

L10: Dimensionality Reduction

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Free tutorial suggestions (next week)

1) Program a game in python: "Rock, Paper, Scissors"

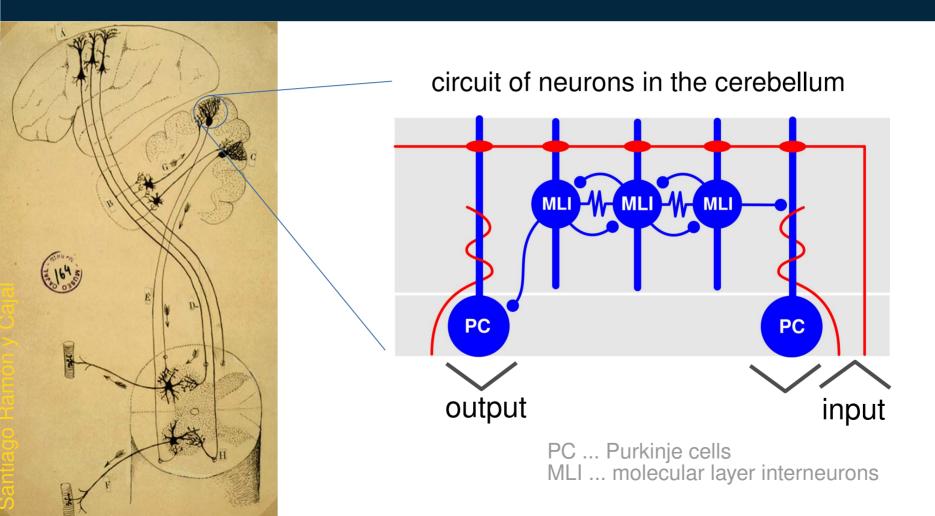


2) Resolve typical programming errors (bugs) in form of a quiz

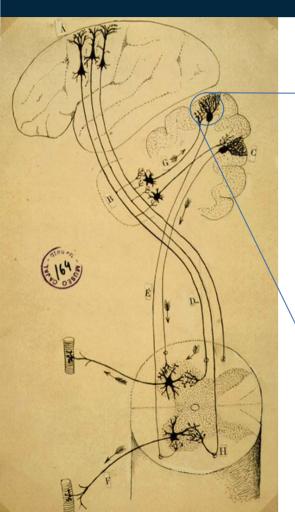
3) Re-work on programming concept/analysis concept suggested by YOU.

4) Start to work on a "fictive" end-of-course project : discuss approach, tips, tricks, typical problems

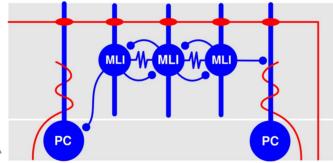
Cerebellum and motor control



Cerebellum and motor control

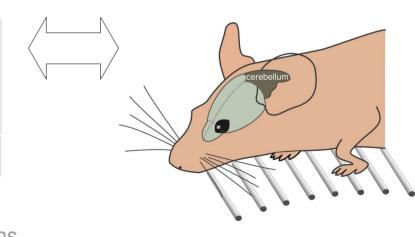


circuit of neurons in the cerebellum



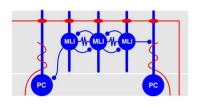
PC ... Purkinje cells MLI ... molecular layer interneurons

