

CPC 2022: Introduction to Computational Psychiatry

Klaas Enno Stephan



Translational Neuromodeling Unit

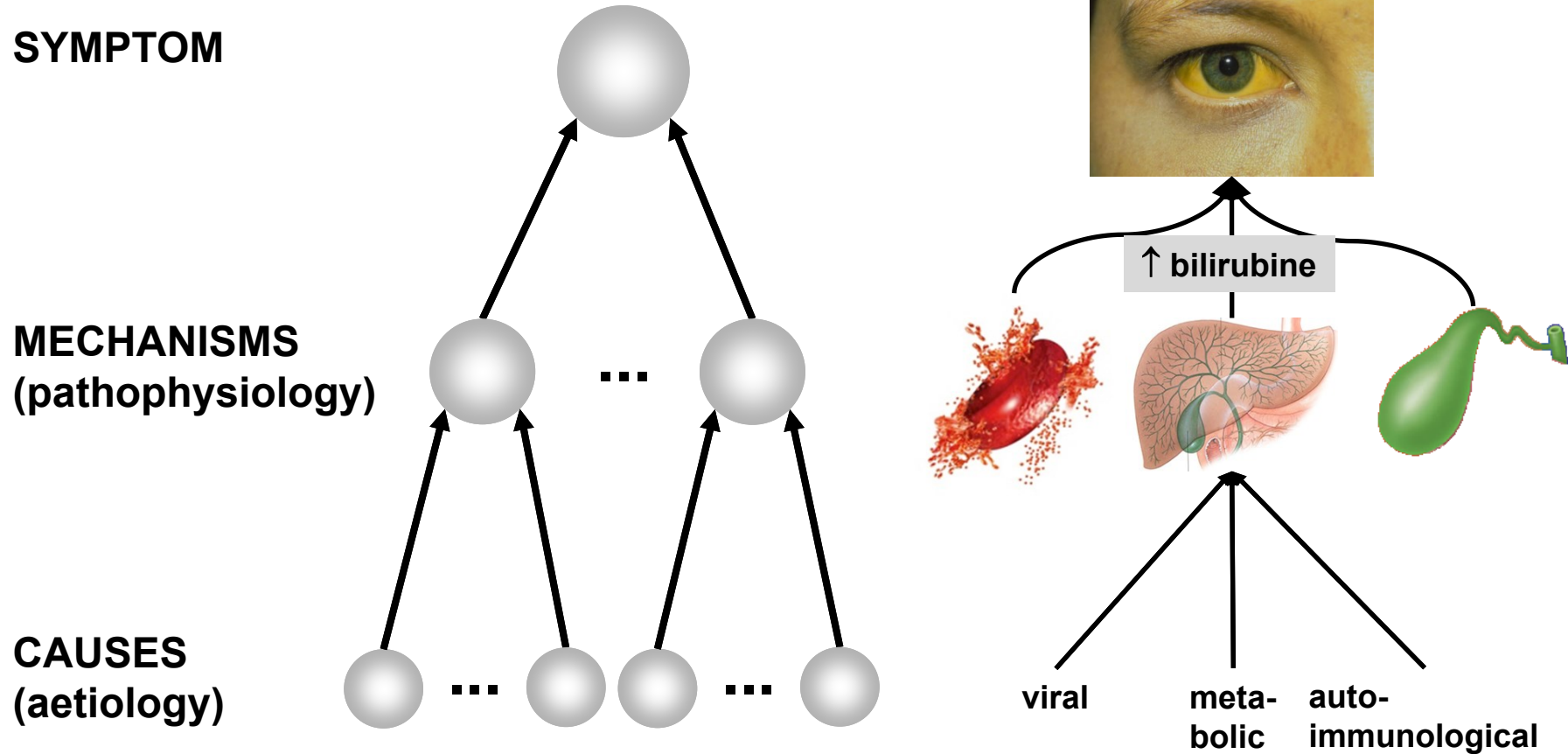


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Zürich^{UZH}

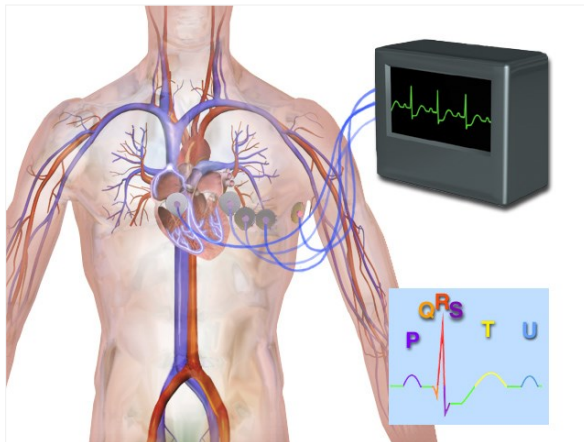
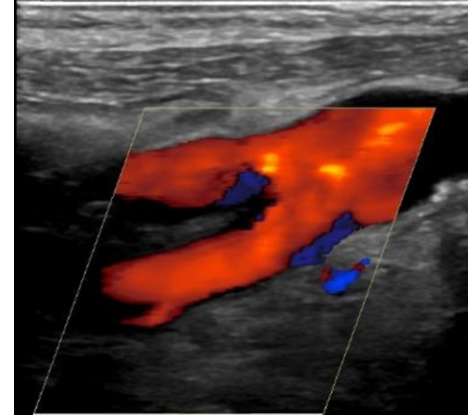
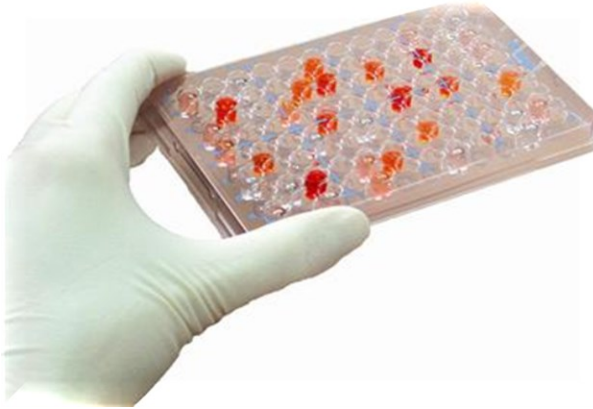


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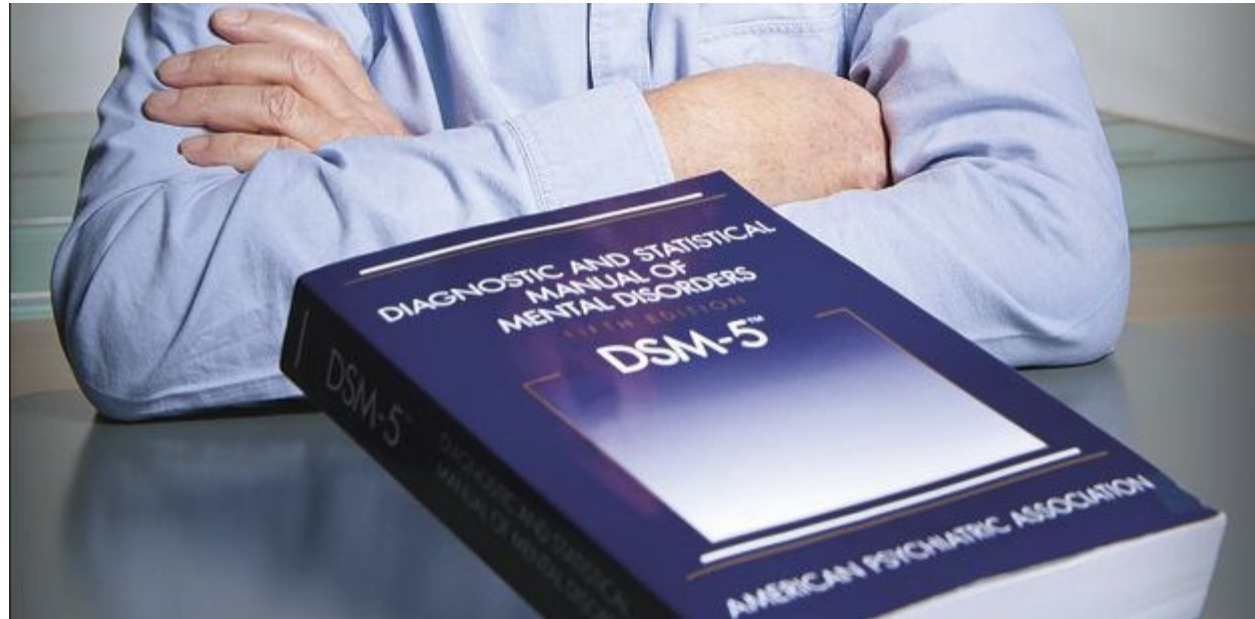
From differential diagnosis to nosology



>3,000 clinical tests in medicine

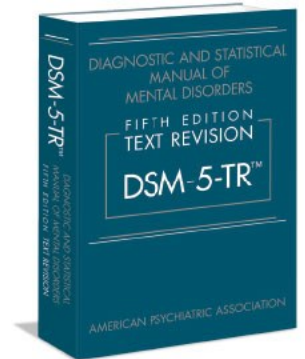
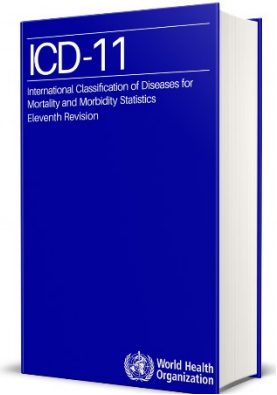


