

Demixed PCA, or how to tie dimension reduction to experimental conditions

Mihály Bányai
Population Activity Study Group
MTA KOKI, 04.15. 2016.

Demixed Principal Component Analysis

Demixed principal component analysis of population activity in higher cortical areas reveals independent representation of task parameters

Wieland Brendel

Ecole Normale Supérieure,
Champalimaud Neuroscience
Lisbon, Portugal

Ecole

Champalimaud

Dmitry Kobak^{1,*}, Wieland Brendel^{1,2,*}, Christos Constantinidis³,
Claudia E. Feierstein¹, Adam Kepecs⁴, Zachary F. Mainen¹,
Ranulfo Romo⁵, Xue-Lian Qi³, Naoshige Uchida⁶, and Christian K. Machens¹

¹Champalimaud Centre for the Unknown, Lisbon, Portugal

²École Normale Supérieure, Paris, France

³Wake Forest University School of Medicine, Winston-Salem, NC, USA

⁴Cold Spring Harbor Laboratory, NY, USA

⁵Universidad Nacional Autónoma de México, Mexico

⁶Harvard University, Cambridge, MA, USA

*Equal contribution

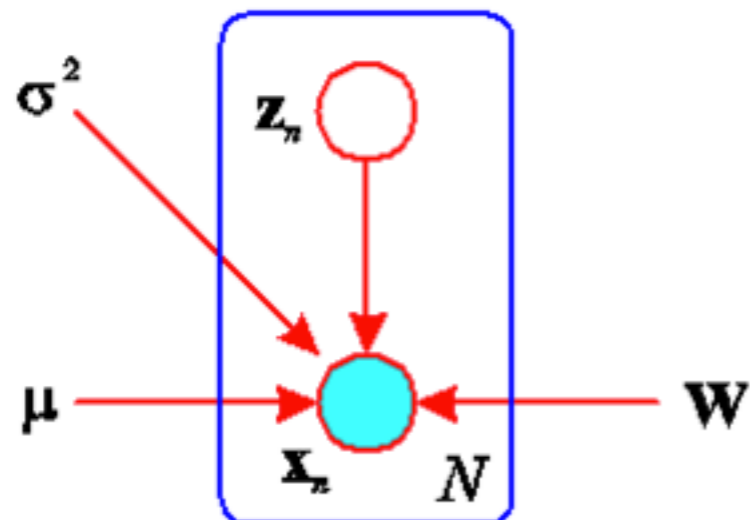
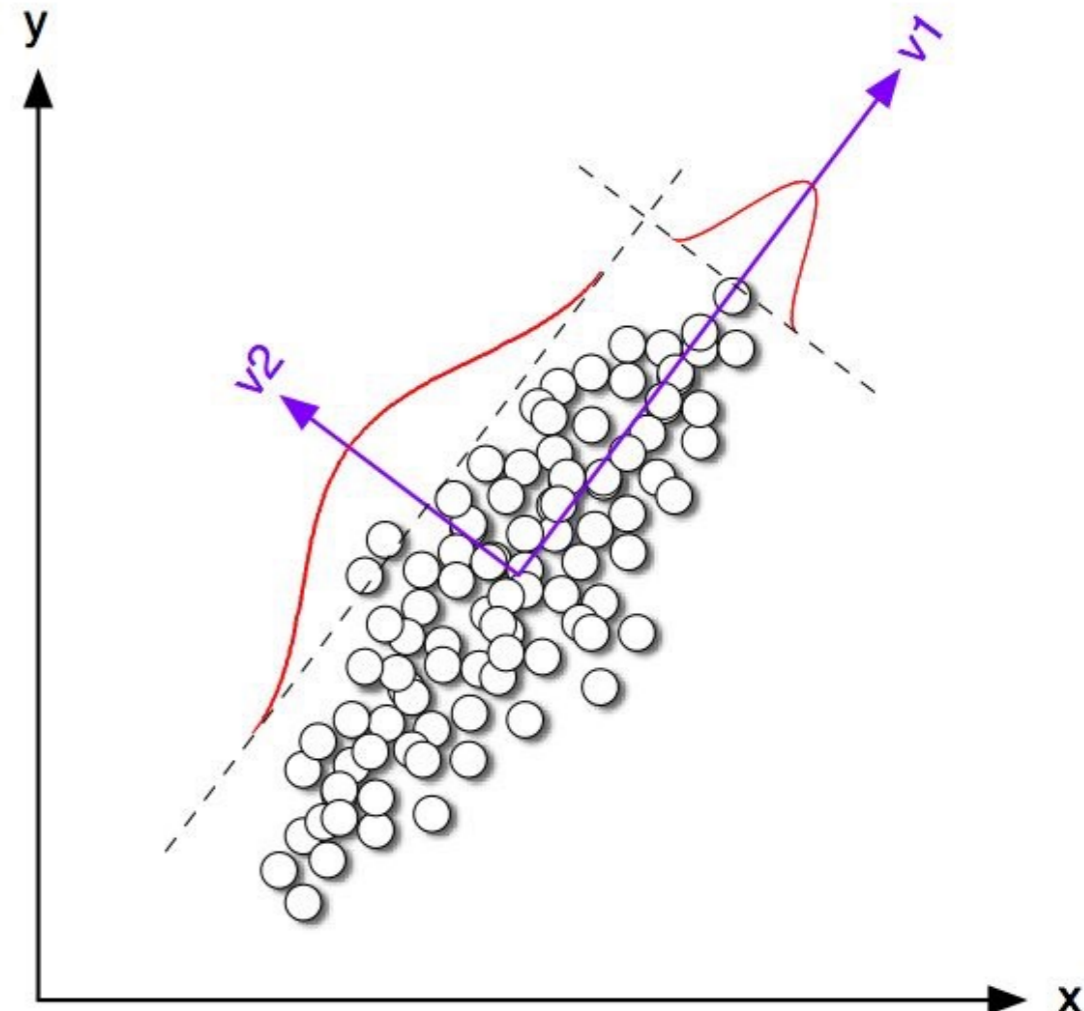
October 2014

