

Module 24:

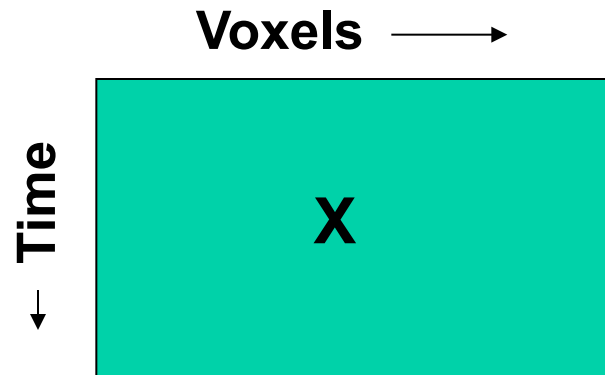
Multivariate Decomposition Methods

Decomposition Methods

- We often use **multivariate decomposition methods** to study functional connectivity.
 - Provides a decomposition of the data into separate components.
 - Can be used to find coherent brain networks.
 - Provides information on how different brain regions interact with one another.
- The most common decomposition methods are **principal components analysis** and **independent components analysis**.

Data Structure

- Throughout we organize the fMRI data in an $M \times N$ matrix **X**.
 - The row dimension is the number of time points and the column dimension the number of voxels.



Principal Components Analysis

- **Principal Components Analysis** (PCA) is a multivariate procedure concerned with explaining the variance-covariance structure of a high dimensional random vector.
- In PCA, a set of **correlated** variables are transformed into a set of **uncorrelated** variables, ordered by the amount of variability in the data that they explain.

Principal Components Analysis

- In fMRI **principal components analysis** involves finding spatial modes, or **eigenimages**, in the data.
 - These are the patterns that account for most of the variance-covariance structure in the data.
 - They are ranked in order of the amount of variation they explain.
- The eigenimages can be obtained using **singular value decomposition (SVD)**, which decomposes the data into two sets of orthogonal vectors that correspond to patterns in space and time.

Singular Value Decomposition

- Singular value decomposition (SVD) is an operation that decomposes a matrix \mathbf{X} as:

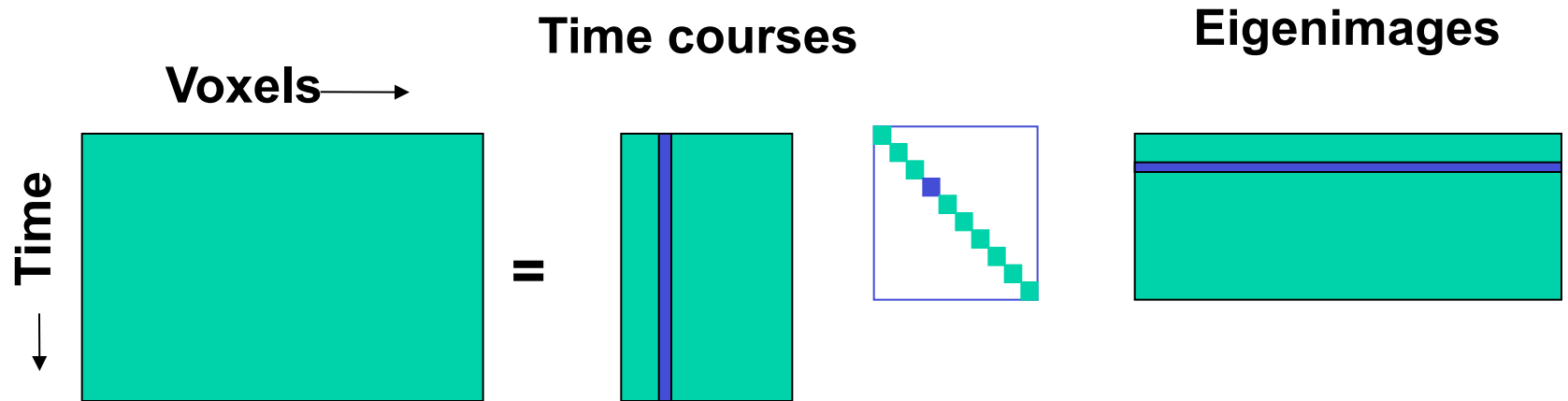
$$\mathbf{X} = \mathbf{U}\mathbf{S}\mathbf{V}^T$$