1 diagnostic instrument in psychiatry



Contemporary psychiatric classifications: ICD and DSM

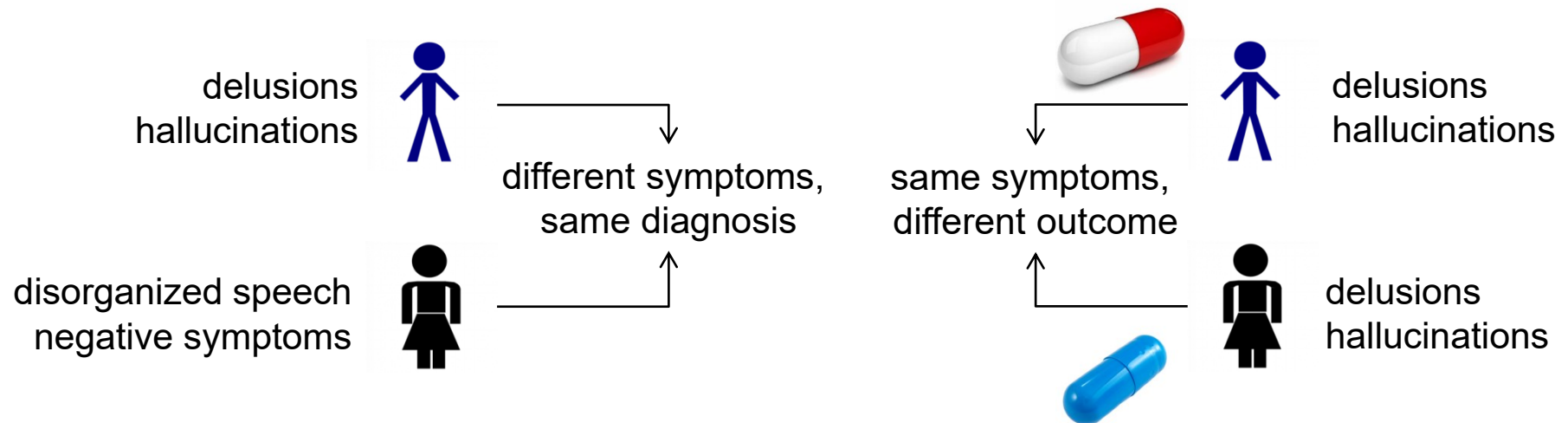
- **International Classification of Diseases (ICD):**
 - curated by the World Health Organization (WHO)
 - freely available
 - presently in its 10th revision (ICD-10); ICD-11 will come into effect in 2022
- **Diagnostic and Statistical Manual of Mental Disorders (DSM)**
 - published by the American Psychiatric Association (APA)
 - not free
 - presently: fifth edition (DSM-5); text revision (TR) published in 2022
- **both schemes**
 - define mental disorders as syndromes
 - reflect the consensus (or compromise) of expert committees
 - are descriptive (without reference to mechanisms)



DSM-5: Schizophrenia

- Positive symptoms:
 - Delusions
 - Hallucinations
 - Disorganized speech
 - Grossly disorganized or catatonic behavior
 - Negative symptoms (e.g., flat affect, anhedonia, avolition, asociality)
- + social or occupational dysfunction
+ continuous signs of the disturbance for at least six months

≥ 2 symptoms
(at least one pos. symptom)
over ≥ 1 month



Heterogeneity of psychiatric disorders




polygenetic basis
gene-environment interactions
environmental variation

**variability in clinical
trajectory and treatment
response**

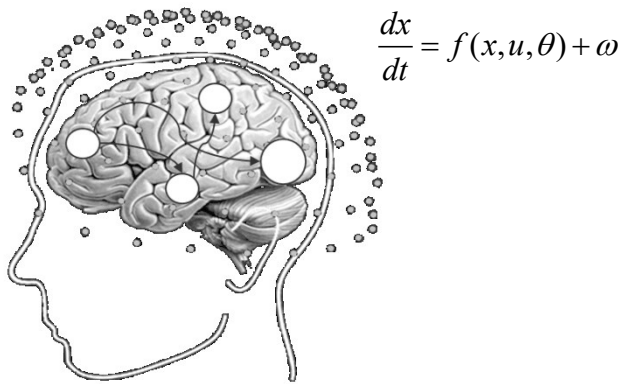
multiple disease mechanisms

PERSPECTIVE

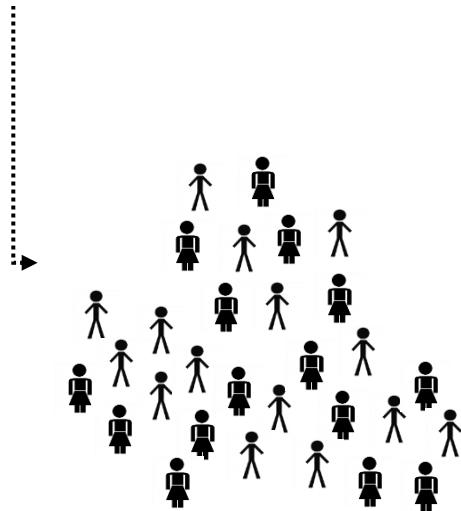
Why has it taken so long for biological psychiatry to develop clinical tests and 

S Kapur¹, AG Phillips² and TR Insel³

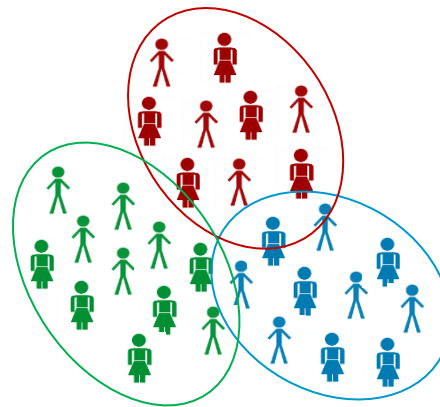
1 Developing computational assays of neuronal and cognitive processes



2 Application to brain activity and behaviour of individual patients



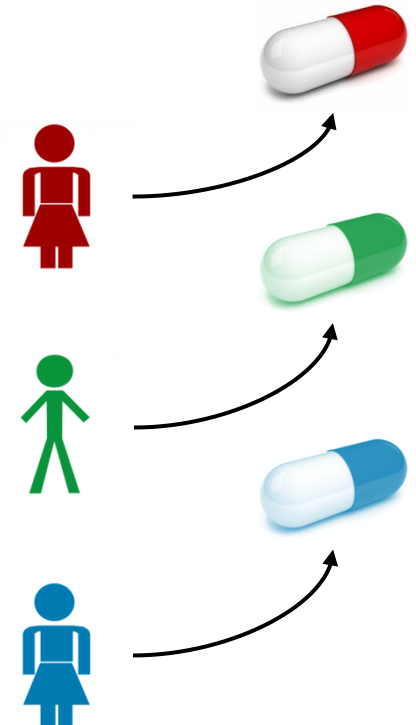
3 Differentiating patients based on inferred mechanisms



- disease mechanism A
- disease mechanism B
- disease mechanism C

Translational Neuromodeling & Computational Psychiatry

4 Individual treatment prediction

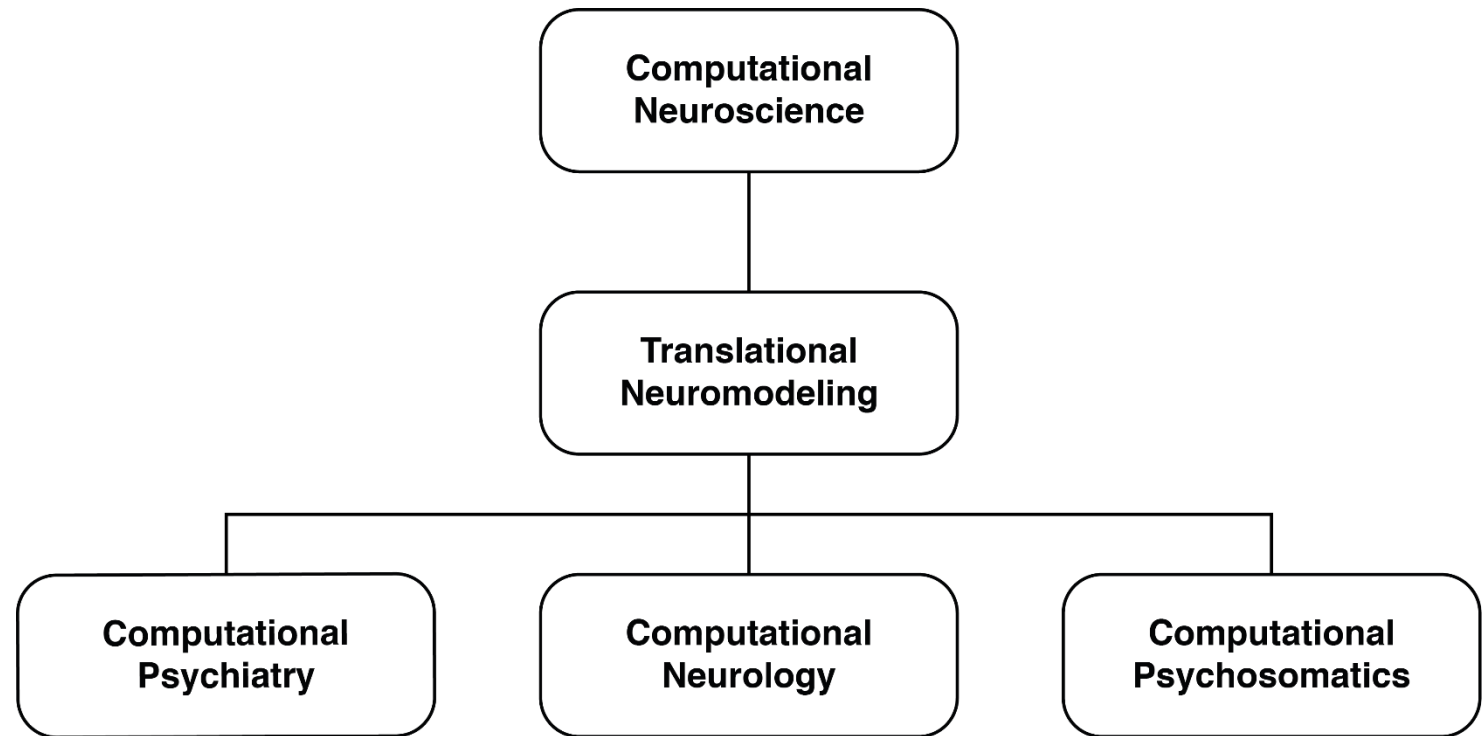


A taxonomy of computational clinical neuroscience

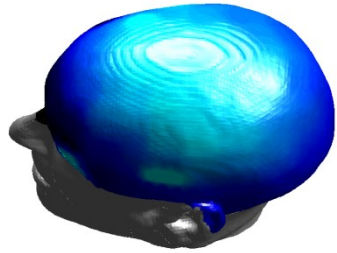
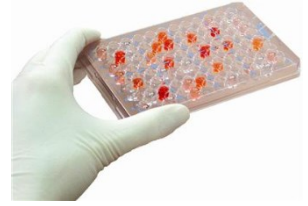
Understanding how/what
the brain computes

