

Center of Functionally Integrative Neuroscience  
Aarhus University / Aarhus University Hospital - DENMARK



## GLHMM toolbox: Out-of-sample prediction

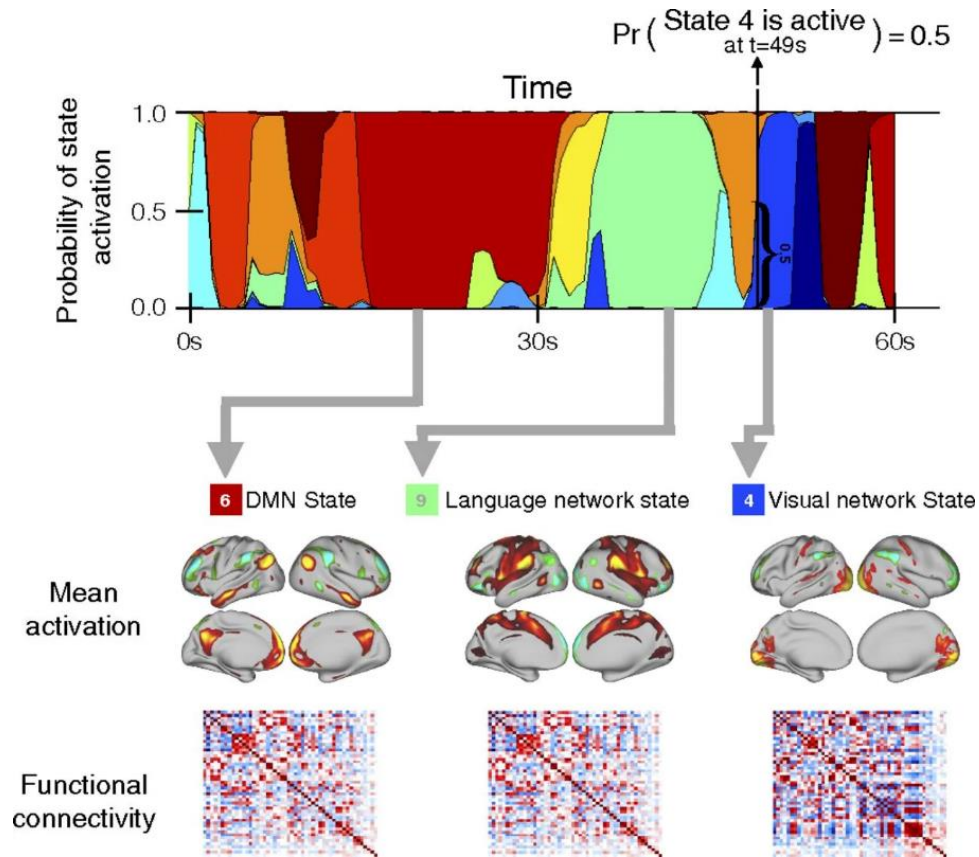
Christine Ahrends, PhD

Analysis workshop CFIN – 26.05.2025



*How can we predict an individual's traits or behaviour from the patterns in which their brain function changes over time?*