Augmenting dimensionality reduction

- When looking for a lower dimensional representation of your data, you typically enforce generic principles about the latent variables
 - independence, orthogonality, sparsity, etc.
 - these may correspond to stimulus features or experimental conditions, but not necessarily (especially given the structure of your model, e.g. linear mappings)
- Another approach is to put conditions explicitly in the model

Review of PCA

- goal: explain as much variance with as few variables (components) as possible
- parameter estimation: eigenvectors of data covariance matrix
 - or an iterative Expectation Maximisation parameter estimation algorithm

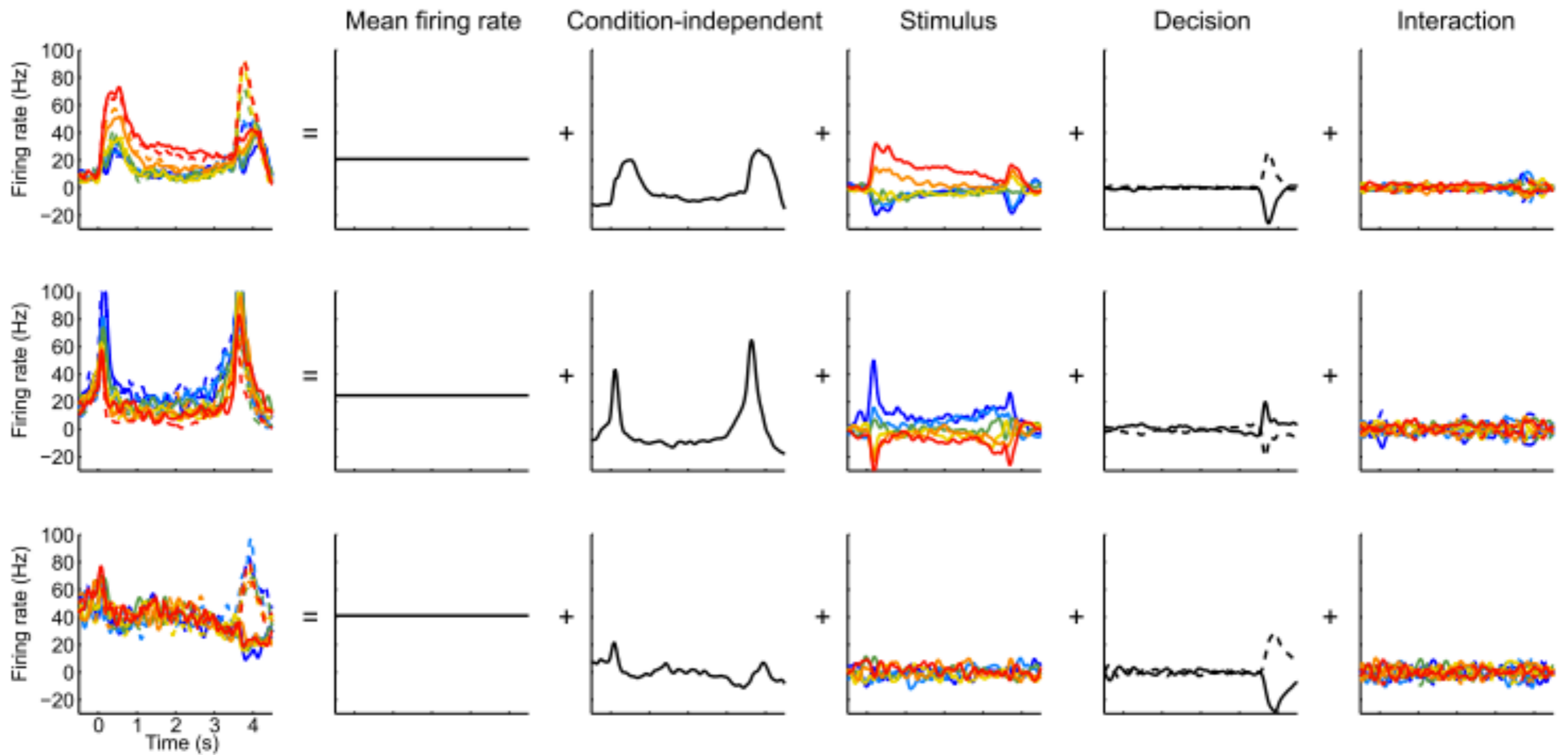


$$p(z) = \mathcal{N}(z; 0, \mathbf{I})$$
$$p(x | z) = \mathcal{N}(x; \mathbf{W}z + \mu, \sigma^2 \mathbf{I})$$

Data

- four different datasets from the labs of Romo, Constantinidis and Mainen
- monkey PFC, rat OFC
 - decision tasks, in each trial there is a stimulus (tactile, visual or olfactory) and the animal makes a binary decision
- they always use the peristimulus time histogram as data
 - they average over trials (needed because cells are not recorded simultaneously, and have diff. number of trials too)
 - data dimension: $\#cells \times (\#stim \cdot \#dec \cdot \#timebin)$

Demixing



Decomposition of variance

- If we decompose the data by marginalising over different parameters, the covariance matrix can be given as a sum of the decomposed covariances