Develops/validates
mathematical models for
solving clinical problems

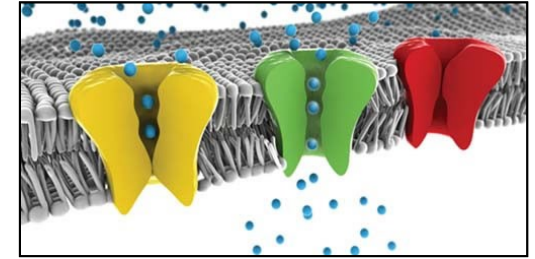
Application within
specific medical fields



Generative models as "computational assays"



$$\begin{array}{c} \xleftarrow{p(y | \theta, m) \cdot p(\theta | m)} \\ \xrightarrow{p(\theta | y, m)} \end{array}$$



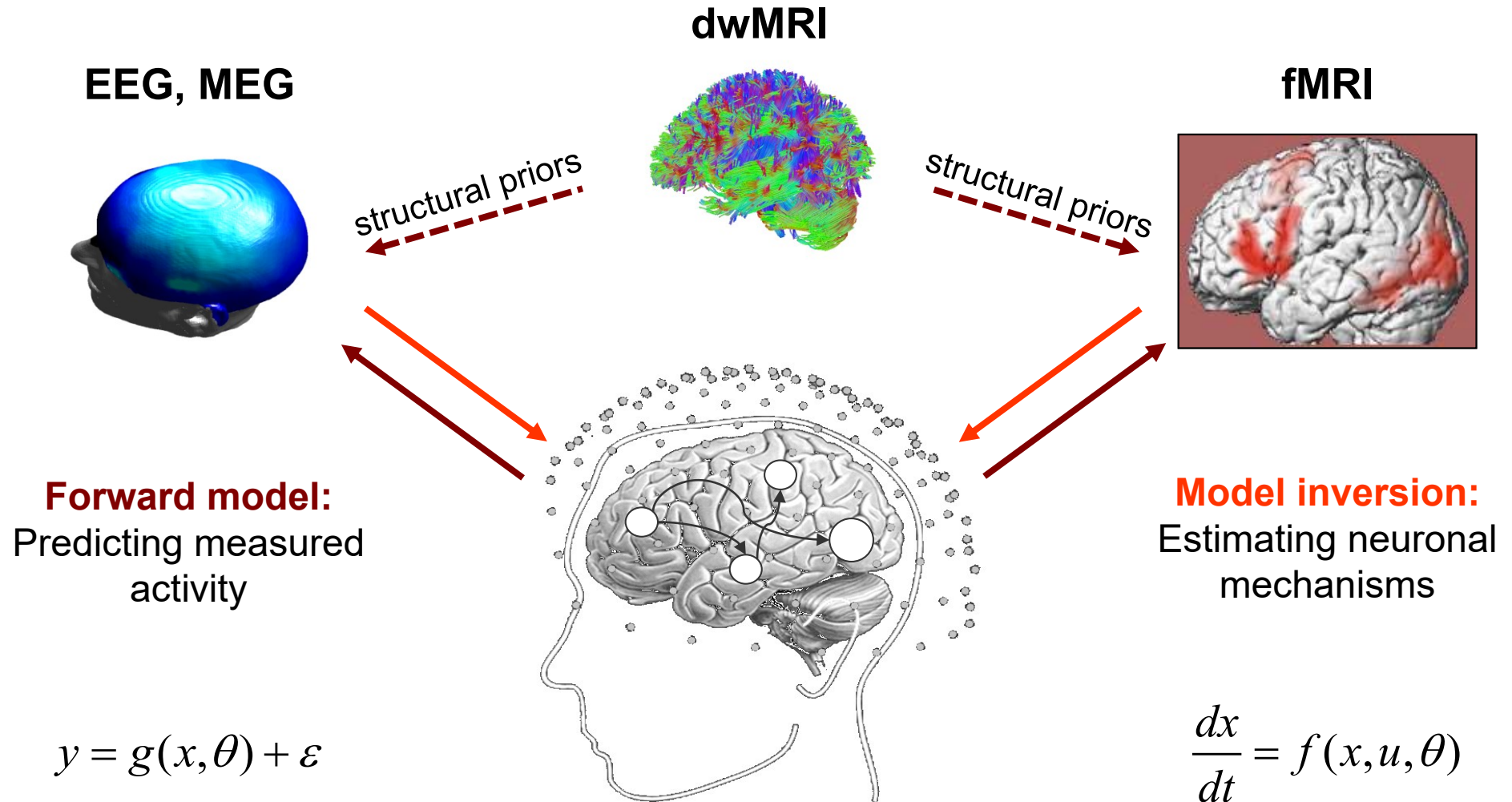
y = data, θ = parameters, m = model



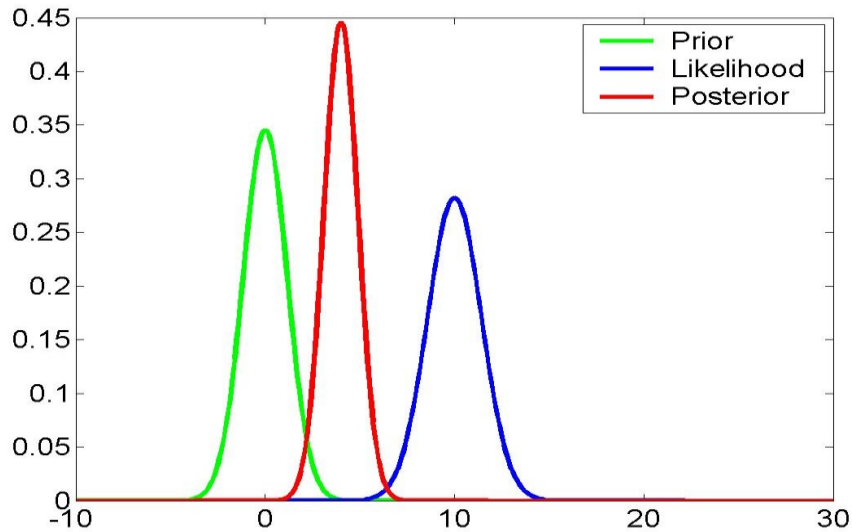
$$\begin{array}{c} \xleftarrow{p(y | \theta, m) \cdot p(\theta | m)} \\ \xrightarrow{p(\theta | y, m)} \end{array}$$



Example: Dynamic causal models (DCMs)

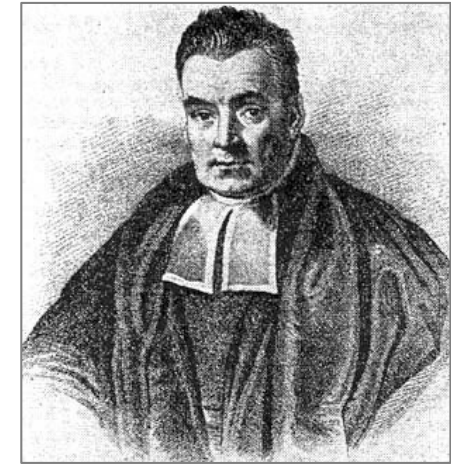


Bayes' rule



$$\text{Posterior (inference)} = \frac{\text{Likelihood (data)} \times \text{Prior (prediction)}}{\text{Evidence (normalisation term)}}$$

θ : parameters
 y : data

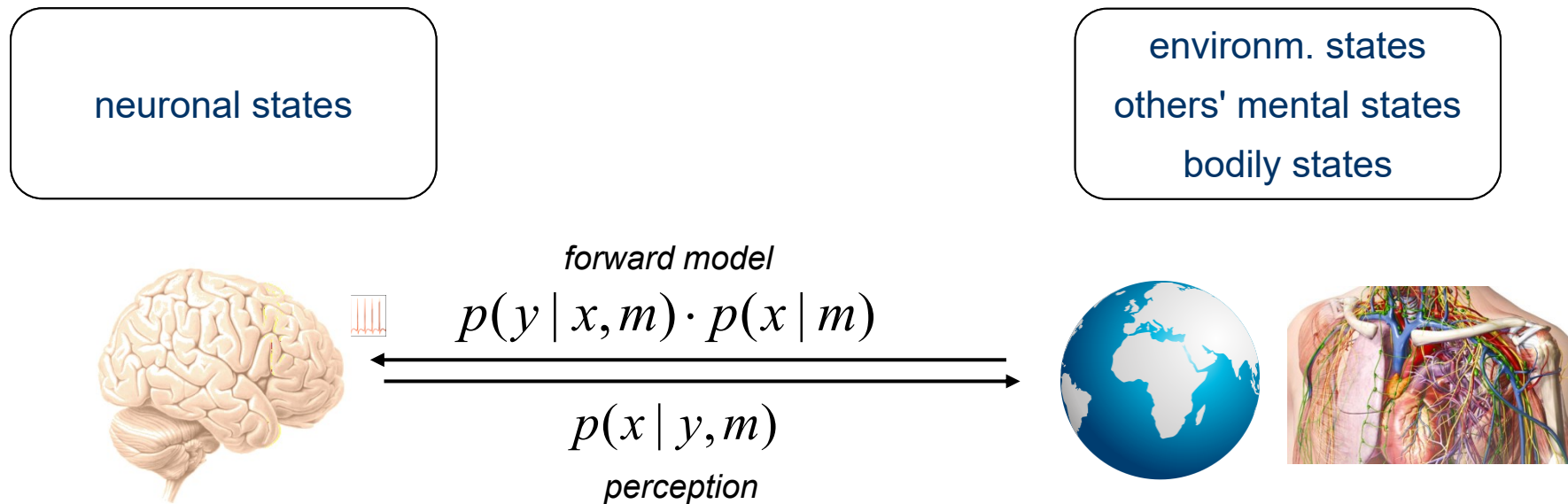


The Reverend Thomas Bayes
(1702-1761)

“... the theorem expresses how a ... degree of belief should rationally change to account for availability of related evidence.”

