

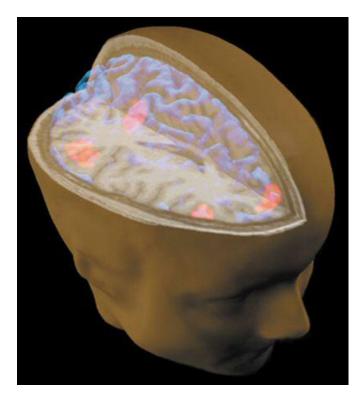




Introduction to fMRI Data Analysis

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Research Interests

- Functional Neuroimaging (fMRI)
- MRI Physics & Pulse Sequence Developments
- Medical Signal & Image Processing
- Pattern Recognition & Neural Networks







Outline

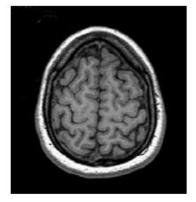
- Introduction
- Preprocessing
- Data Analysis
- Result Representation

sMRI & fMRI

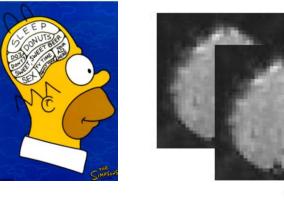
- sMRI studies brain anatomy.
 - High spatial resolution
 - Can distinguish different types of tissue

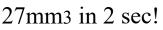
- fMRI studies brain function.
 - Functional images
 - Lower spatial resolution/ Higher temporal resolution





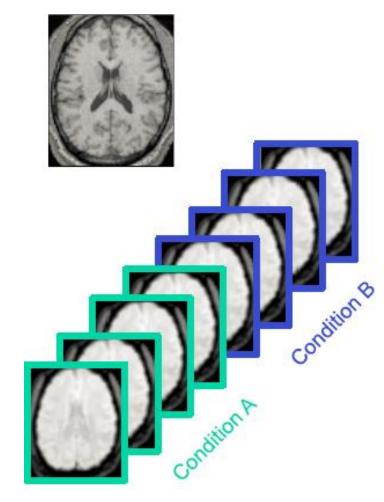
1mm3 in 4 min!





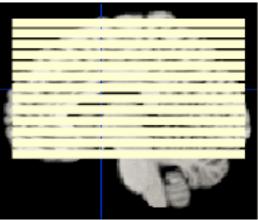
fMRI Data Acquisition

- Structural (T1) images:
 - High spatial resolution
 - Low temporal resolution
 - Can distinguish different types of tissue
- Functional (T2*) images:
 - Lower spatial resolution
 - Higher temporal resolution
 - Can relate changes in signal to an experimental manipulation



Terminology

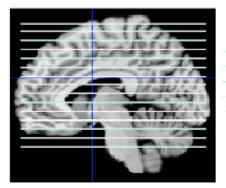
- MRI images are typically acquired in axial slices one at a time.
- This can be performed in either a sequential or interleaved manner.
- Together the slices make up a 3 dimensional brain volume.



Terminology

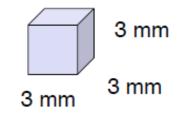
- Field of View (FOV)
 - (e.g. 192 mm)
- Matrix Size
 - (e.g., 64 x 64)
- In-plane resolution
 - 192 mm / 64 = 3 mm
- Slice thickness
 - (e.g., 3 mm)
- Voxel Size