# Challenge: HMMs for phenotype prediction

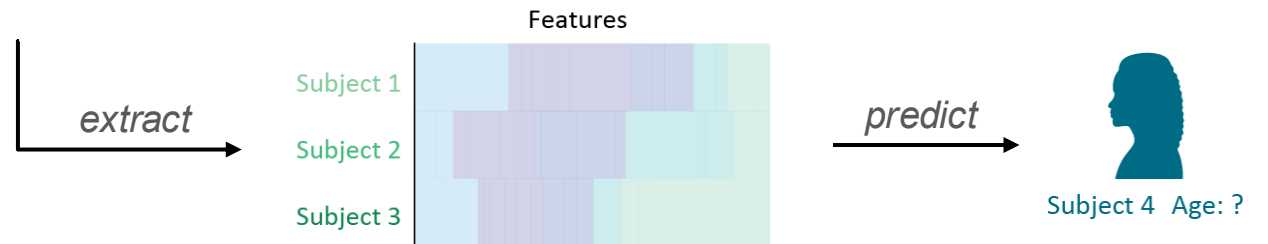
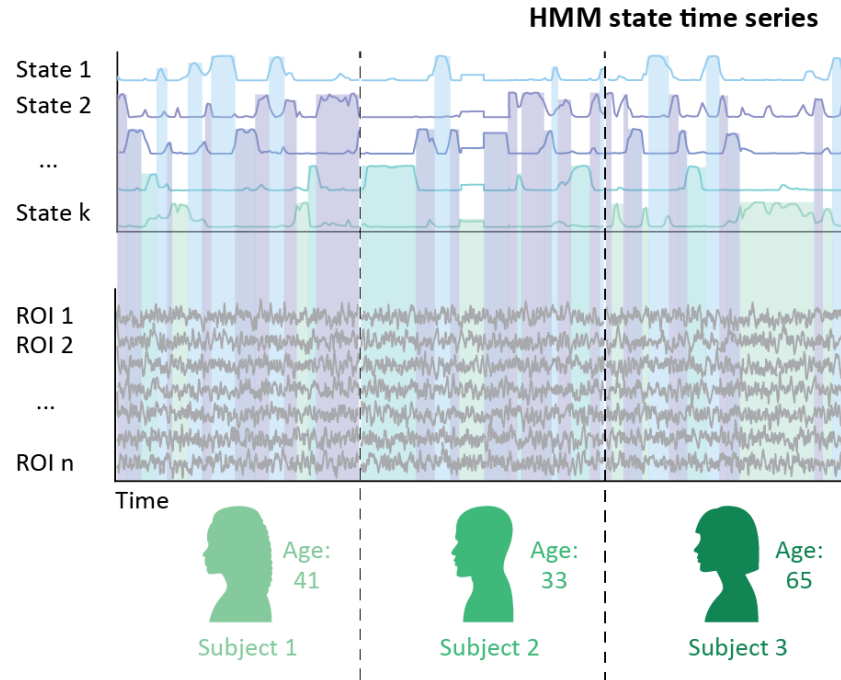


Age?