Wikipedia

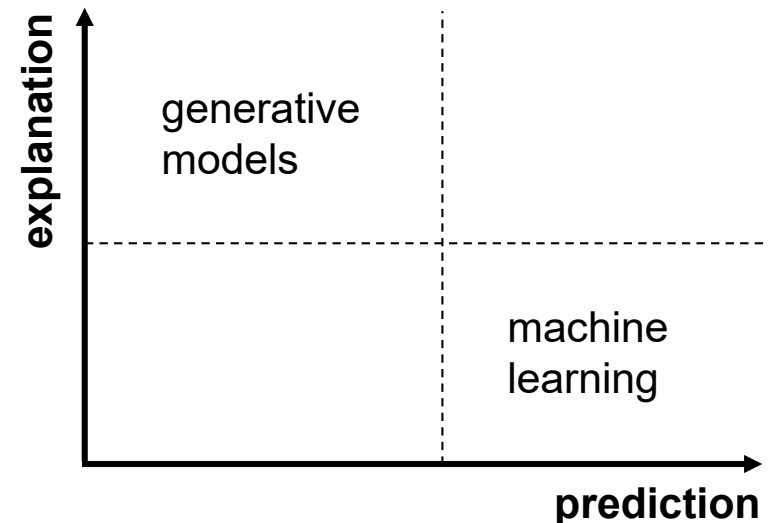
Generative models as a concept for brain function: the "Bayesian brain" hypothesis



perception = inference = inversion of a generative model

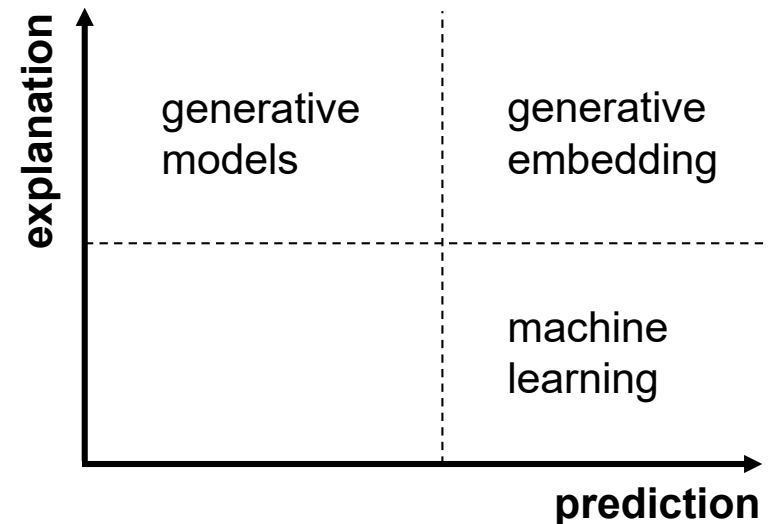
The “Two Cultures of Computational Psychiatry”

- **explanation:** generative models
 - data-generating process is of central interest
 - goal: identify the mechanisms underlying clinical symptoms
- **prediction:** machine learning (ML)
 - data-generating process is treated as a black box
 - goal: prediction of clinically relevant outcomes, e.g. treatment response, remission, relapse



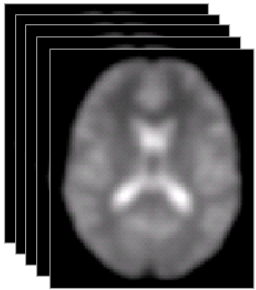
The “Two Cultures of Computational Psychiatry” ... and Generative Embedding as their bridge

- **explanation:** generative models
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- **generative embedding:**
 - applies ML to estimated quantities from generative models

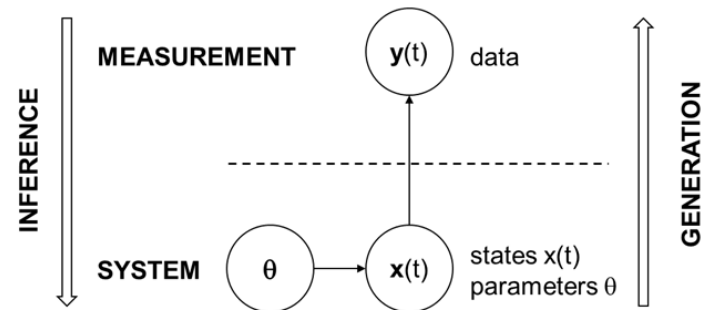


Generative embedding

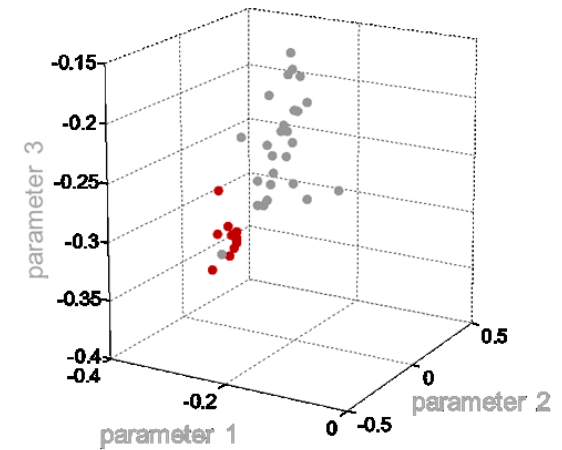
high-dimensional data



generative model



mechanistic interpretation



theory-driven
dimensionality reduction

posterior densities →
features for machine learning

Computational assays: key clinical questions

SYMPTOMS

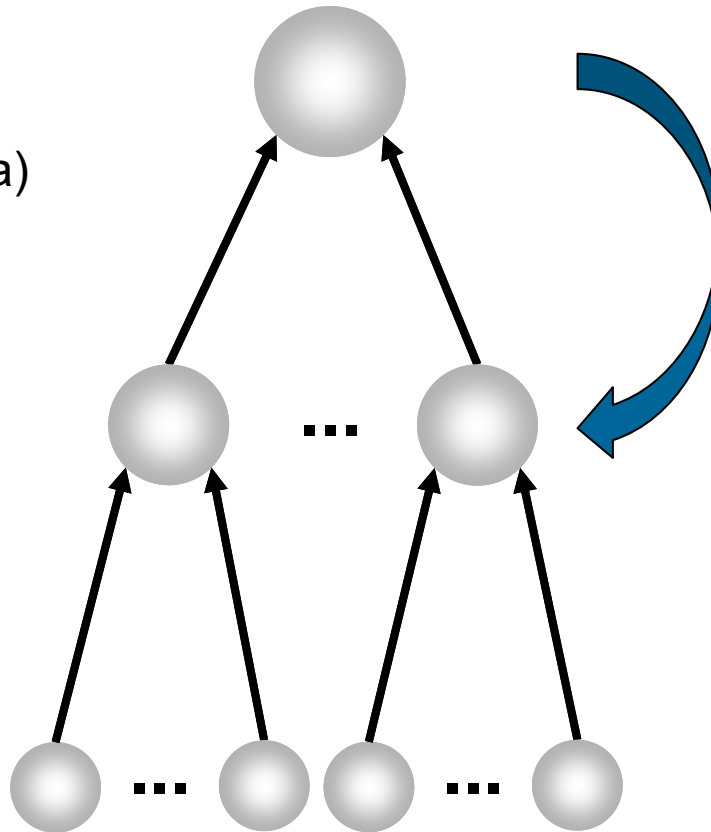
(behavioural or physiological data)

MECHANISMS

(computational, physiological)

CAUSES

(aetiology)