where

$$\mathbf{V}^T \mathbf{V} = \mathbf{I}$$

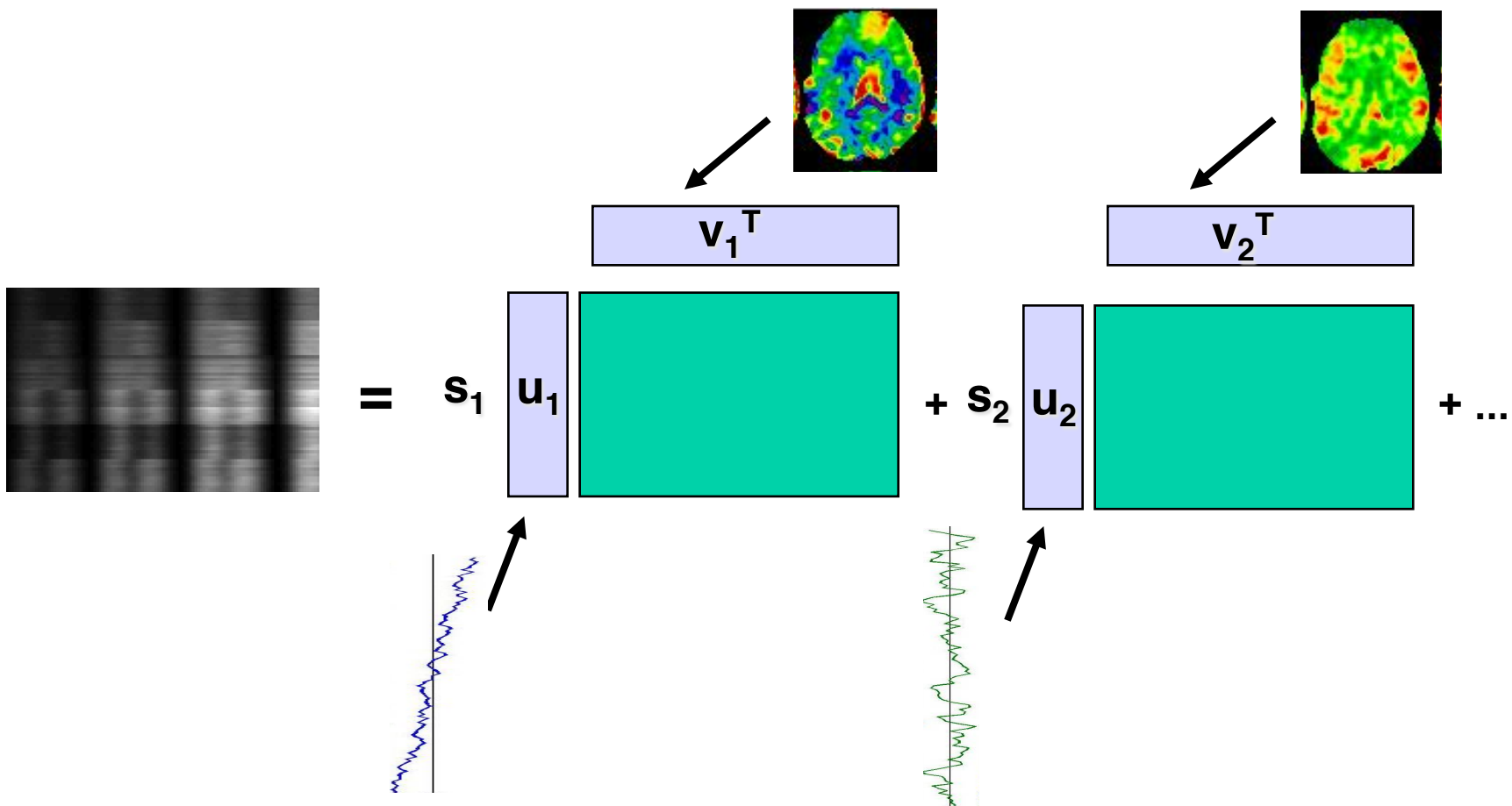
$$\mathbf{U}^T \mathbf{U} = \mathbf{I}$$