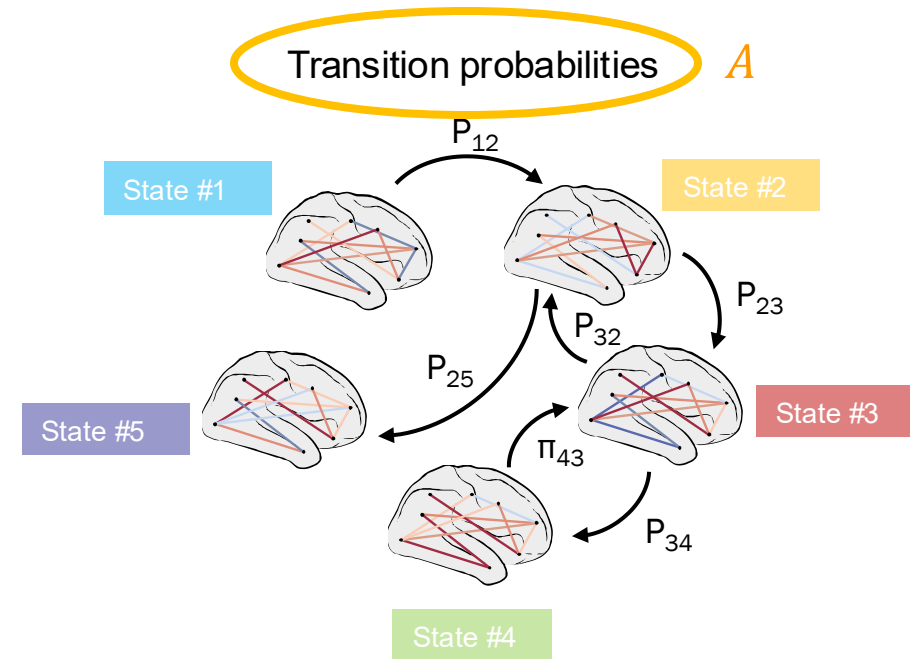
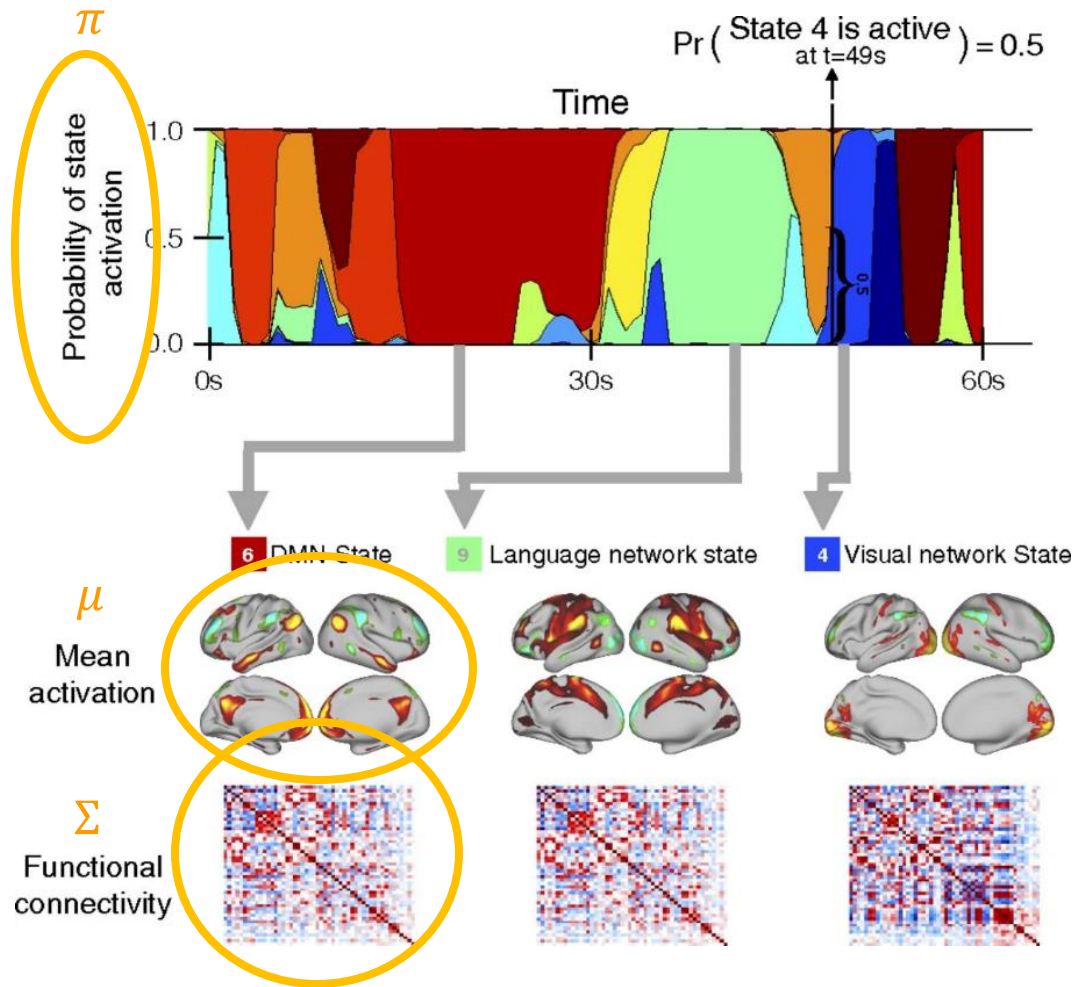
Model of brain dynamics (Hidden Markov Model)

# Summary features

- **Feature engineering approach:**
  - Extract features of interest from HMM
    - Fractional occupancies
    - Switching rates
    - State lifetimes
    - ...
  - Use features to predict individual traits/performance/...
- + Hypothesis-driven
- Losing information



# Challenge: HMMs for phenotype prediction

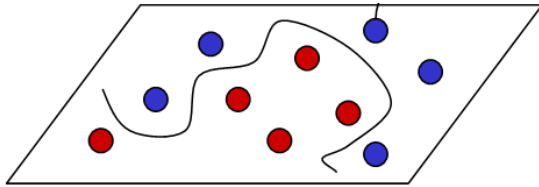


HMM parameters:

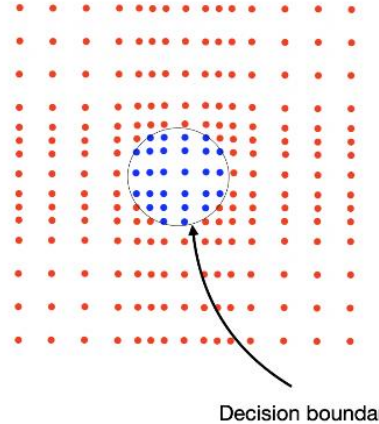
$$\theta = [\pi, A, \mu, \Sigma]$$

→ construct kernel

# Kernel methods



**Input Space**



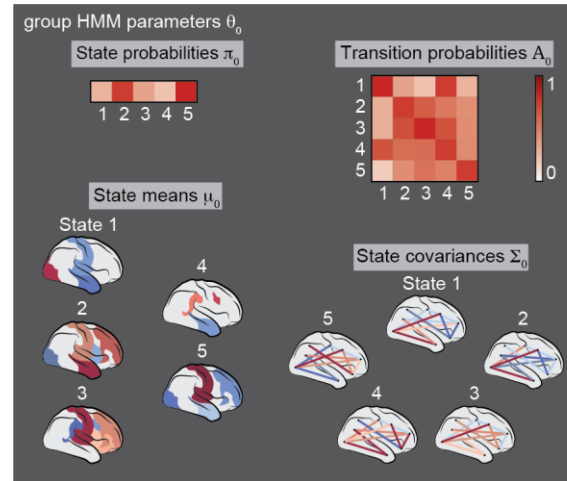
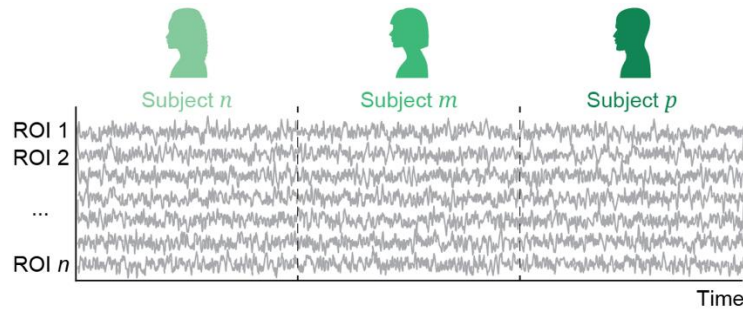
- Find nonlinear decision boundaries using linear model
  - Transform input space into another embedding space
  - Find linearly separating hyperplane in embedding space
- Kernel trick: use kernel function instead of each data point in embedding space
- Similarity functions (e.g. similarity between pair of subjects)
- Computationally efficient, no feature selection