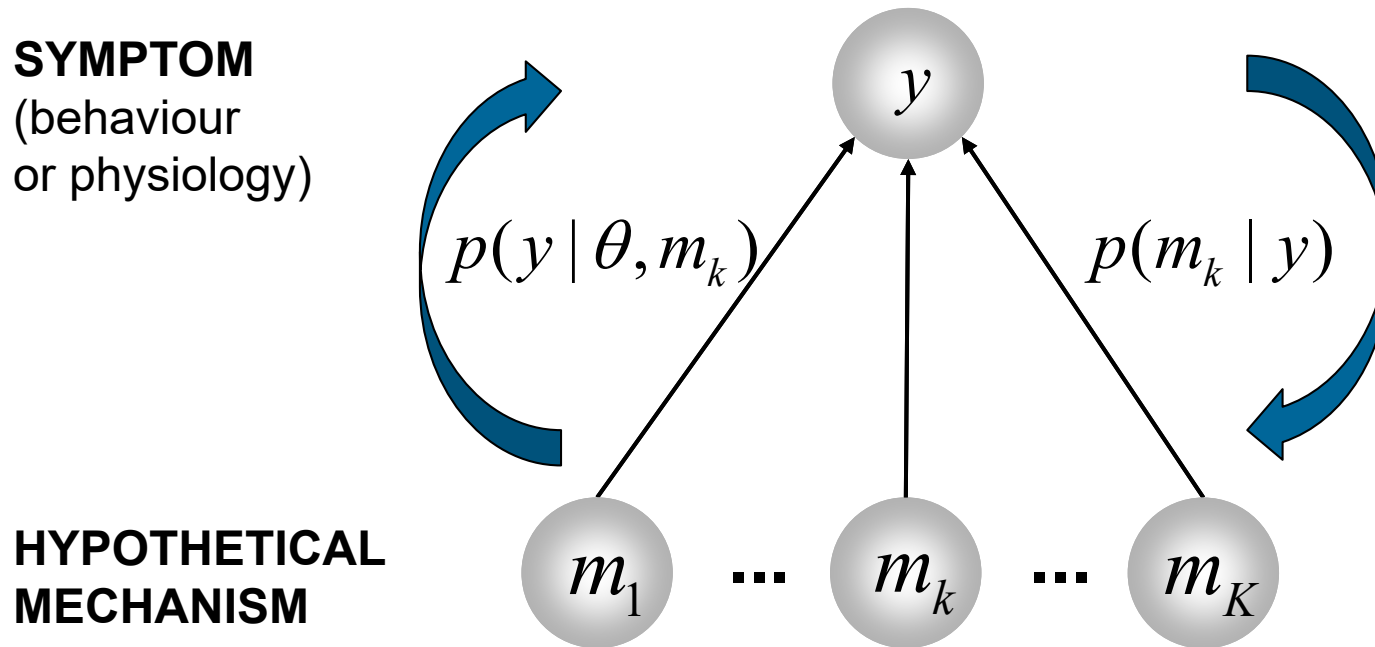


① differential diagnosis of alternative disease mechanisms

② stratification / subgroup detection into mechanistically distinct subgroups

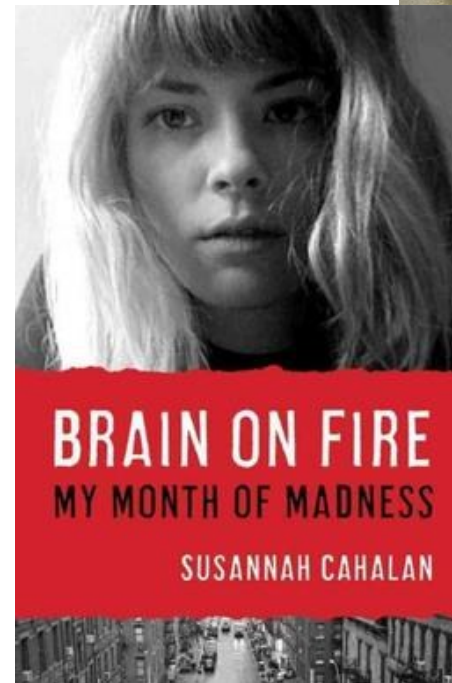
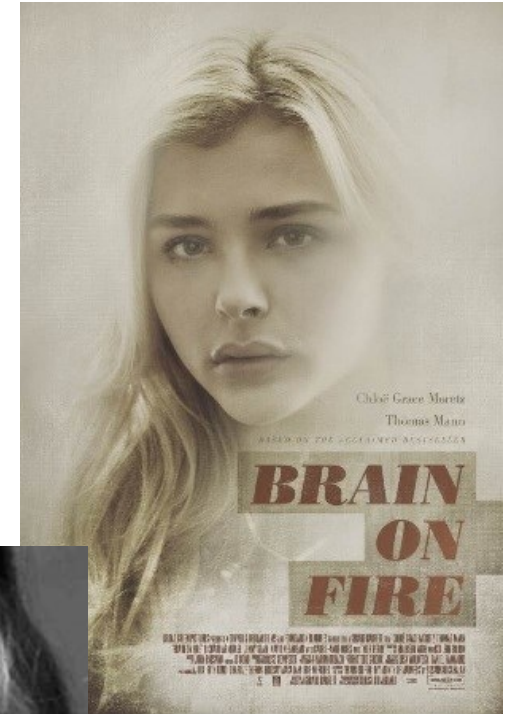
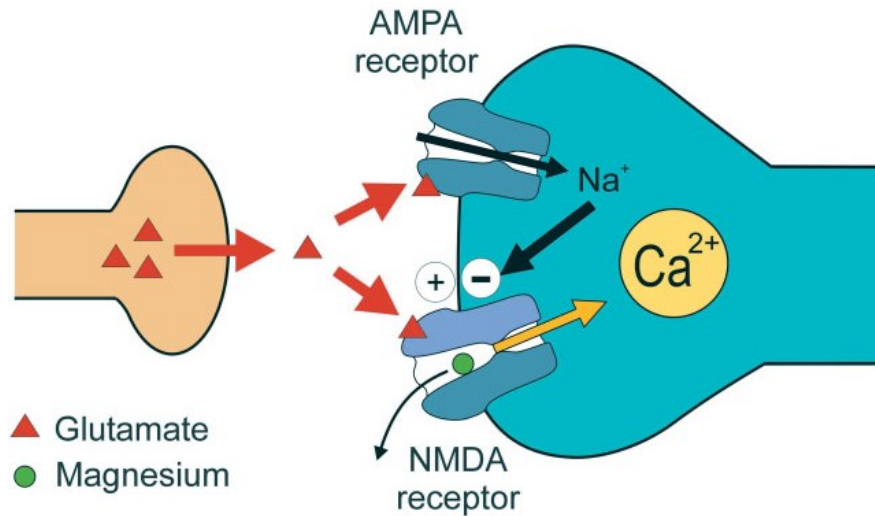
③ prediction of clinical trajectories and treatment response

❶ Differential diagnosis: model selection



$$p(m_k | y) = \frac{p(y | m_k) p(m_k)}{\sum_k p(y | m_k) p(m_k)}$$

NMDA receptor antibody encephalitis



Generative modeling of seizure activity in NMDAR antibody encephalitis

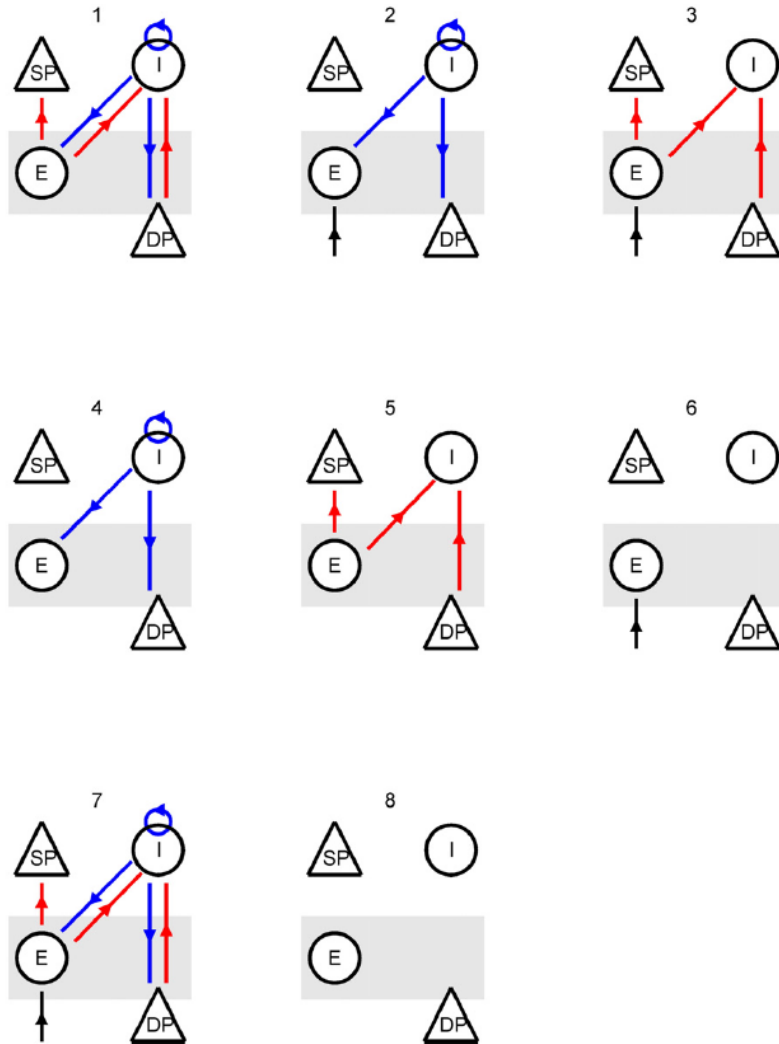


Table 2

The variance described and the free energy for the different models inverted for each patient. Note that the winning model (highest free energy) also had the best fit and these were the same models for both patients. The free energies are expressed relative to the null model.

Model	Patient 1		Patient 2	
	Variance explained	Free energy	Variance explained	Free energy
Inhibitory + excitatory + endogenous	0.97	1430	0.95	1740
Inhibitory + excitatory	0.97	1380	0.94	1650
Inhibitory + endogenous	0.97	1320	0.94	1600
Excitatory + endogenous	0.96	1310	0.94	1680
Inhibitory	0.90	860	0.91	1130
Excitatory	0.91	1010	0.92	1460
Endogenous	0.91	950	0.91	1230
Null	0.50	0	0.75	0

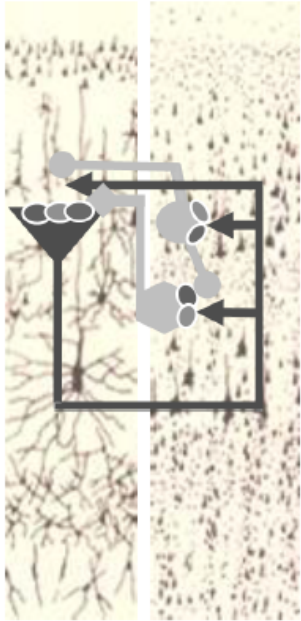
Example of how model selection can serve to infer on pathophysiological processes in single patients.