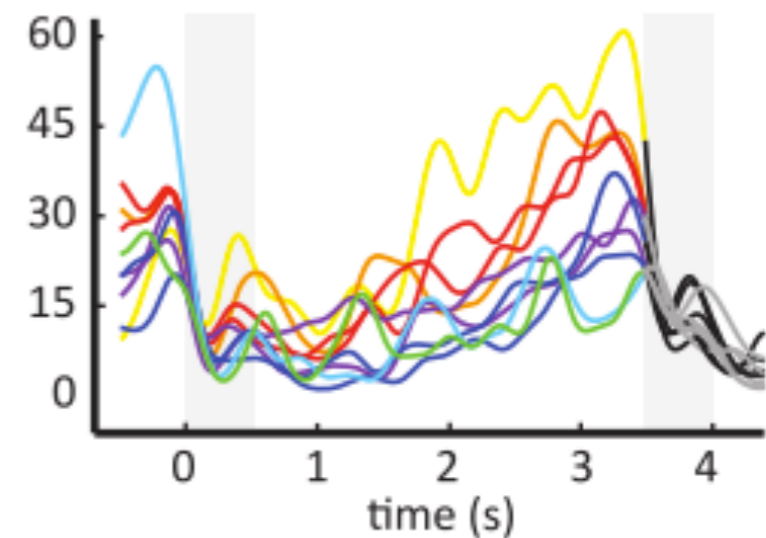
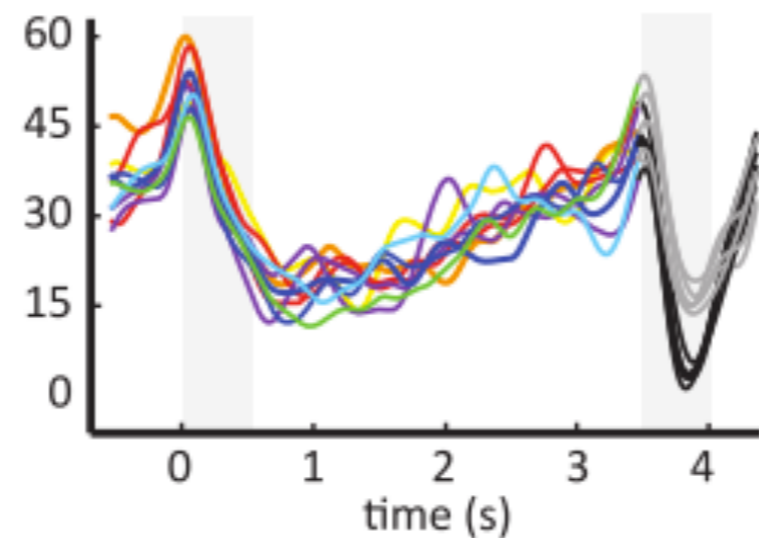
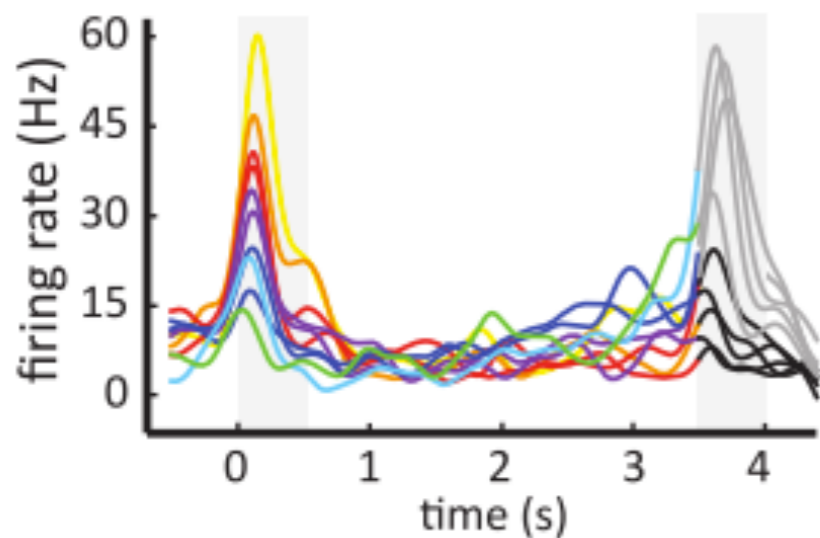
$$x(t, s) = \mu + z(t) + z(s) + z(t, s).$$

$$z(t) = \langle x(t, s) - \mu \rangle_s. \quad z(s) = \langle x(t, s) - \mu \rangle_t. \quad z(t, s) = x(t, s) - \mu - z(t) - z(s).$$

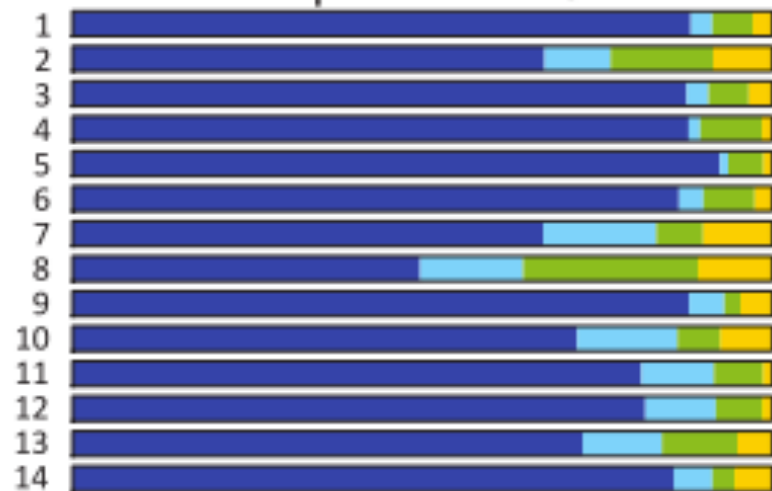
$$\text{Var}(x(t, s)) = \text{Var}(z(t)) + \text{Var}(z(s)) + \text{Var}(z(t, s)).$$

- Instead of regular principal components, we are looking for ones that describe variance only in the direction of the eigenvectors of one of the decomposed covariances