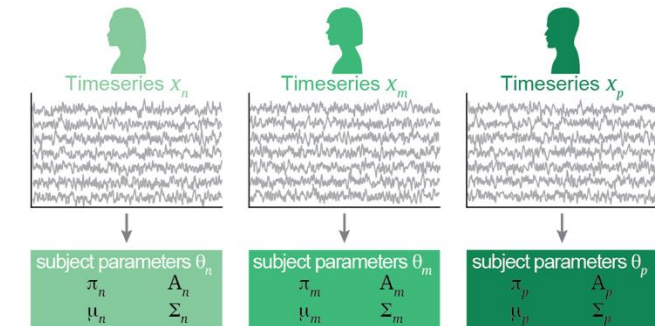
# Kernels from HMM: General approach

## 2. Estimate Hidden Markov model (group-level HMM)

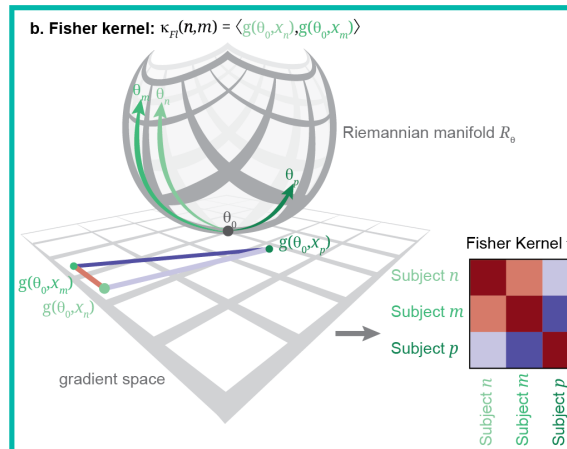
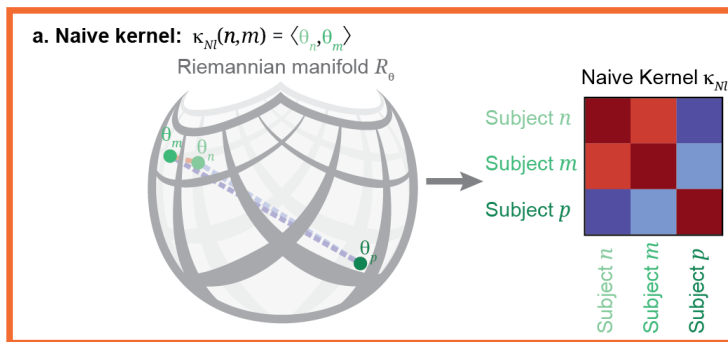