"Free energy" in table above

= negative free energy

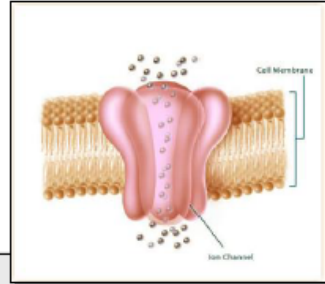
= approximation to log model evidence

① Differential diagnosis: inferring synaptic processes



- inhibitory interneurons
- excitatory interneurons
- pyramidal cells

AMPA, NMDA, GABA_A receptors

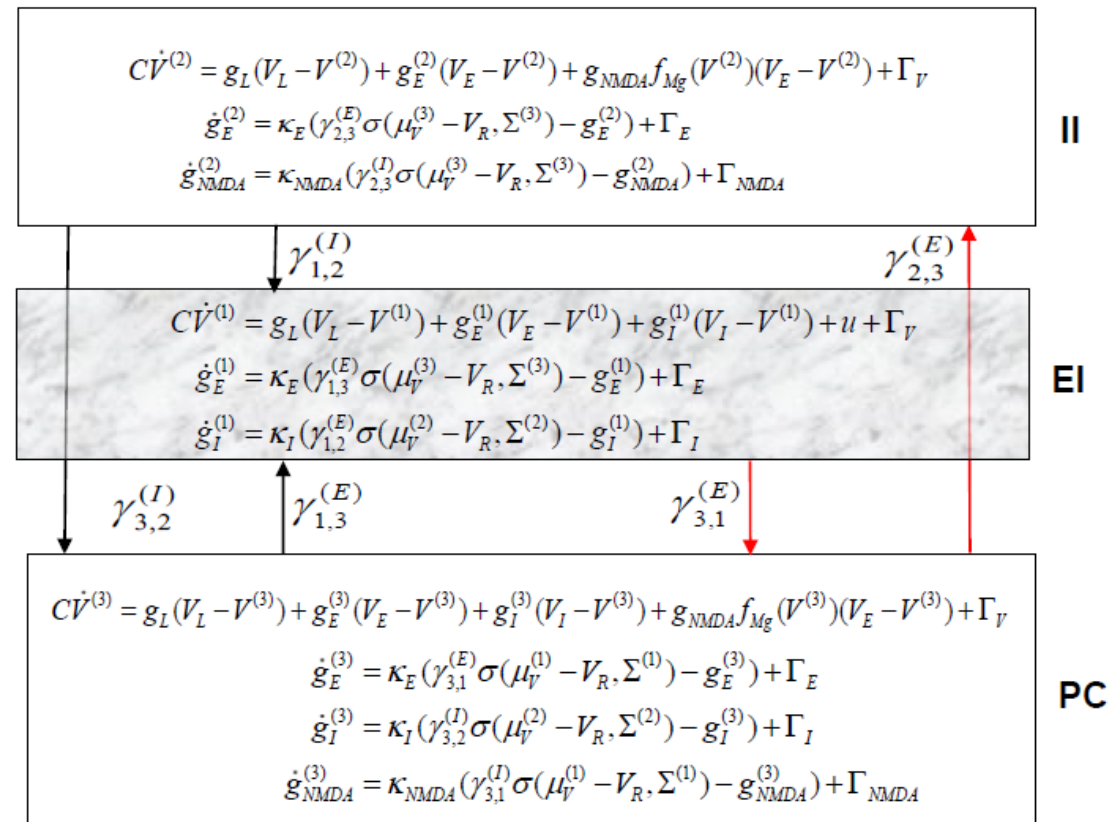


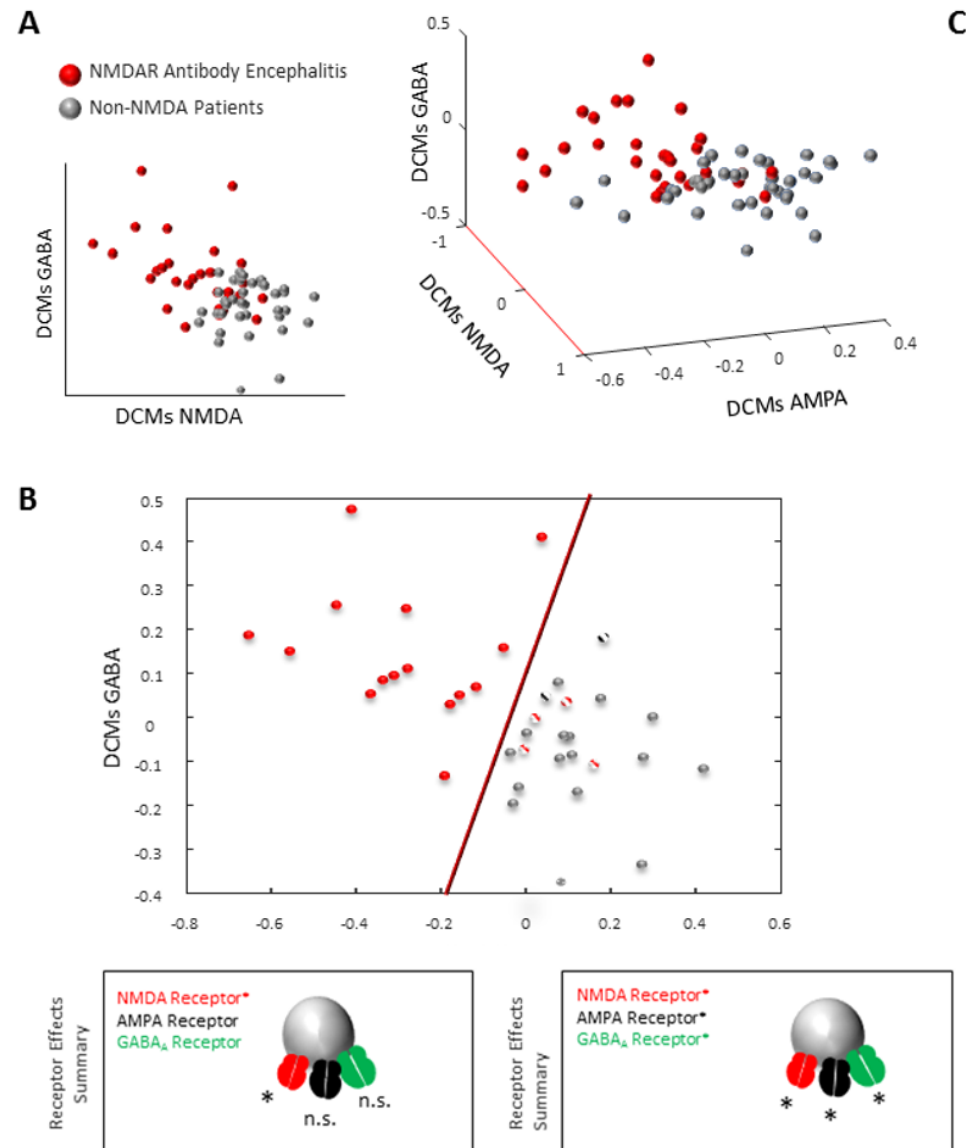
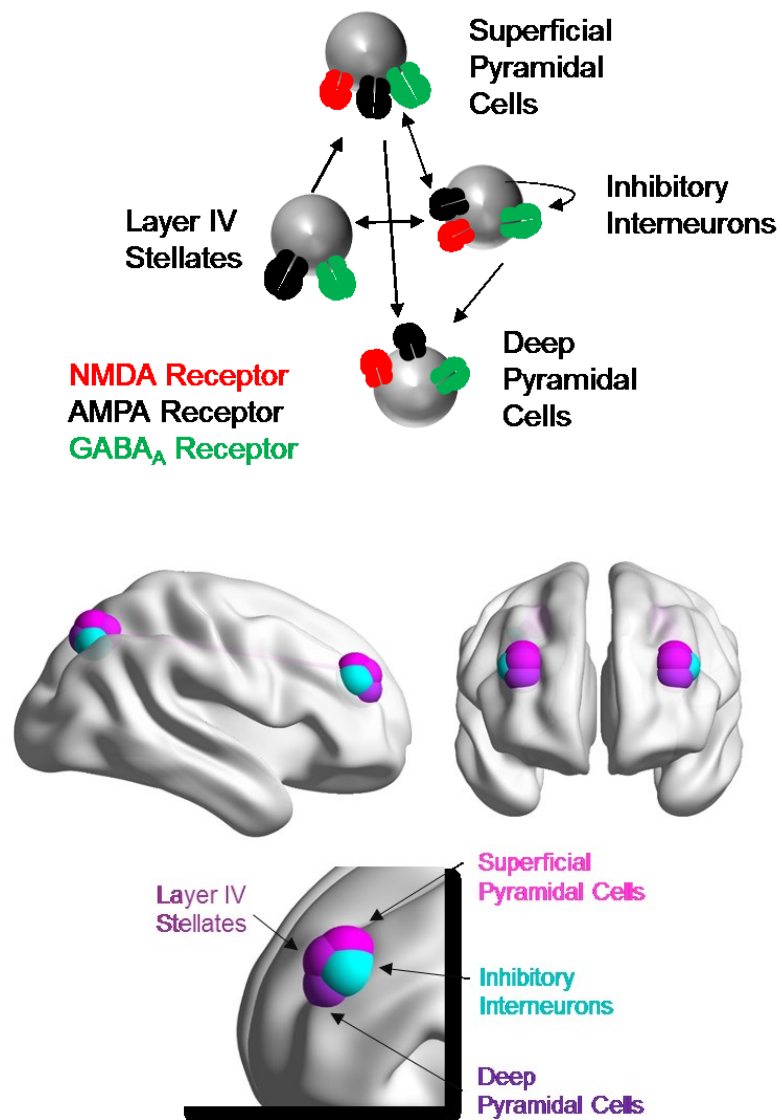
$$C\dot{V} = \sum g_i (V_i^0 - V)$$

$$\dot{g}_k = \kappa (u_{ij} - g_k)$$

$$u_{ij} = \gamma_{ij} \sigma(\mu_V^{(j)} - V_R, \Sigma^{(j)})$$

u_{ij} = presynaptic input from ensemble j to i
 σ = CDF of presynaptic depolarization density around threshold potential V_R





29 patients with NMDAR-antibody encephalitis
 18 control patients (with inflammatory/metabolic encephalopathy)

③ Prediction: Two-year outcome in depression

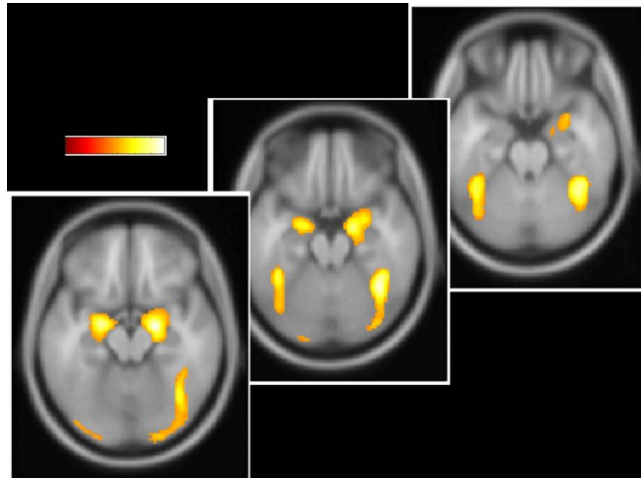
N=85 MDD patients from NESDA study (Schmaal et al. 2015, Biol. Psychiatry)