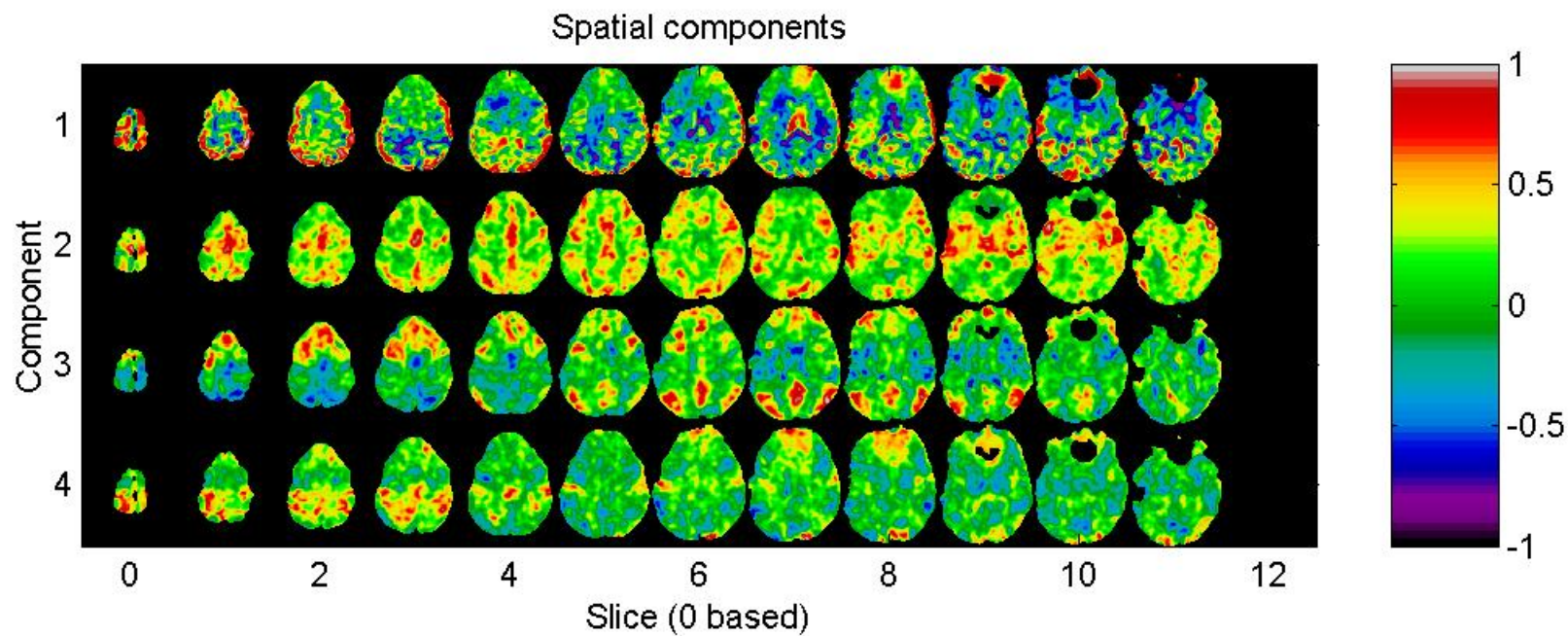
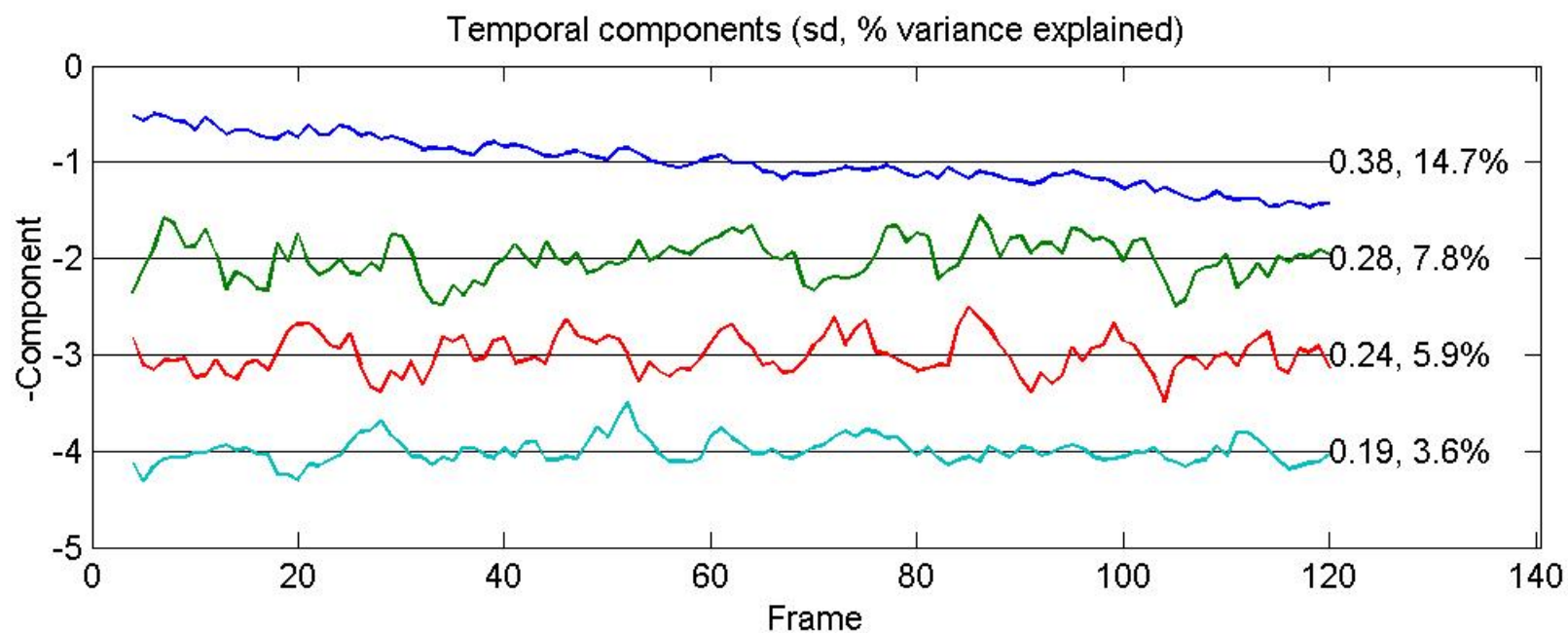
and \mathbf{S} is a diagonal matrix whose elements are called singular values.