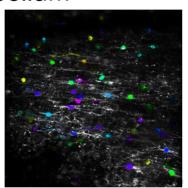
animal walking on treadmill with rungs

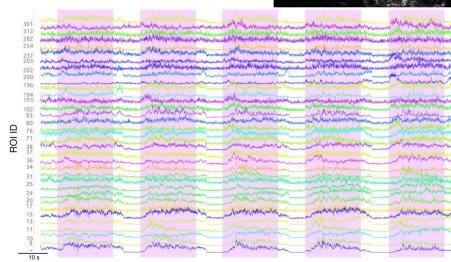


Investigating link between activities and walking

Ca imaging: activity of neurons in the cerebellum

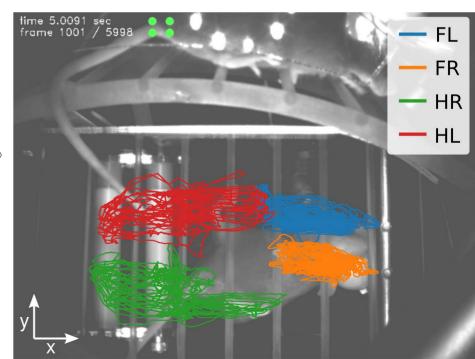








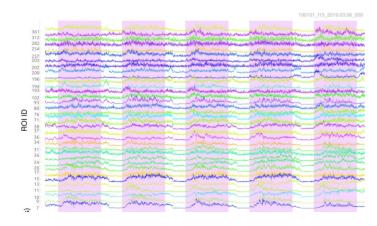
Video: animal walking on treadmill with rungs



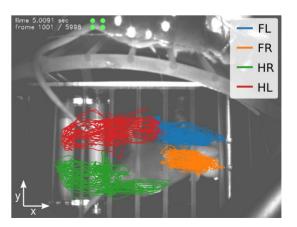
Intuition behind dimensionality reduction

Dimensionality reduction is applied in settings in which there are D measured variables, but one suspects that these variables covary according to a smaller number of explanatory variables K, where K

D measured variables (e.g. neural activities)

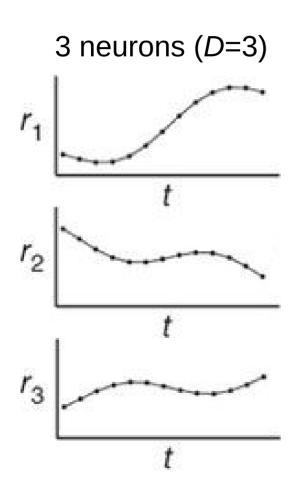


K explanatory variables (e.g. paw movement)

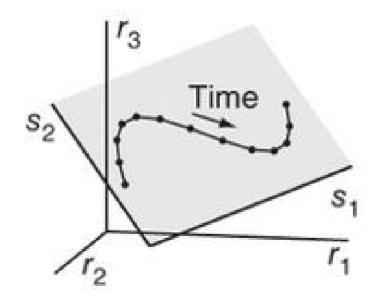


Dimensionality reduction

- Dimensionality reduction methods discover and extract these K explanatory variables
- explanatory variables are called *latent variables* because they are not directly observed
- any data variance not captured by the latent variables is considered to be noise
- example: neural population activity recorded neurons belong to common network and are likely not independent of each other
 - → fewer latent variables might be needed to explain population activity

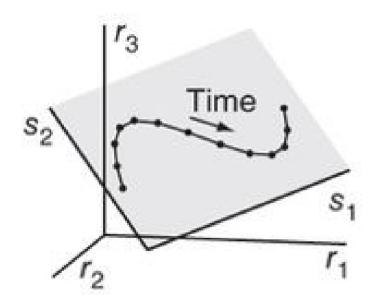


Population space



population activity as trajectory in D-dim space

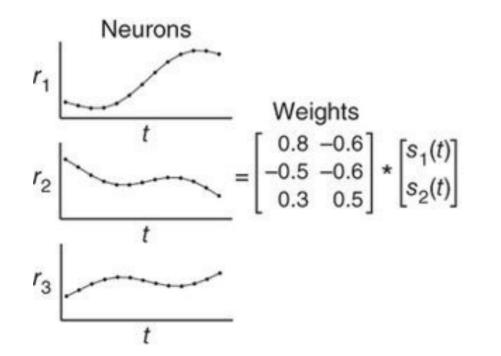
Population space



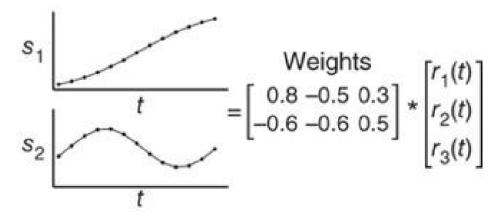
- each point represents the population activity at a particular time
- the points trace out a trajectory over time, time is not plotted, time evolves implicitly along axis
- population activity lies in a plane
- each point can be equivalently referred to using the high-dimensional coordinates
 [r₁,r₂,r₃] or the low-dimensional coordinates [s₁,s₂]
- the points trace out a trajectory over time, time is not plotted, time evolves implicitly along axis

neural activities can be reconstructed from weighted combination of the latent variables