### 1. Group neuroimaging timeseries (e.g. fMRI, MEG, EEG)



### 3. Dual estimation (subject-level HMM)



### 4. Map examples into feature space and construct kernel

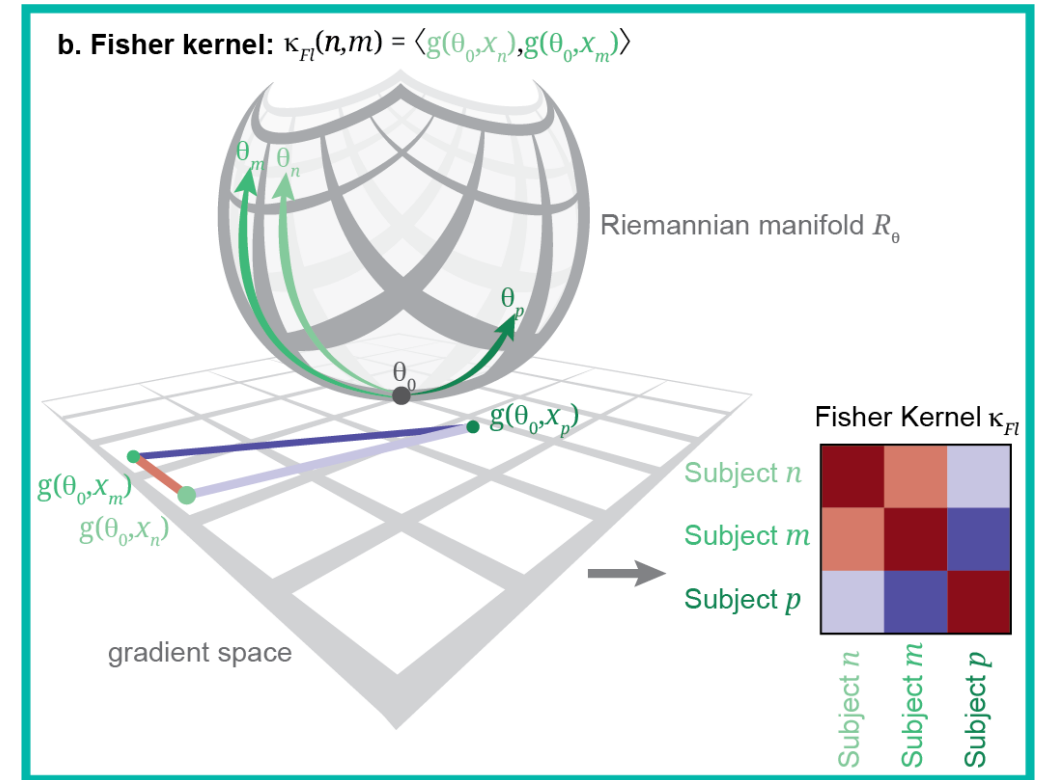
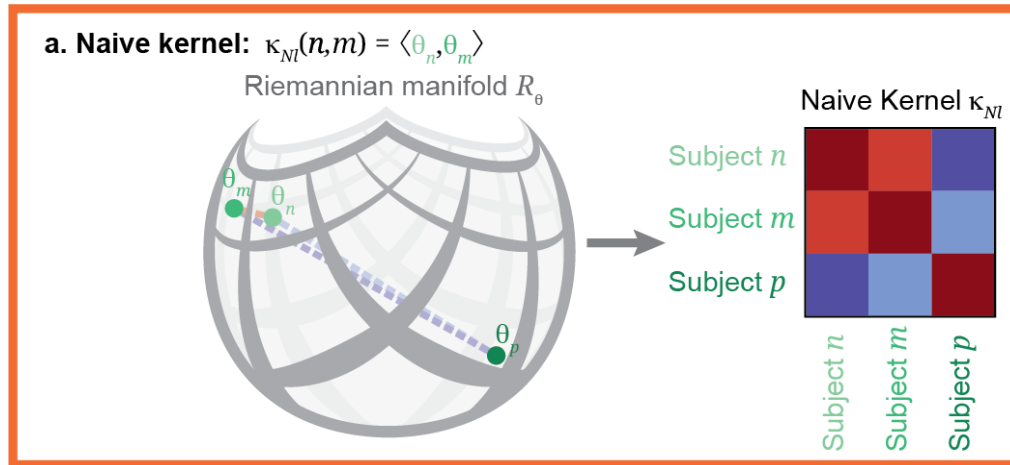


