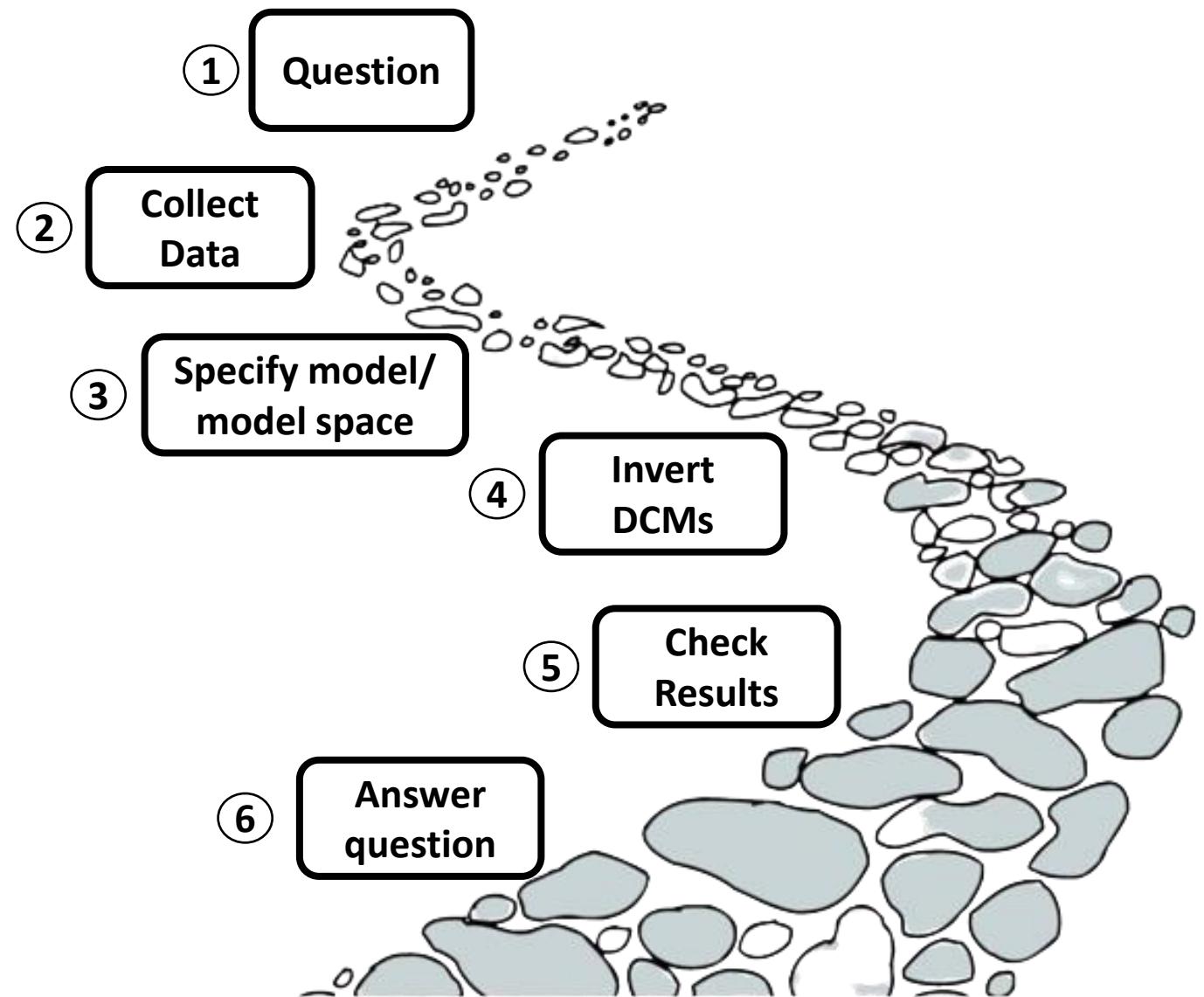
# Example 4

What is the role of muscarinic receptor function for MMN generation?