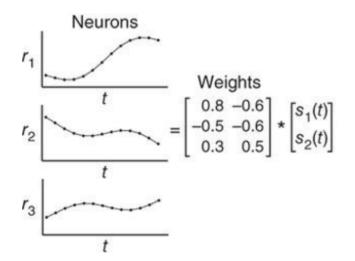
latent variables can be obtained by taking weighted combination of population activity



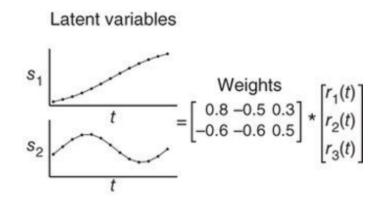
Latent variables



neural activities can be reconstructed from weighted combination of the latent variables

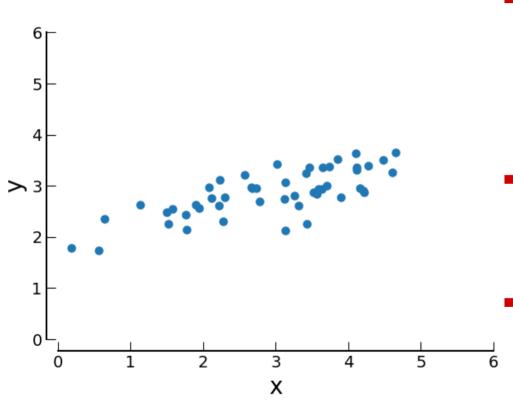


latent variables can be obtained by taking weighted combination of population activity



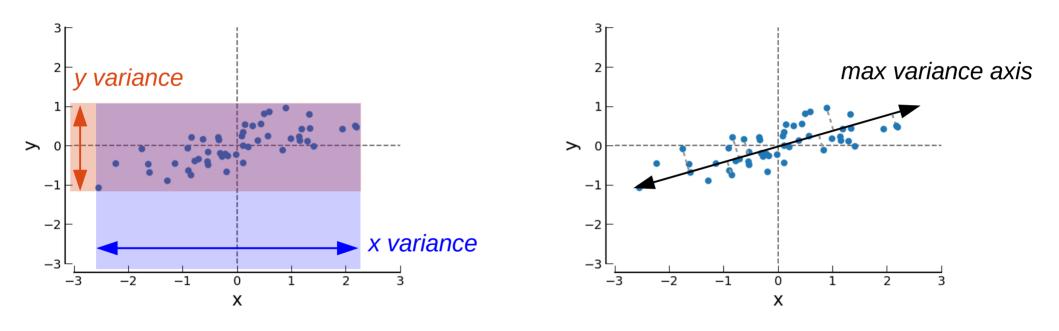
→ Weights and latent variables are determined by the dimensionality reduction method

How does Principal Component Analysis work?



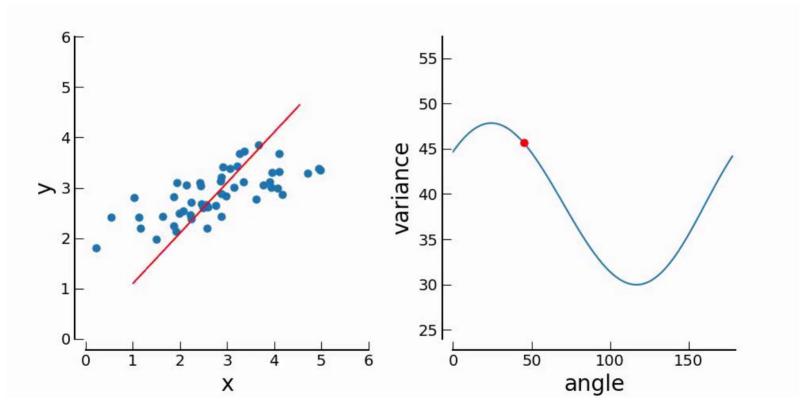
- Principal Component Analysis (PCA) is transformation method creating combinations of the original variables. PCA comes form the field of linear algebra.
- Aim: the new combination of variables will capture as much variance in the data as possible while eliminating correlations
 - **Procedure:** PCA creates new variables by transforming the original observations to new dimensions using eigenvector and eigenvalues calculated from the covariance matrix of the original variables

How does Principal Component Analysis work?