### 5. Kernel-based prediction

	Age	Item A	Item B	Item C	Item D
Subject <i>n</i>	28	102	115	5	87
Subject <i>m</i>	32	110	113	4	33
Subject <i>p</i>	54	99	125	6	18

Ahrends, Woolrich & Vidaurre (eLife, 2025)

# Kernels from HMM: General approach



Ahrends, Woolrich & Vidaurre (elife, 2025)



# The Fisher Kernel

- Fisher score: gradient of the log-likelihood w.r.t. each model parameter:

$$g(\theta^0, x^n) = \left( \frac{\partial \log \mathcal{L}_{\theta^0}(x^n)}{\partial \theta^0} \right) \quad g \in \mathbb{R}^{1 \times (K + K * K + K * M + K * M * M)}$$

- how we need to change the group-level model to better explain an individual's timeseries

- (Practical) linear Fisher kernel:

$$\kappa_F(n, m) = g(\theta^0, x^n)^T g(\theta^0, x^m) \quad \kappa_F \in \mathbb{R}^{N \times N}$$

- Gaussian Fisher kernel:

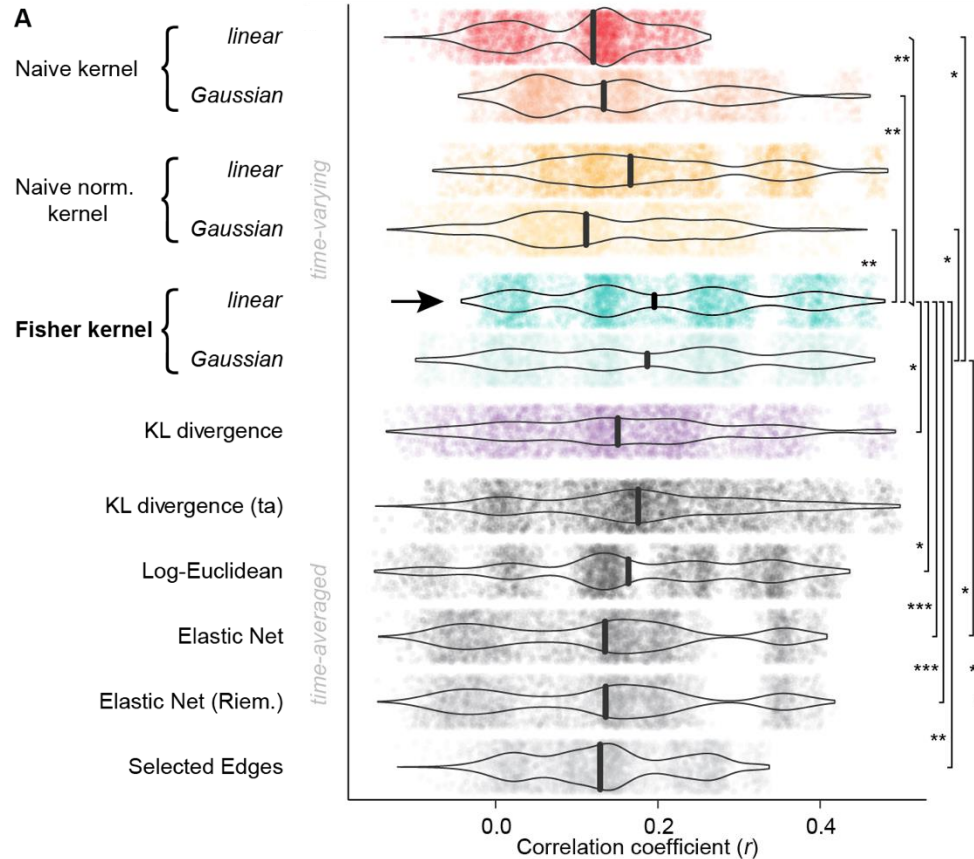
$$\kappa_{Fg}(n, m) = \exp\left(-\frac{\|g(\theta^0, x^n) - g(\theta^0, x^m)\|^2}{2\tau^2}\right) \quad \kappa_{Fg} \in \mathbb{R}^{N \times N}$$

Jaakkola & Haussler (1998) *NIPS*  
Jaakkola et al. (1999) *PICISMB*

# Fisher kernel: Accuracy and reliability

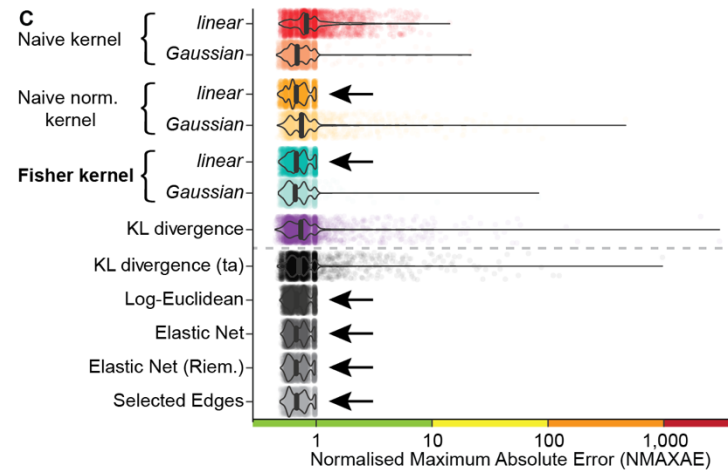
## Prediction accuracy:

## Correlation between predicted and true

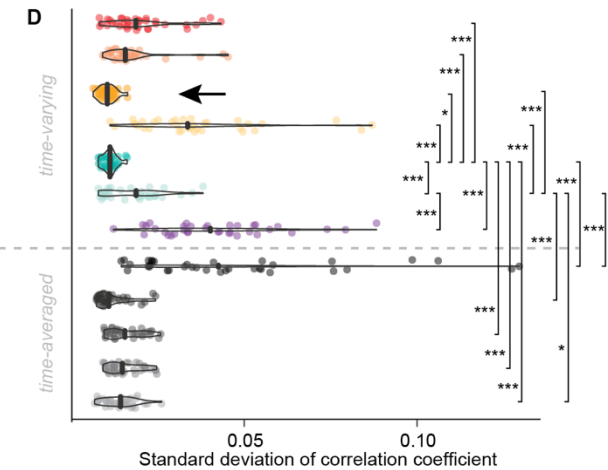


## Reliability:

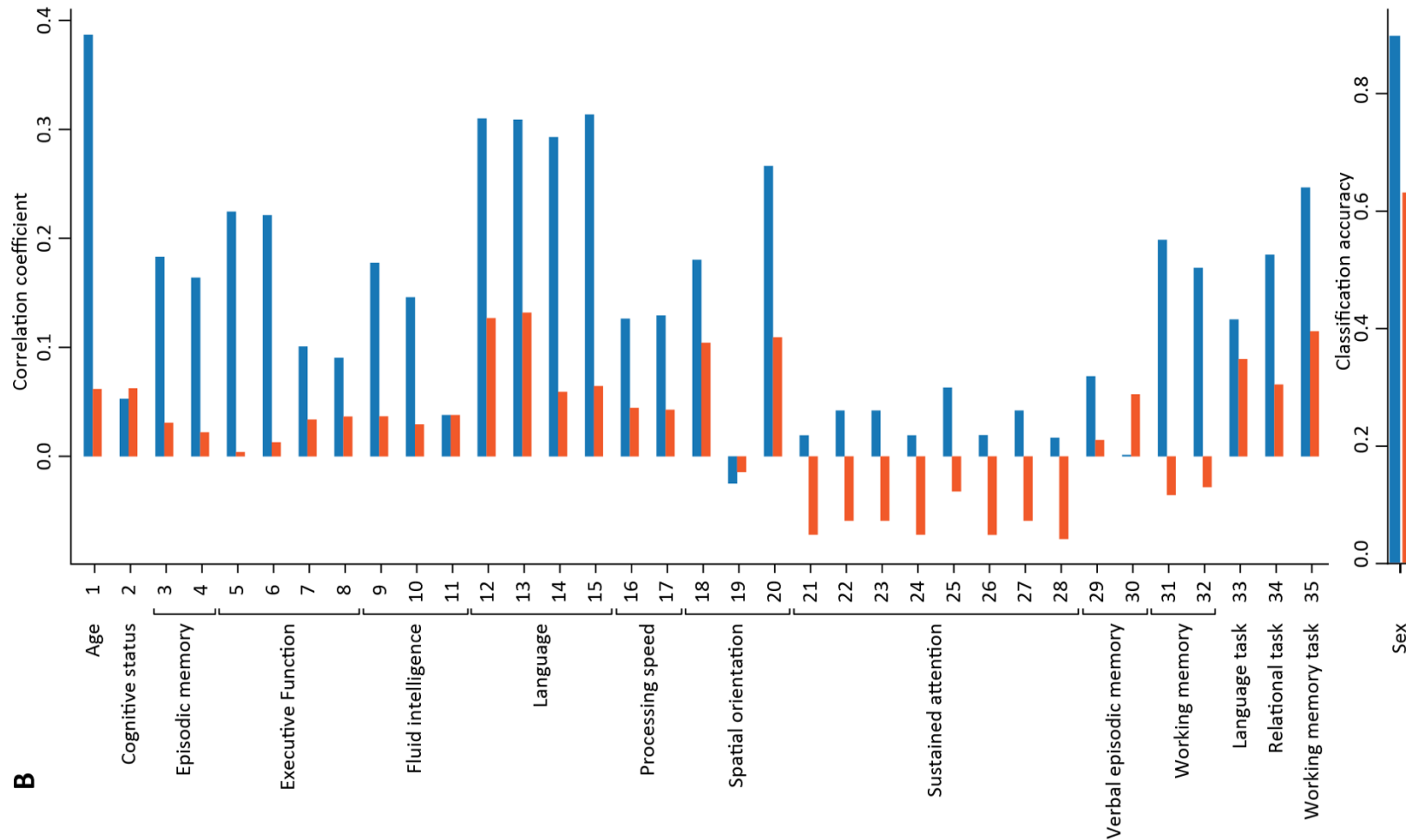
## Risk of excessive errors



## Robustness



# Kernels vs. feature engineering



Fisher kernel

Summary features

# Try it yourself:

- Notebooks:
  - Gaussian HMM 2\_1\_GaussianHMM.ipynb
  - Out-of-sample prediction 2\_2\_Predction.ipynb

## Additional reading:

Ahrends, Woolrich & Vidaurre (*elife*, 2025) Predicting individual traits from models of brain dynamics accurately and reliably using the Fisher kernel.

## Code:

<https://github.com/ahrends/FisherKernel>