$$\mathbf{X} = \mathbf{U}\mathbf{S}\mathbf{V}^T$$

$$\mathbf{X} = s_1 \mathbf{u}_1 \mathbf{v}_1^T + s_2 \mathbf{u}_2 \mathbf{v}_2^T + \dots + s_N \mathbf{u}_N \mathbf{v}_N^T$$





Independent Components Analysis

- Independent Components Analysis (ICA) is a family of techniques used to extract independent signals from some source signal.
- ICA provides a method to blindly separate the data into spatially independent components.
- The key assumption is that the data set consists of p spatially independent components, which are linearly mixed and spatially fixed.

Independent Components Analysis

- The ICA Model:

$$\mathbf{X} = \mathbf{AS}$$

- Here \mathbf{A} is referred to as the **mixing matrix** and \mathbf{S} as the **source matrix**.
- Our goal is to find an **un-mixing matrix** \mathbf{W} such that

$$\mathbf{Y} = \mathbf{WX}$$

provides a good approximation to \mathbf{S} .

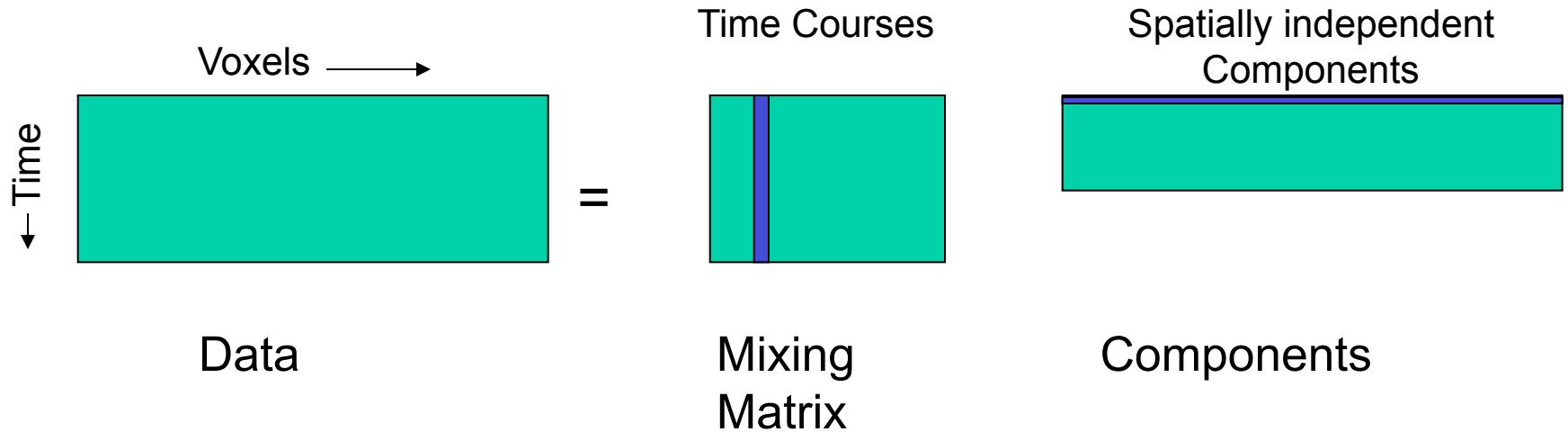
Assumptions

- If the mixing matrix is known, the problem is straight forward.
- However, ICA solves this problem **without** knowing the mixing parameters.
- Instead it exploits some key assumptions:
 - Linear mixing of sources.
 - The components s_i are statistically independent.
 - The components s_i are non-Gaussian.