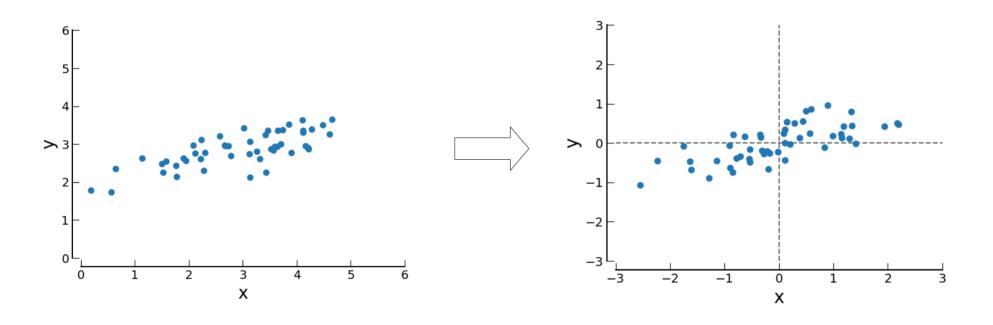


- PCA tries to find successively axes which account for the maximal variance in the data.
 - data is projected onto an axis and orientation varied to maximize variance

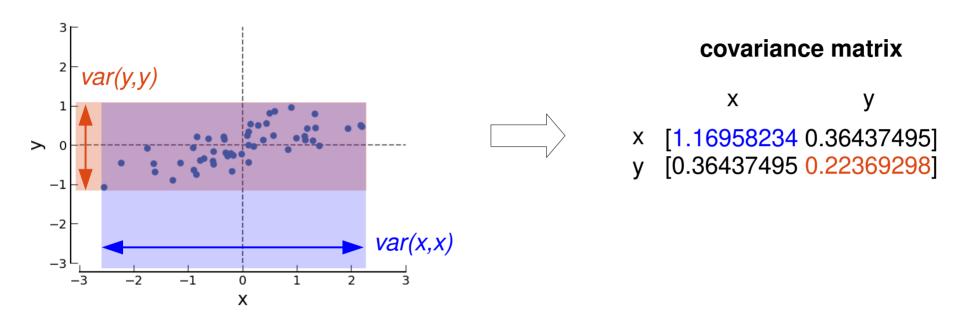
How does Principal Component Analysis work?



 PCA tries to find successively axes which account for the maximal variance in the data.



1) Centering the values of the input variables, i.e. substracting the mean (sometimes the data has to be **normalized** too if one considers variables of different units or scale)