• 3*3*3 mm





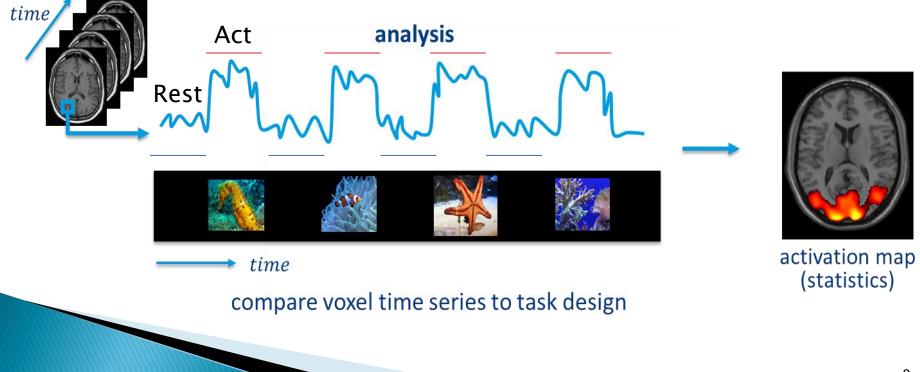




64*64*35 Voxels

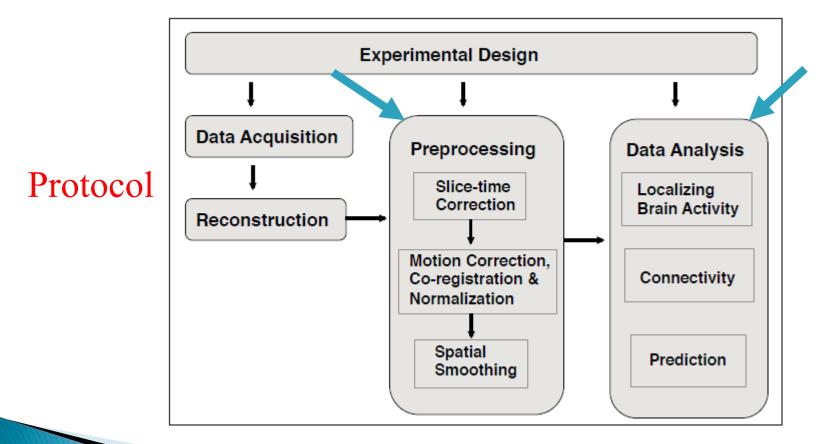
fMRI Experiment

 An fMRI experiment consists of a sequence of individual MR images, where one can study oxygenation changes in the brain across time



Pipeline

Task



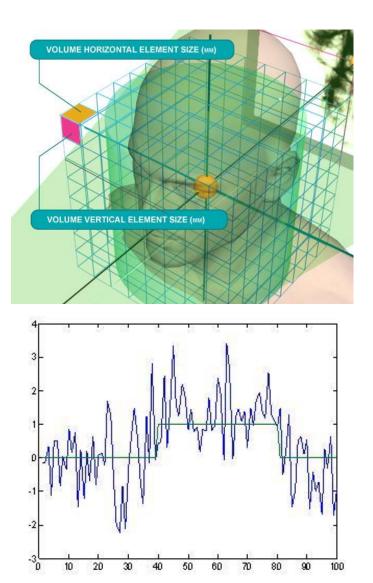
Processing

Pre-processing

- Prior to analysis, fMRI data undergoes a series of preprocessing steps aimed at identifying and removing artifacts and validating model assumptions.
- The goals of preprocessing are
 - To minimize the influence of data acquisition and physiological artifacts;
 - To check statistical assumptions and transform the data to meet assumptions;
 - To standardize the locations of brain regions across subjects to achieve validity and sensitivity in group analysis.

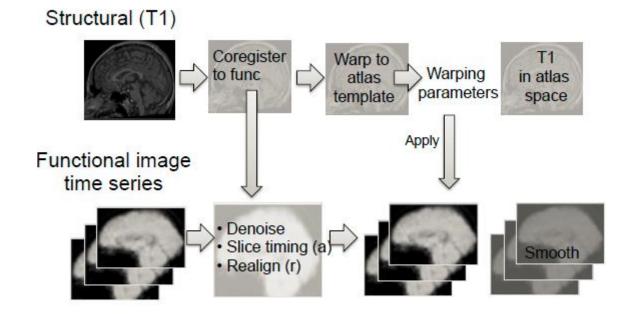
fMRI Noise

- Sources of noise
 - Thermal motion of free electrons in the system.
 - Low frequency signal drift.
 - Patient movement during the experiment.
 - Physiological effects: subject's heartbeat and respiration.



Pre-processing Pipeline

• Preprocessing is performed both on the fMRI data and structural scans collected prior to the experiment.

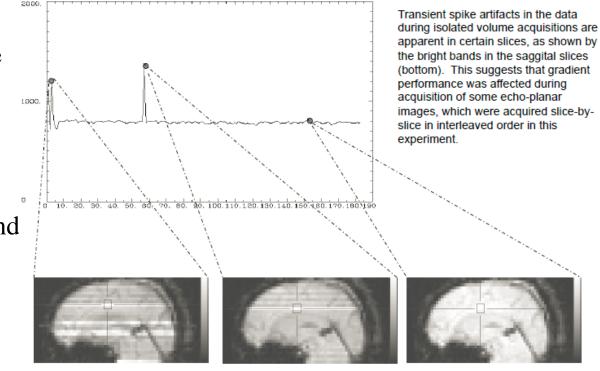


Pre-processing Steps

- Visualization and Artifact Removal
- Slice Time Correction
- Motion Correction
- Physiological Corrections
- Co-registration
- Normalization
- Spatial Filtering
- Temporal Filtering

Visualization & Artifact Removal

- Investigate the raw image data and detect possible problems and artifacts.
- fMRI data often contain transient spike artifacts and slow drift over time.



Easy but Important!