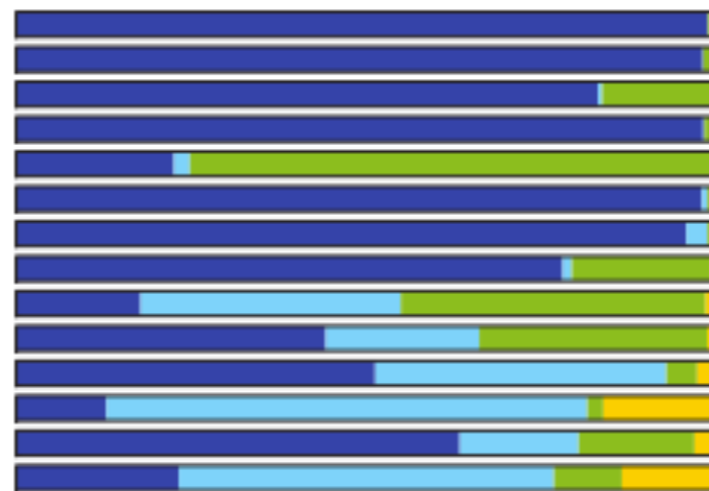
Selectivity of neurons and principal components



sample neurons

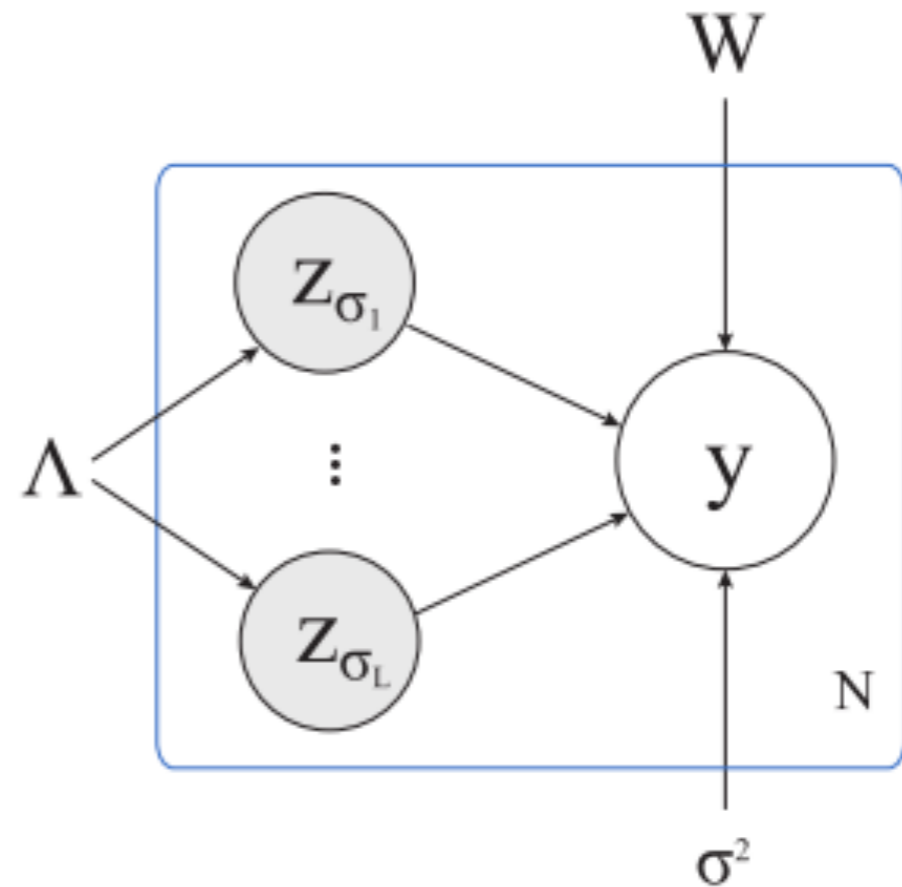
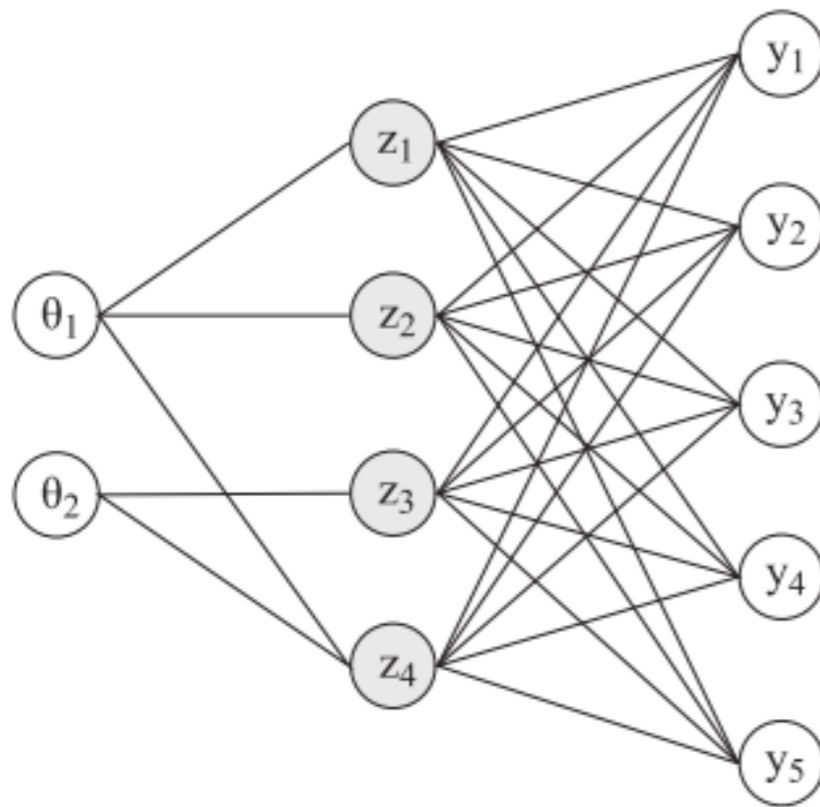