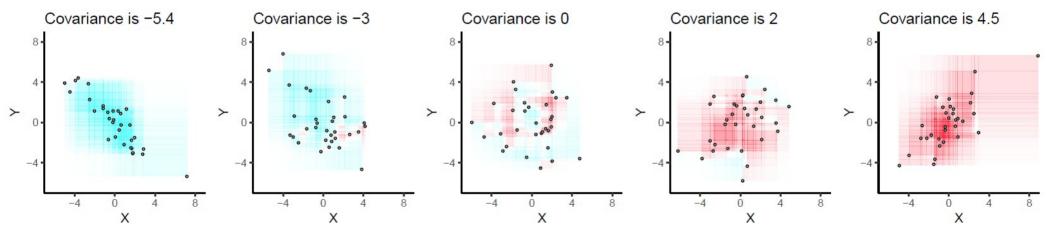


2) Calculating the covariance matrix

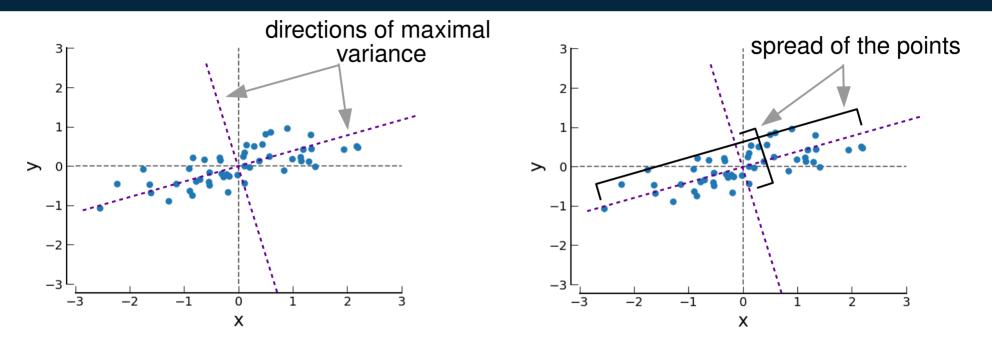
- symmetric matrix as covariance has no direction

Covariance: measures joint variability of variables

$$cov(x, y) = \frac{1}{n} \sum_{i=1}^{n} (x_i - \overline{x})(y_i - \overline{y})$$

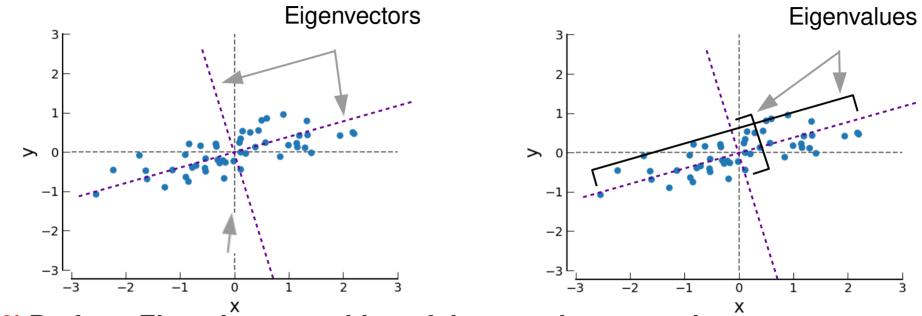


- a large covariance (positive or negative) indicates a strong linear relationship btw. the variables
- covariance ~0 indicate weak or non-existing linear relationship



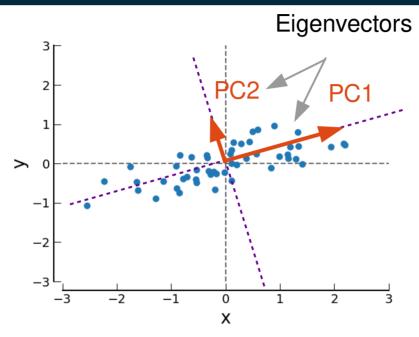
3) Perform Eigendecomposition of the covariance matrix this is where PCA find Eigenvectors and Eigenvalues of the data-set

- Eigenvectors of the covariance matrix are the axes of the principal components
- Eigenvalues describe the magnitude of the eigenvector (spread of the points)



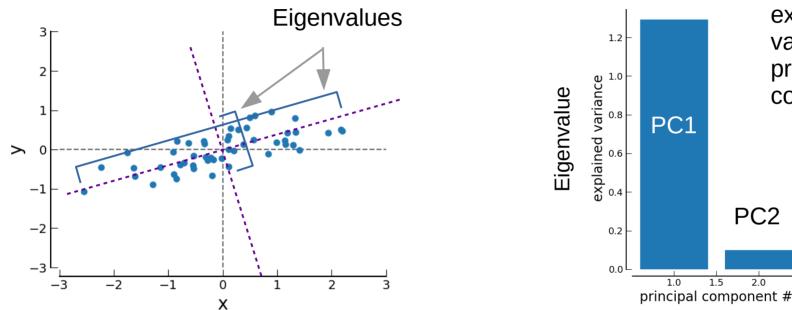
3) Perform Eigendecomposition of the covariance matrix

$$A x = \lambda x$$
 covariance matrix (symmetric) Eigenvector Eigenvalue (scalar)