Three distinct trajectories:

chronic (CHR): n = 15

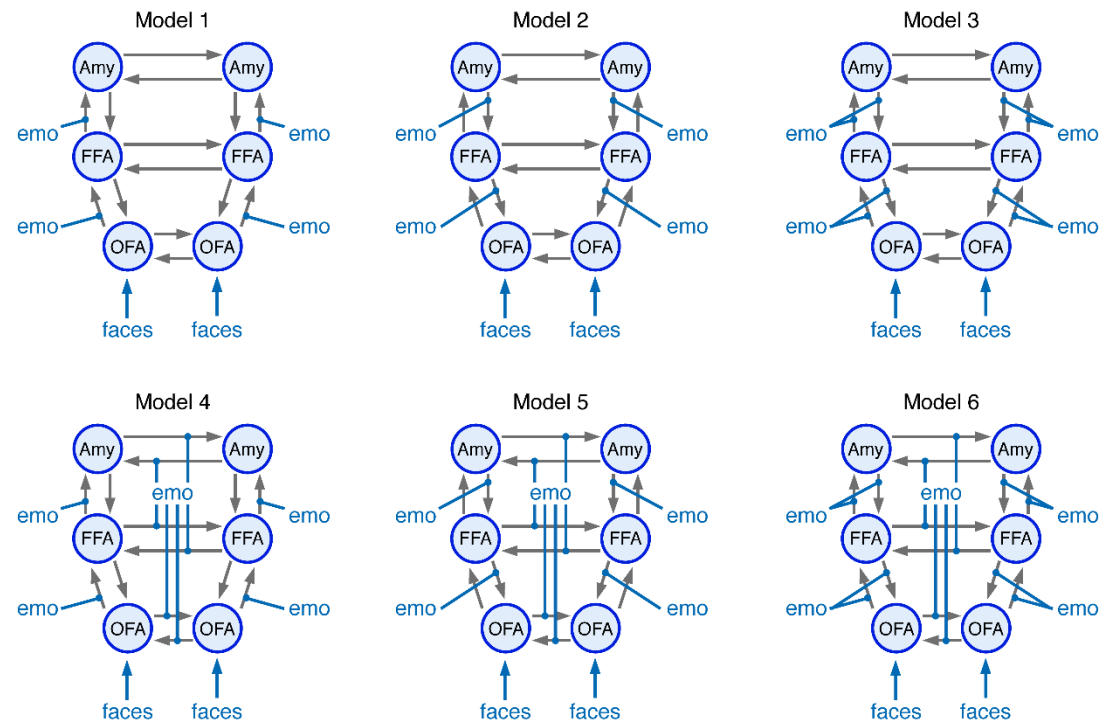
gradually improving (IMP): n = 31

remission (REM): n = 39

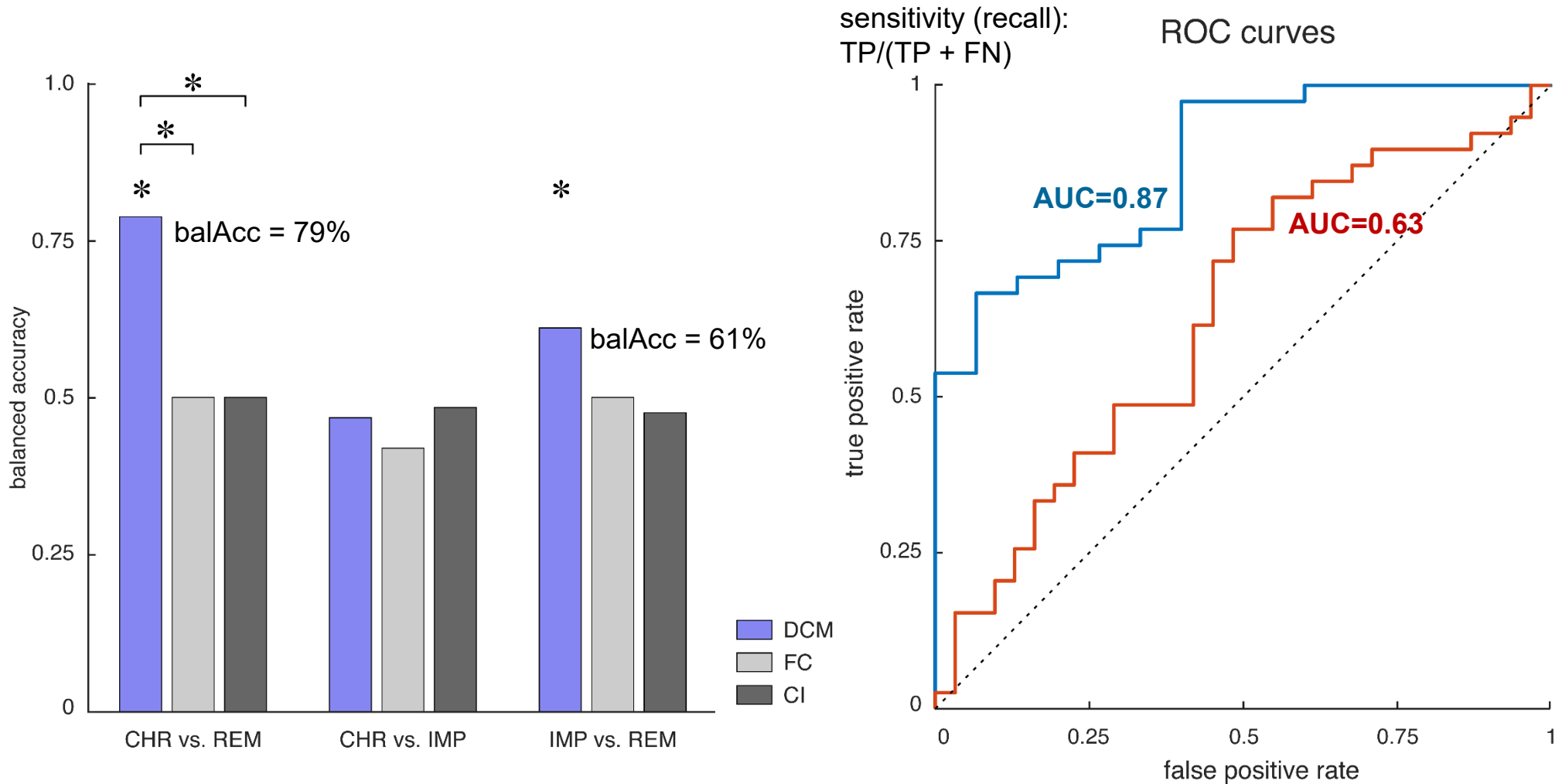


emotional faces > scrambled faces

DCM + BMA (emotional face processing)