PCA



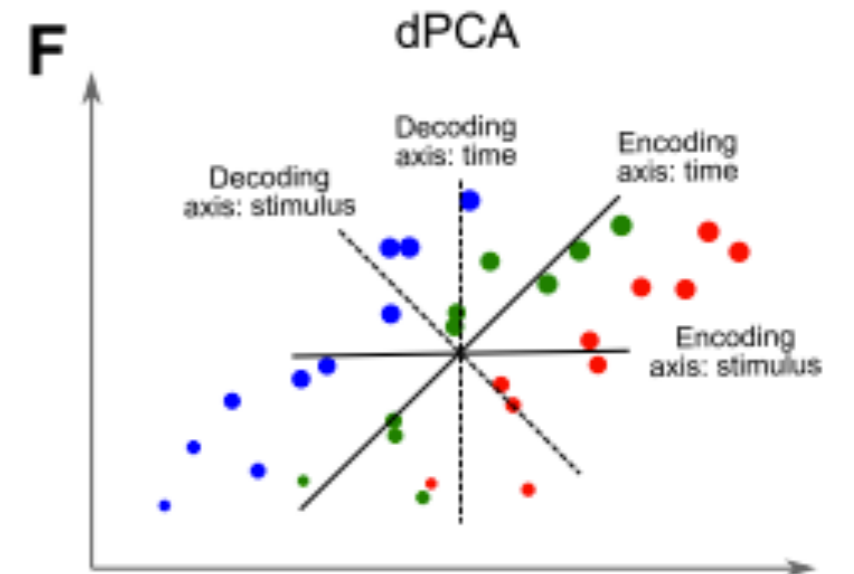
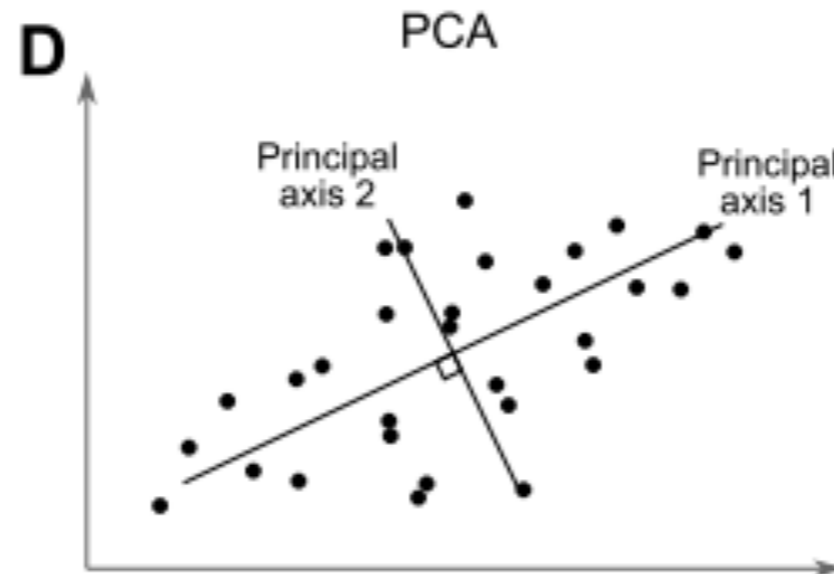
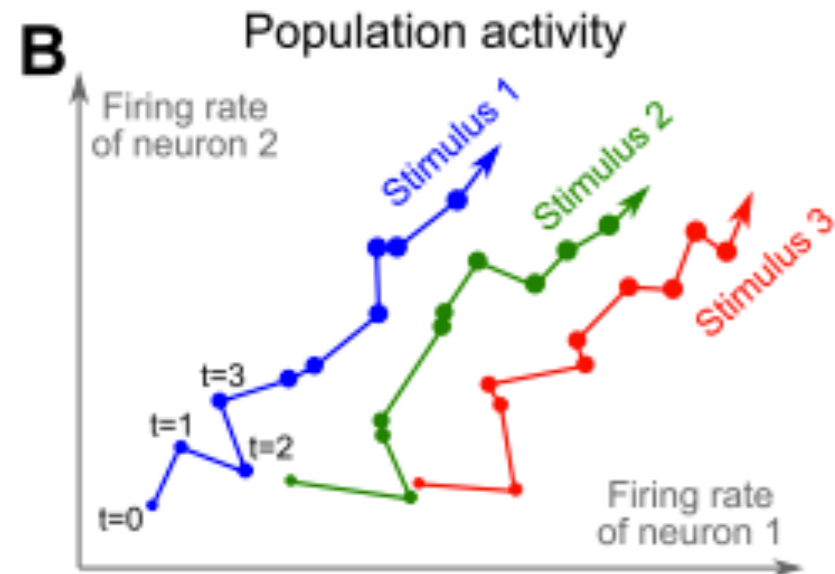
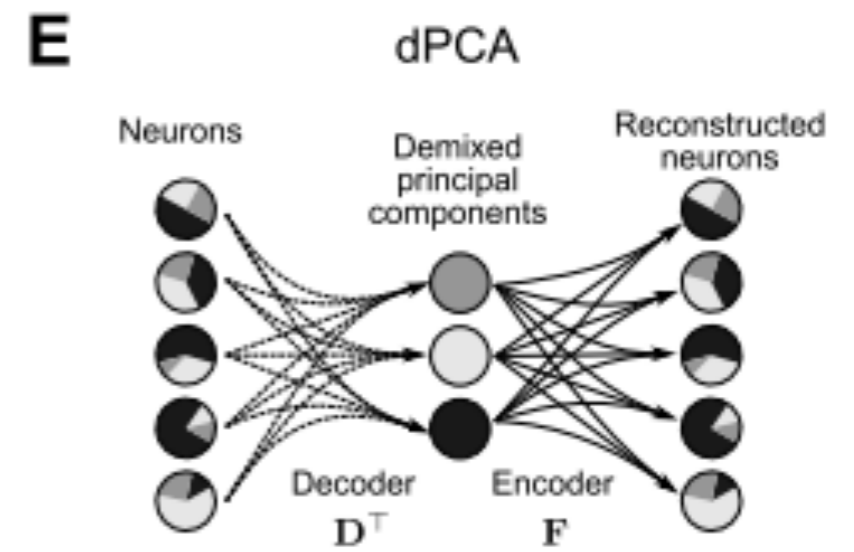