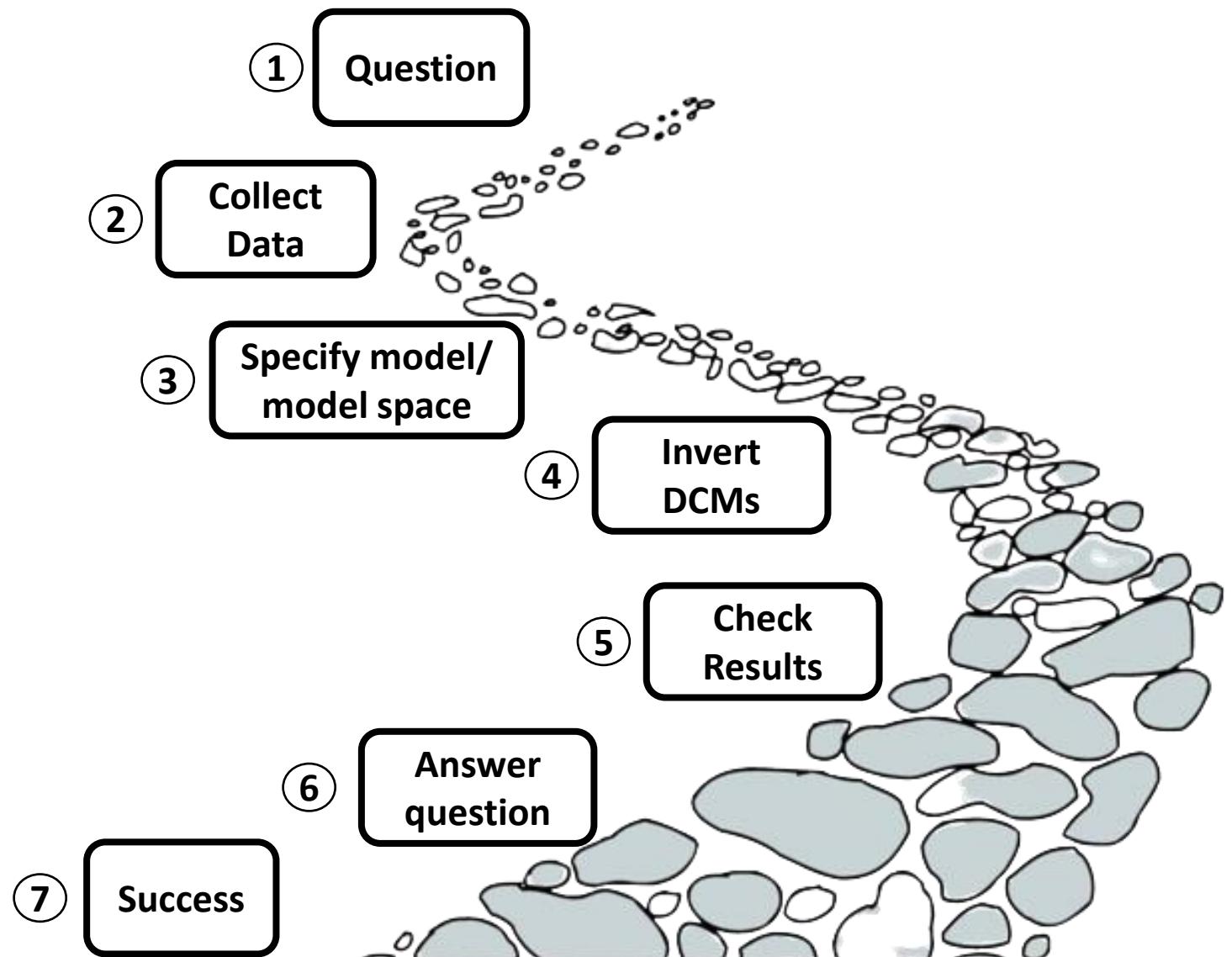
# The DCM analysis path

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# The DCM analysis path

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# Resources

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## Overview Papers

- Moran, Pintosis, & Friston (2013), Neural masses and fields in dynamic causal modeling, *Front. Comput. Neurosci.*
- Pereira et al. (2021), Conductance-based dynamic causal modeling: A mathematical review of its application to cross-power spectral densities, *NeuroImage*

## Books

- SPM manual: <https://www.fil.ion.ucl.ac.uk/spm/doc/manual.pdf>

## Videos

- Previous SPM courses: <https://www.fil.ion.ucl.ac.uk/spm/course/>
- Zurich CPC courses: <https://www.tnu.ethz.ch/de/teaching/cpcourse>
- KCNI Summer School: <https://www.crowdcast.io/e/kcni-summer-school-2021>

## In-depth reading

- Garrido, M.I., Kilner, J.M., Kiebel, S.J. and Friston, K.J., 2007. Evoked brain responses are generated by feedback loops. *Proceedings of the National Academy of Sciences*, 104(52), pp.20961-20966.
- David, O., Harrison, L. and Friston, K.J., 2005. Modelling event-related responses in the brain. *NeuroImage*, 25(3), pp.756-770.
- Hodgkin, A.L. and Huxley, A.F., 1952. A quantitative description of membrane current and its application to conduction and excitation in nerve. *The Journal of physiology*
- Jansen BH, Rit VG (1995) Electroencephalogram and visual evoked potential generation in a mathematical model of coupled cortical columns. *Biol Cybern*
- Marreiros, A.C., Kiebel, S.J., Daunizeau, J., Harrison, L.M. and Friston, K.J., 2009. Population dynamics under the Laplace assumption. *Neuroimage*
- Litvak, V., Jafarian, A., Zeidman, P., Tibon, R., Henson, R.N. and Friston, K., 2019, October. There's no such thing as a 'true' model: the challenge of assessing face validity. In 2019 IEEE
- Zeidman P, Friston K, Parr T (2022) A primer on Variational Laplace
- Pinotsis, D.A., Leite, M. and Friston, K.J., 2013. On conductance-based neural field models. *Frontiers in computational neuroscience*
- Pinotsis, D.A. and Friston, K.J., 2014. Neural fields, masses and Bayesian modelling. In *Neural fields* (pp. 433-455). Springer, Berlin, Heidelberg.
- Marreiros, A.C., Pinotsis, D.A., Brown, P. and Friston, K.J., 2015. DCM, conductance based models and clinical applications. *Validating Neuro-Computational Models of Neurological and Psychiatric Disorders*, pp.43-70.  
[https://www.researchgate.net/publication/274078765\\_DCM\\_Conductance\\_Based\\_Models\\_and\\_Clinical\\_Applications](https://www.researchgate.net/publication/274078765_DCM_Conductance_Based_Models_and_Clinical_Applications)

# Acknowledgments

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- Rick Adams
- Julia Rodriguez-Sanchez
- Hope Oloye
- Ingrid Martin
- Victorita Neacsu
- Lioba Berndt
- Vladimir Litvak
- Dimitris Pinotsis

# Thank you for your attention!

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# Questions & Discussion

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