ICA for fMRI

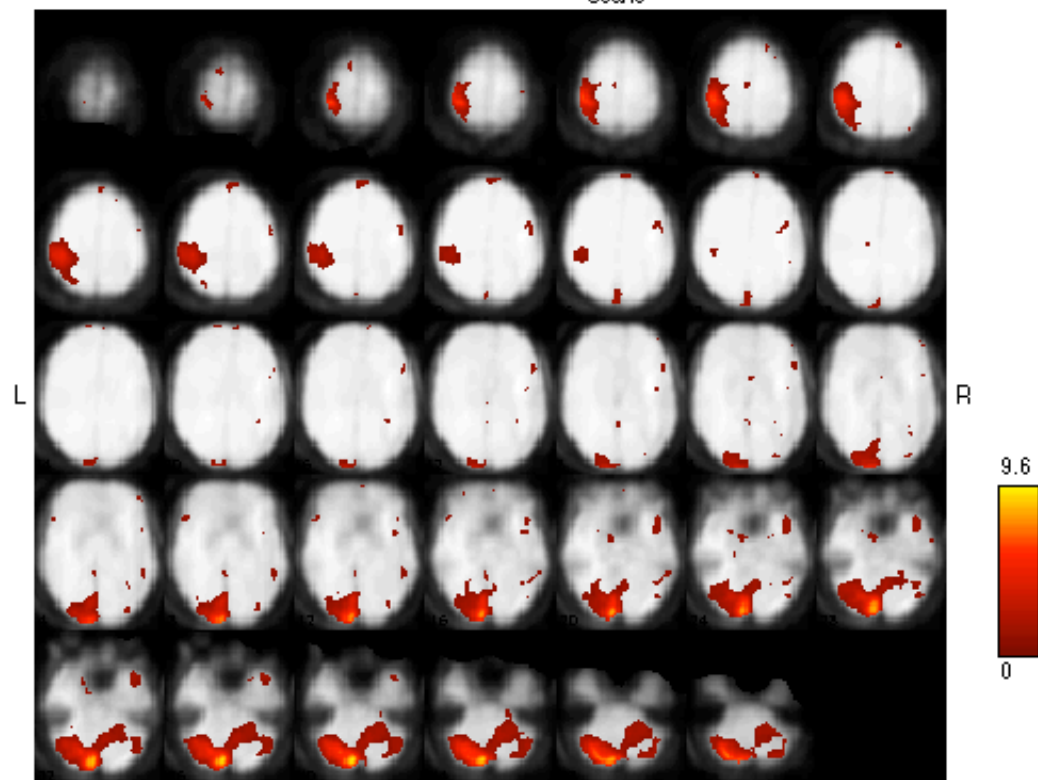
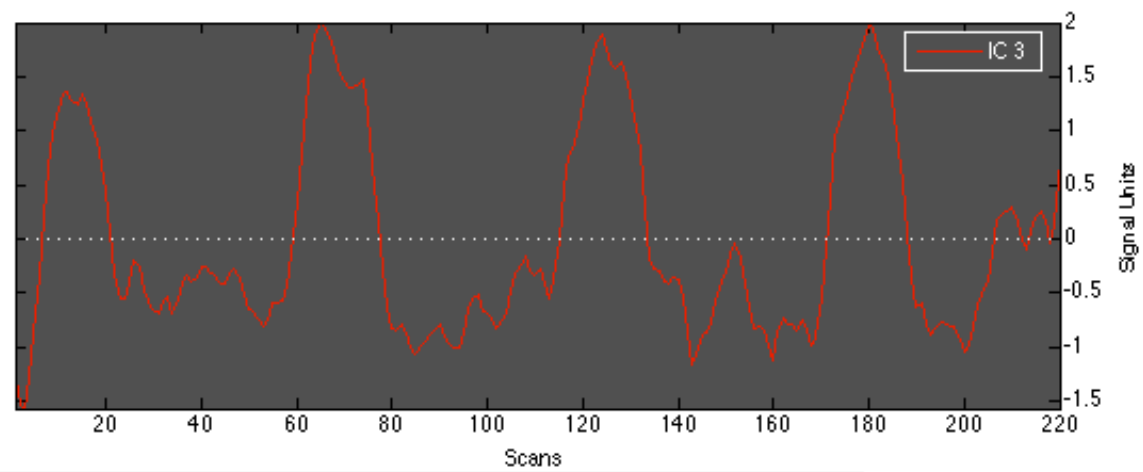
- It is assumed that the fMRI data can be modeled by identifying sets of voxels whose activity both vary together over time and are different from the activity in other sets.
- Decompose the data set into a set of **spatially independent** component maps with a set of corresponding time-courses.