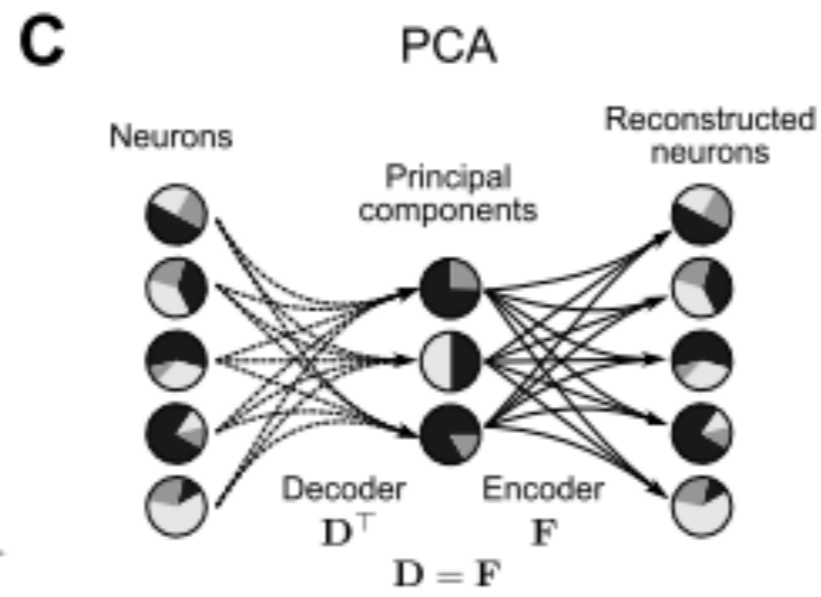
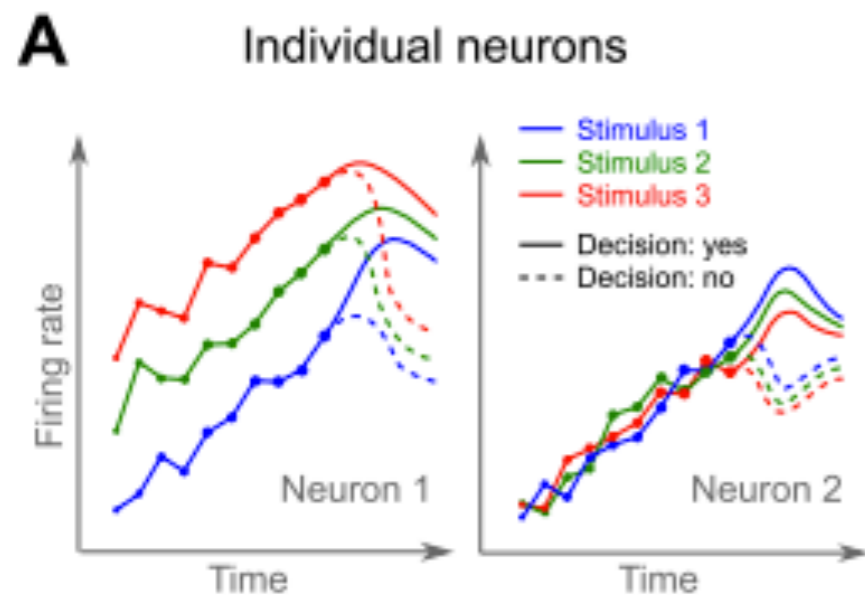
- blue - cond. indep. / time
- green - stimulus
- yellow - decision