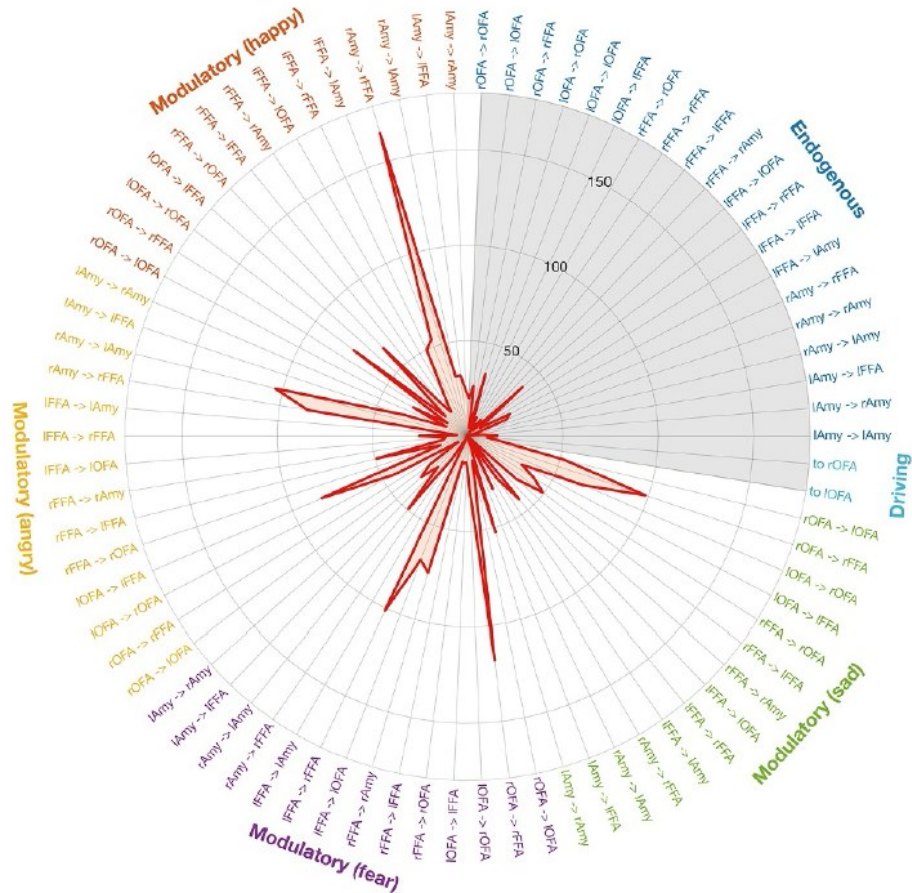


③ Prediction: Two-year outcome in depression

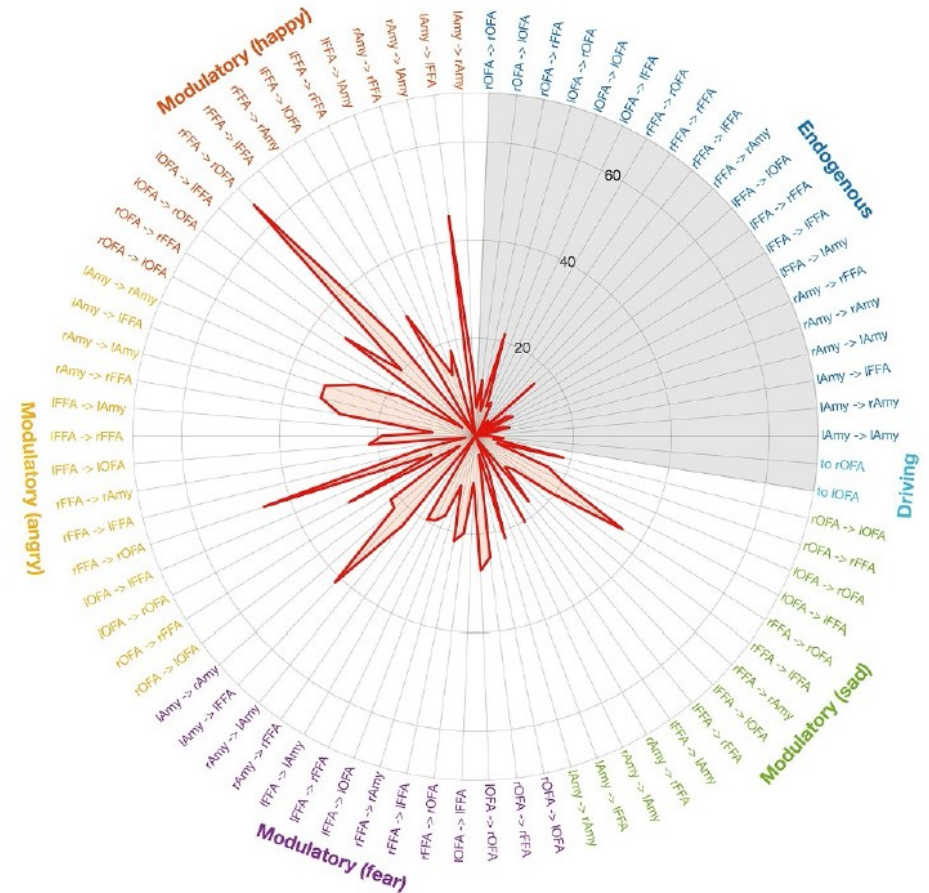


③ Prediction: Two-year outcome in depression

CHR vs. REM

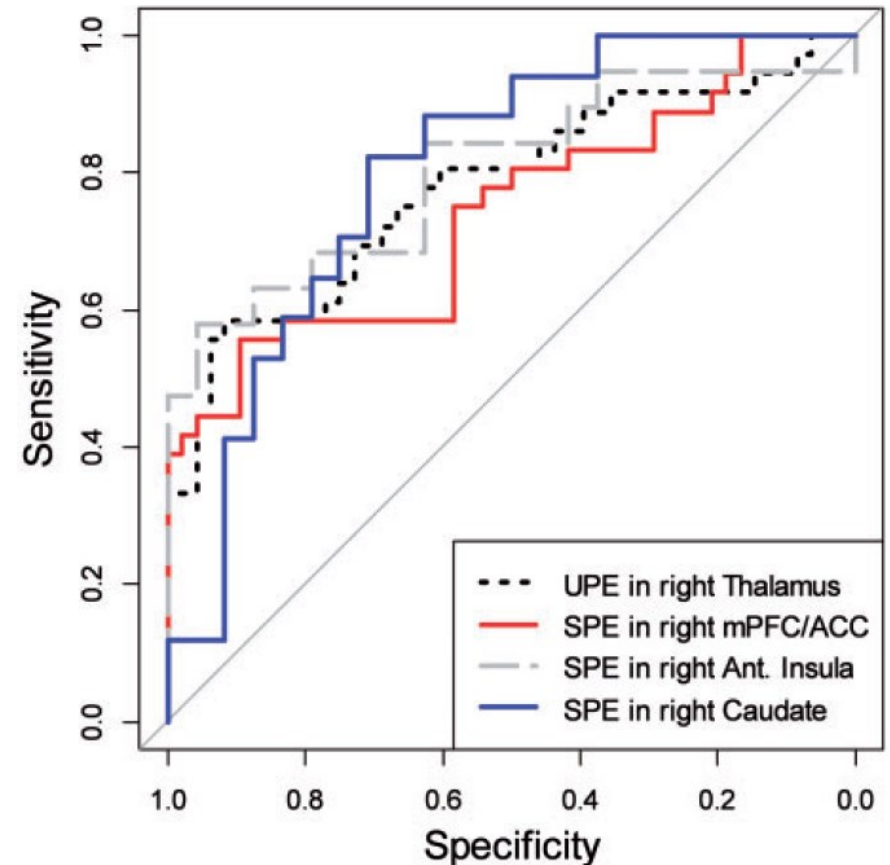


IMP vs. REM



③ Prediction: Future problem use of stimulants

- 88 occasional stimulant users
- "determine whether individual differences in the neural representation of the need to stop in an inhibitory task can predict the development of problem use (i.e. abuse or dependence)"
- fMRI (stop-signal task), Bayesian Hidden Markov Model
- prediction error (PE) activity from 4 brain regions predicted problem use 3 years later
- prediction based on computational variables: sensitivity 62%, specificity 83%
- outperformed predictions based on clinical variables and conventional fMRI analyses



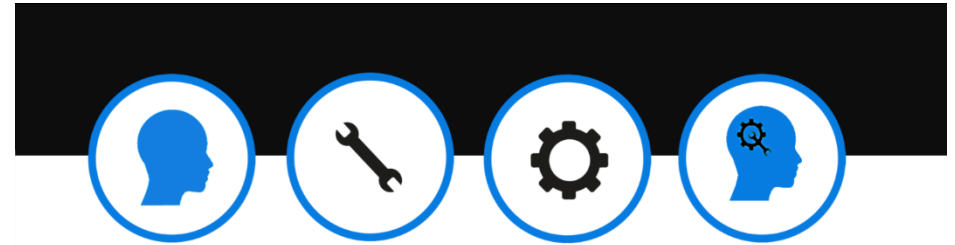
UPE = unsigned PE
SPE = signed PE

Final remark:

What exactly do we mean by "computational"?

- often used in computational psychiatry:
levels of analysis for an information-processing system (David Marr):
 - **computational level:** what problem does the system solve?
 - **algorithmic level:** which representations and operations are used?
 - **implementational level:** how is the system physically realized?
- this is in conflict with the classical concept of "computation" from computer science
 - “**computation**” = **finite sequence of operations (algorithm)** that transform an input set into an output set
- a better terminology might be to replace the "computational level" with "teleological" level

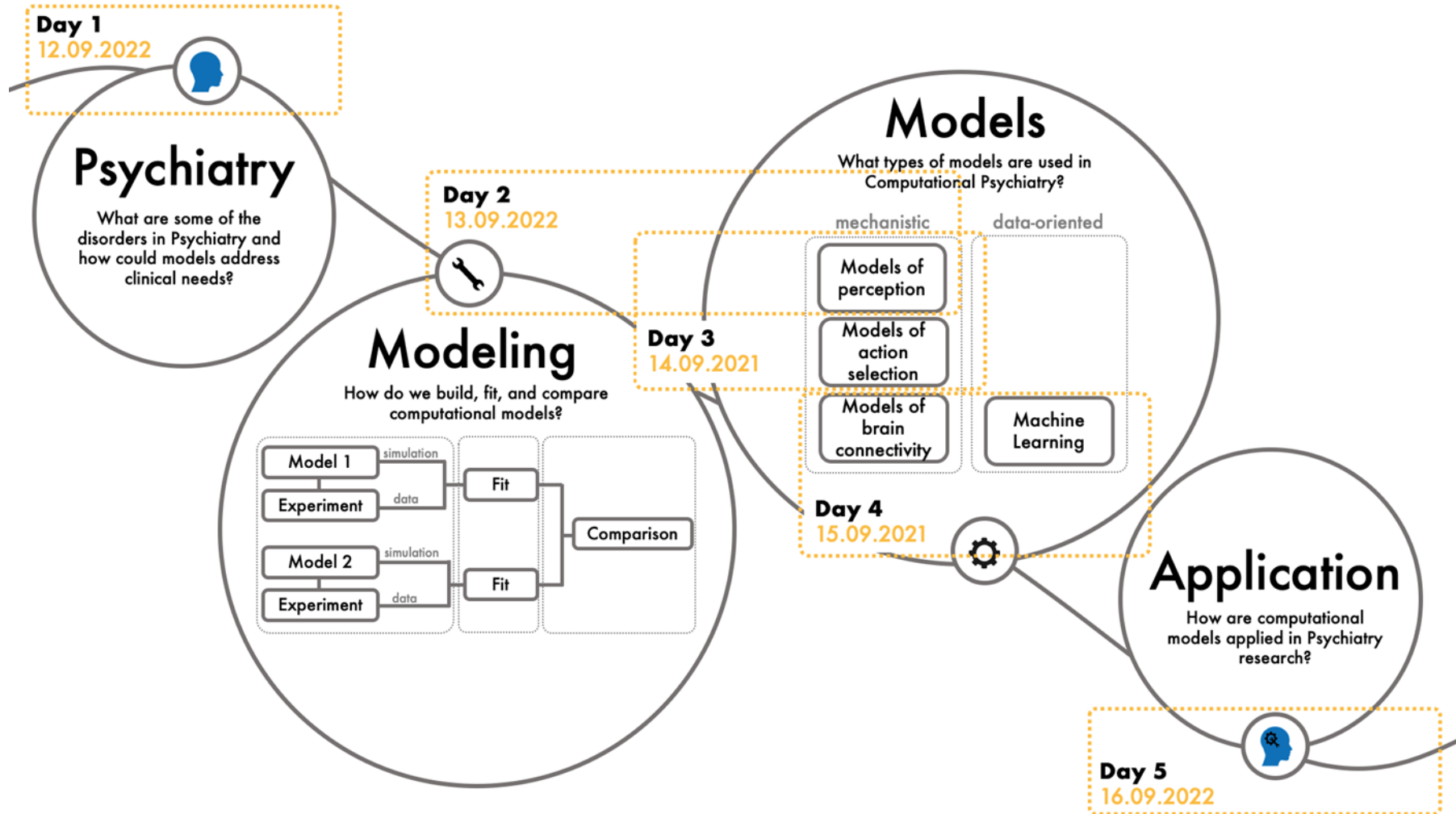
CPC 2022



<http://www.translationalneuromodeling.org/cpcourse/>

- 8th international edition
- originated from our local courses on Computational Psychiatry since 2012
- 2022: first time in hybrid mode
- key features
 - clinical lectures (Monday)
 - methodological lectures (Tuesday – Thursday)
 - application talks (Friday)
 - practical exercises (Saturday) with different open source toolboxes
 - covers models of both neurophysiology and behaviour
 - 46 presenters from 22 international institutions

CPC 2022



Further reading: reviews on computational psychiatry

- Bennett D, Silverstein SM, Niv Y (2019) The Two Cultures of Computational Psychiatry. *JAMA Psychiatry* 76: 563-564.
- Frässle S, Yao Y, Schöbi D, Aponte EA, Heinzle J, Stephan KE (2018) Generative models for clinical applications in computational psychiatry. *Wiley Interdisciplinary Reviews: Cognitive Science* 9: e1460.
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- Petzschner FH, Weber LAE, Gard T, Stephan KE (2017) Computational Psychosomatics and Computational Psychiatry: Toward a joint framework for differential diagnosis. *Biological Psychiatry* 82: 421-430.
- Stephan KE, Mathys C (2014) Computational Approaches to Psychiatry. *Current Opinion in Neurobiology* 25:85-92.
- Stephan KE, Iglesias S, Heinzle J, Diaconescu AO (2015) Translational Perspectives for Computational Neuroimaging. *Neuron* 87: 716-732.
- Stephan KE, Schlagenhauf F, Huys QJM, Raman S, Aponte EA, Brodersen KH, Rigoux L, Moran RJ, Daunizeau J, Dolan RJ, Friston KJ, Heinz A (2017) Computational Neuroimaging Strategies for Single Patient Predictions. *NeuroImage* 145:180-199
- Wang XJ, Krystal JH (2014) Computational psychiatry. *Neuron* 84: 638-654.

**Once again, a very warm welcome –
we hope you will enjoy the CPC 2022!**



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<http://www.translationalneuromodeling.org/cpcourse/>