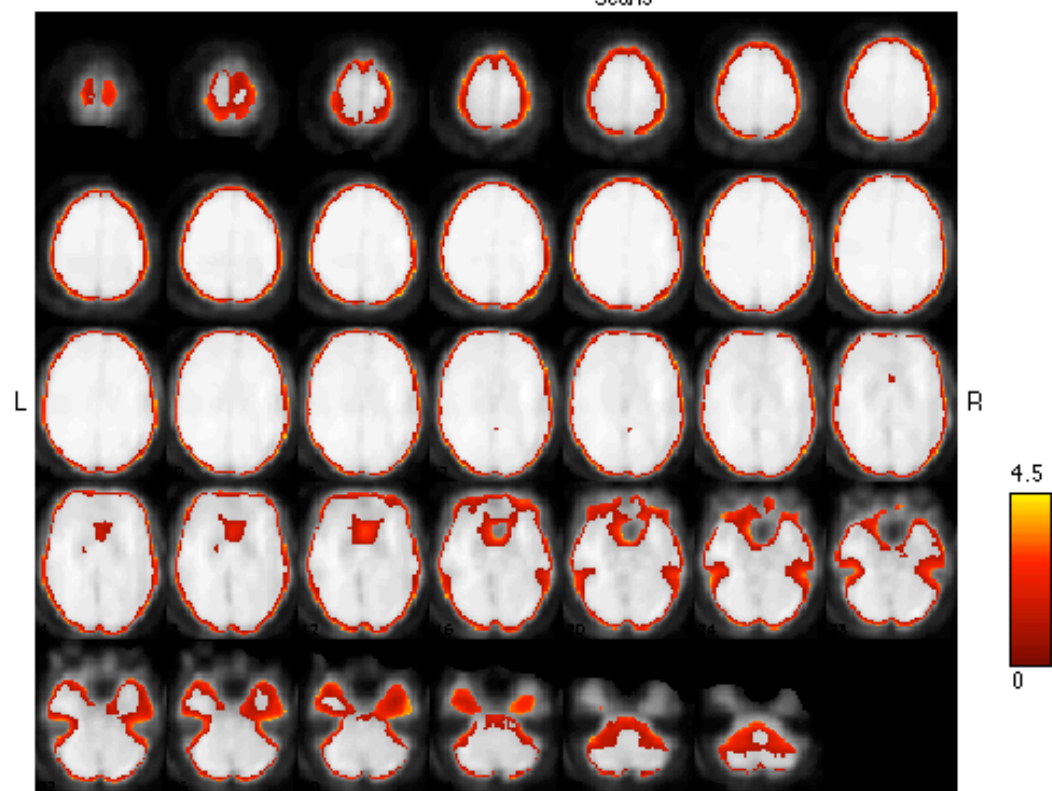
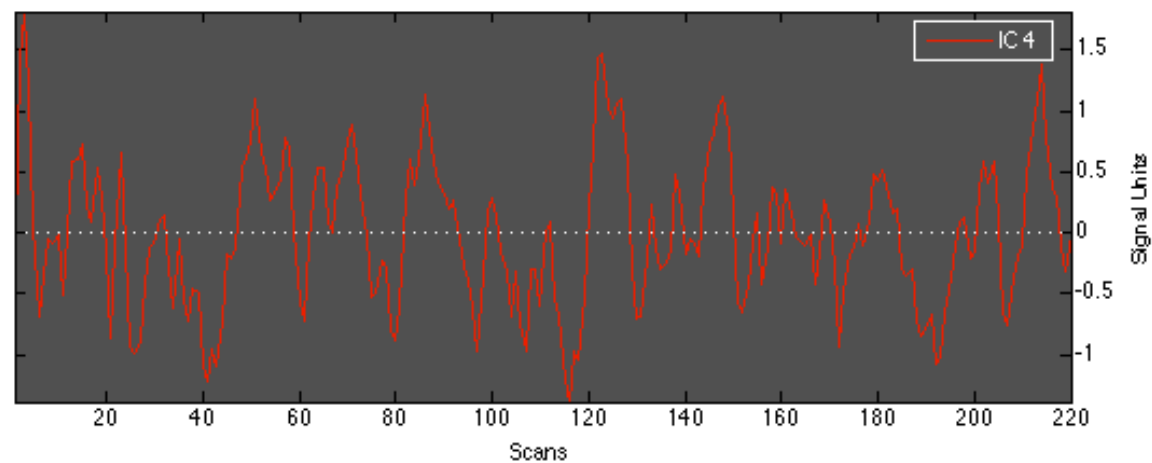
Overview



$$X = AS$$

Use an ICA algorithm to find A and S.





Comments

- Unlike PCA which assumes an orthonormality constraint, ICA assumes statistical independence among a collection of spatial patterns.
 - Independence is a stronger requirement than orthonormality.
- However, in ICA the spatially independent components are not ranked in order of importance as they are when performing PCA.

End of Module



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