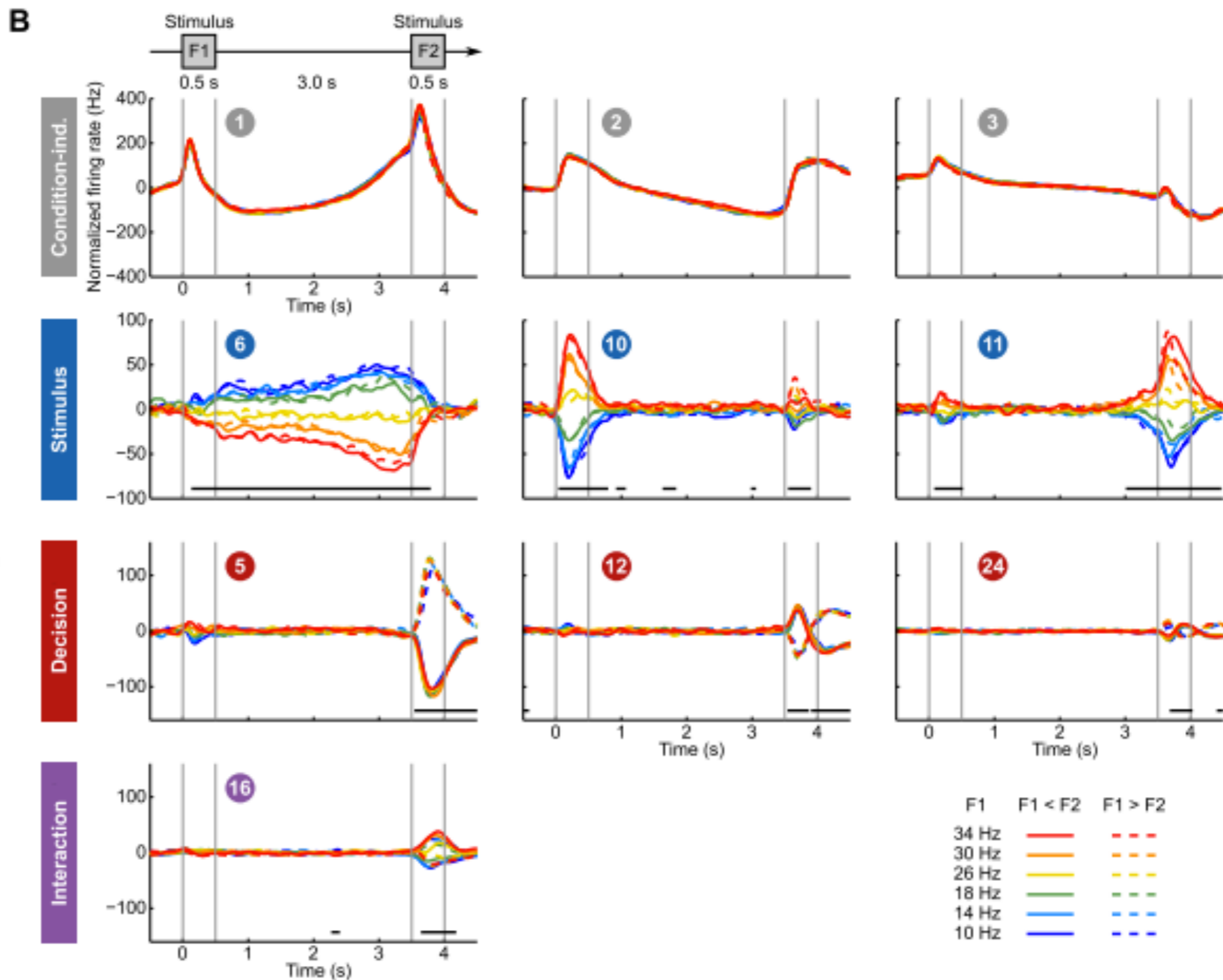
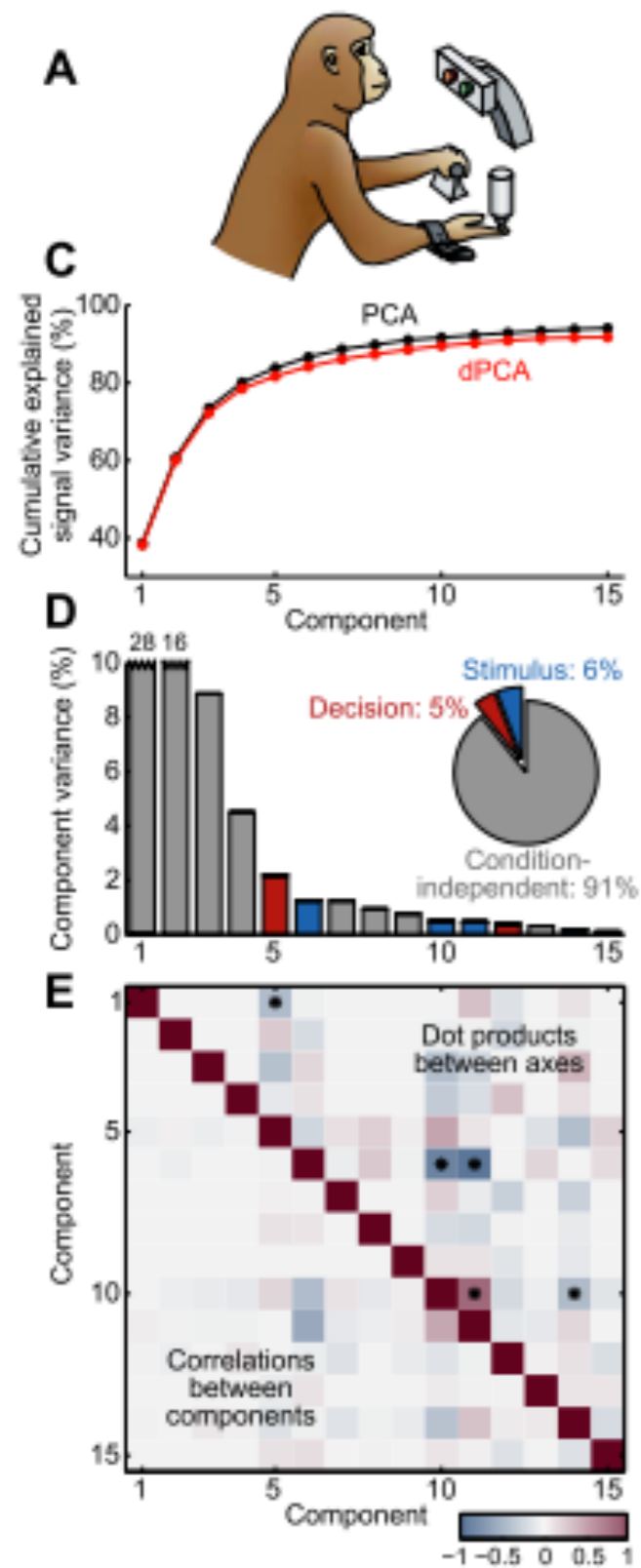
Generative model of dPCA