3) Eigendecomposition of the covariance matrix → Eigenvectors

- the first eigenvector will span the greatest variance found in the dataset
- all subsequent eigenvectors will be perpendicular (orthogonal), *i.e.*, each of the principal components will be uncorrelated with each other



explained variance by the principal components PC2

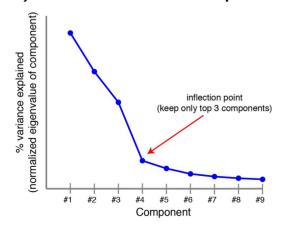
3) Eigendecomposition of the covariance matrix \rightarrow Eigenvalues

- the first eigenvector spans the greatest variance
- principal components are sorted by descending eigenvalues
- principal components with the highest eigenvalues account for most variance in the data

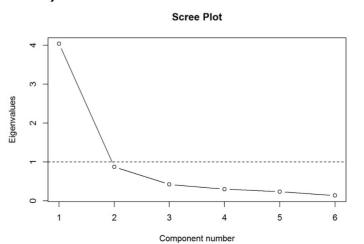
4) Select number of principal components to keep

- PCA returns as many principal components are there are variables
- → How do we know how many of these factors are essential?
- most of the variance will be accounted for by the first principal components
- various ways to select principal components to keep

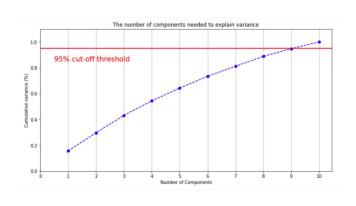
A) look for inflection point

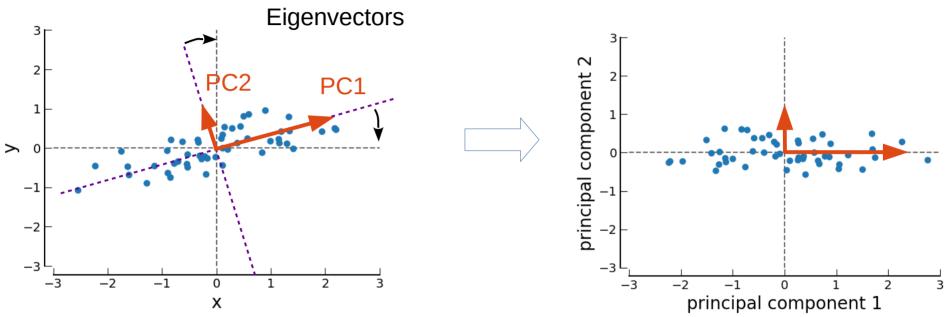


B) Kaiser criterion EV > 1



C) threshold on variance explained



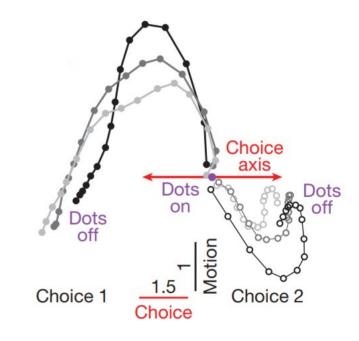


5) Original data is converted to the selected principal components

- projection matrix are simply selected eigenvectors concatenated to a matrix matrix is multiplied with original observations
- → transformed dataset projected onto new space spanned by the principal components

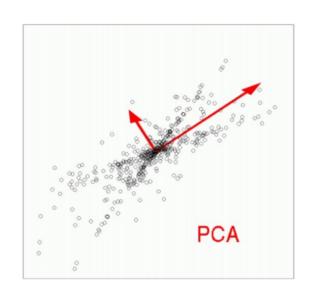
6) Interpreting the meaning of the factors

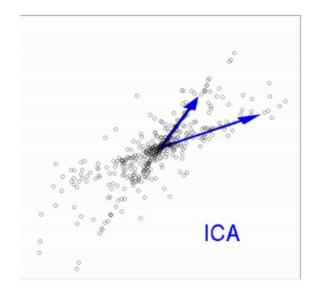
- visualizing temporal evolution of the first principal components
- using principal components to correlate with experimental observables
- classification and cluster analysis



population activity recorded in PFC [Mante et al. Nature 2013]

Other dimensionality reduction method: e.g. ICA





ICA – Independent Component Analysis

- tries to find independent components of data
- all components are equally important
- eigenvectors are not orthogonal