- An EM algorithm can be derived to find the components by adding constraints to the standard PCA inference

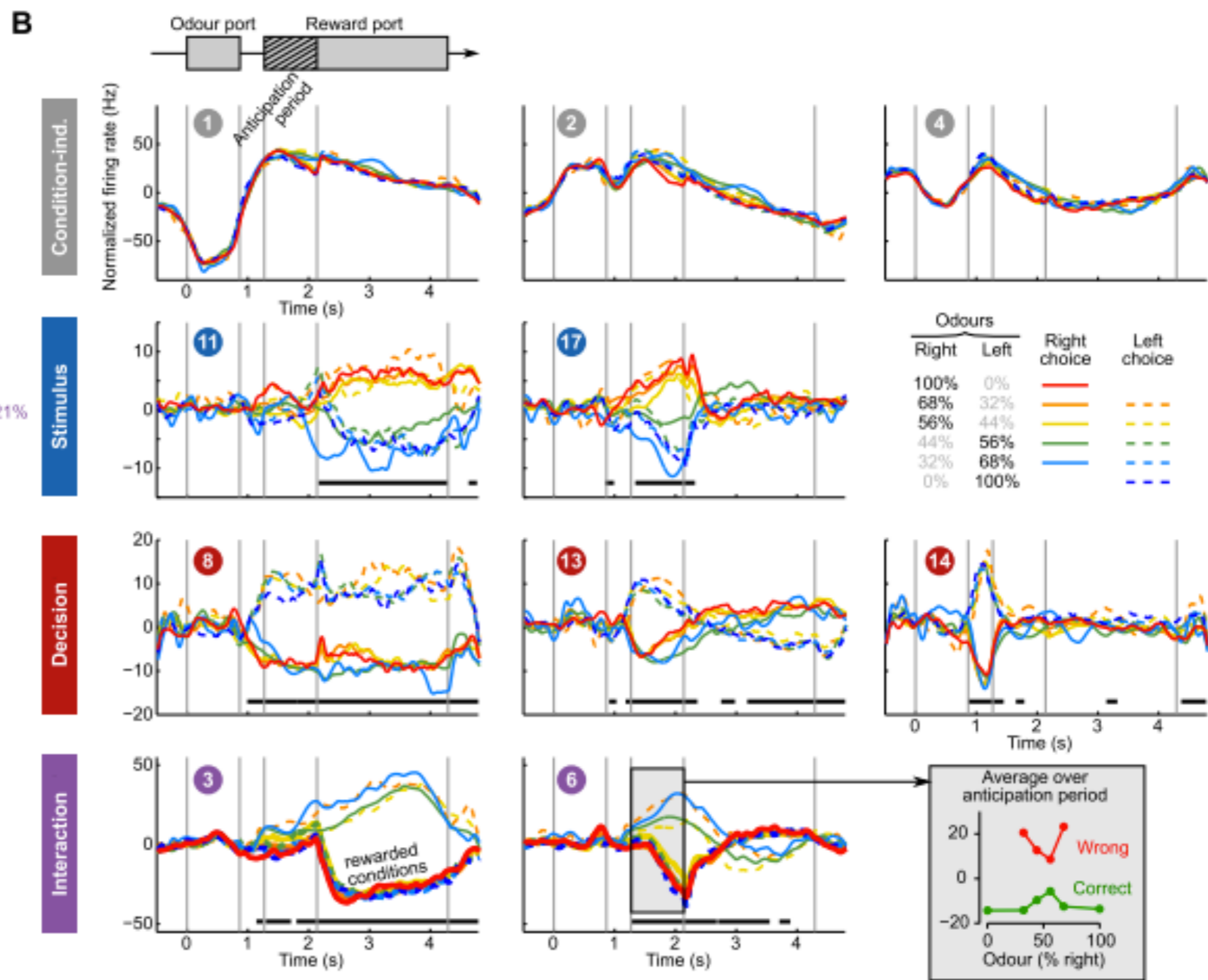
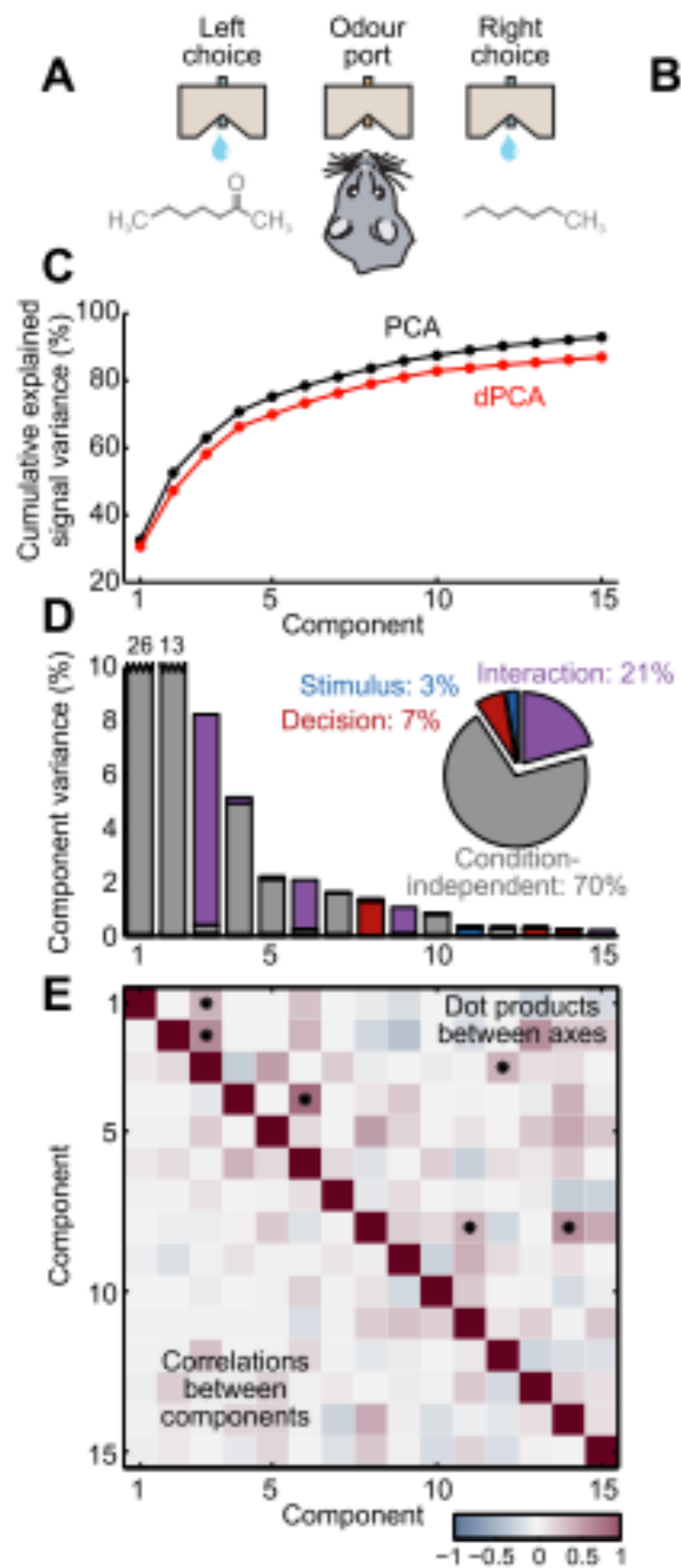
PCA vs. dPCA





Insights

- Linear demixing of the population activity is possible
- Most of the variance is explained by components not related to any conditions
- Some components are the derivatives of each other -> similar activity patterns arise in the population with temporal shifts
- Using decision-related components as fixed linear decoders, one can decode the decision from the population activity with ~75% accuracy



Limitations

- Needs lots of neurons
- All parameter combinations need to be present in the data
- Applied only to PSTH here, in theory it might be applicable to trial-to-trial covariability, but not tested

Functional generative models vs. demixed dimensionality reduction

- Generative models of perception
 - neural activation / latent causes \rightarrow stimulus
 - emphasis on prediction of neural activity statistics
- Demixed dimension reduction
 - stimulus / conditions \rightarrow neural activation
 - emphasis on identification of variable mappings
- Hypotheses about neural representations are formalisable in both frameworks