Slice Time Correction

- Almost all FMRI scanning takes each slice separately
- Each slice is scanned at a slightly different time
- Slice order can be interleaved or sequential (up or down)



Slice Time Correction

Temporal Interpolation

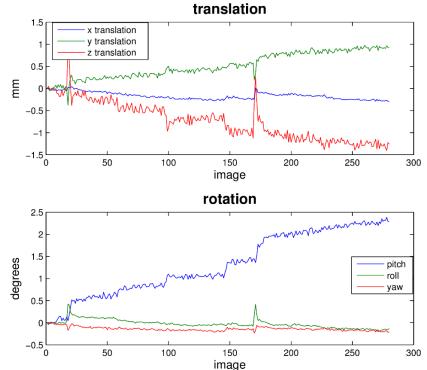
• Use information from nearby time points to estimate the amplitude of the MR signal at the onset of the TR."

Phase Shift

• Slide the time course by applying a phase shift to the Fourier transform of the time course.

Head Motion

- Very small movements of the head during an experiment can be a major source of error if not treated correctly.
- When analyzing the time series associated with a voxel, we assume that it depicts the same region of the brain at every time point
 - Head motion may make this assumption incorrect.
- Can be corrected using a rigid body transformation.



Head Motion

- The goal is to find the best possible alignment between an input image and some target images.
- To align the two images, one of them needs to be transformed.
- A rigid body transformation is used.
- It involves 6 variable parameters, 3 sets of translations and 3 sets of rotations (6 DOF).

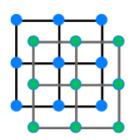
Linear transformations

- Rigid body (6 DOF) translation and rotation
- Similarity (7 DOF) translation, rotation and a single global scaling
- Affine (12 DOF) translation, rotation, scaling and shearing.

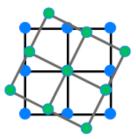
Warping

• Transformations where the equations relating the coordinates of the images are non-linear.

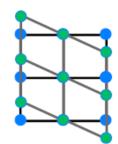
Translation



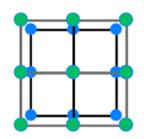
Rotation



Shearing

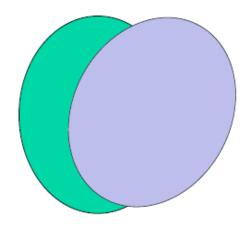






Head Motion

- The target image is usually defined to be the first (or mean) image in the fMRI time series.
- The goal is to find the set of parameters which minimizes some cost function that assesses similarity between the image and the target.
- Examples of cost functions include the sum of squared differences or mutual information.

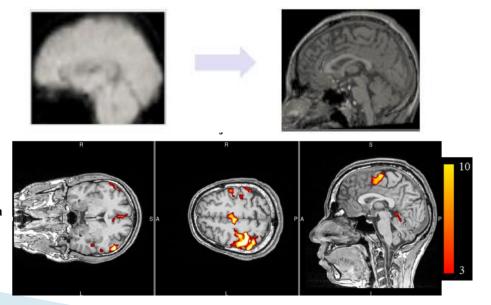


Co-registration

• A structural MRI collected in the beginning of the session is registered to the fMRI images in a process referred to as coregistration.

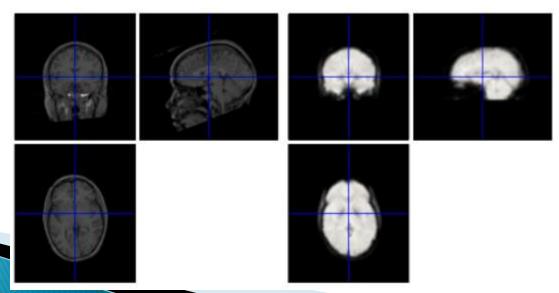
– Allows one to visualize single-subject task activations overlaid on the individual's anatomical information.

– Simplifies later transformation of the fMRI images to a standard coordinate system.



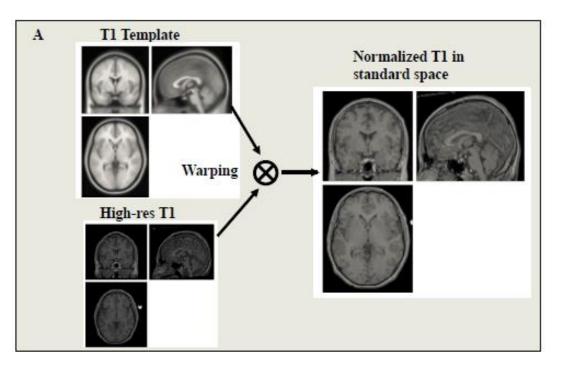
Activation Map

- There are certain key differences between coregistration and motion correction.
 - Functional and structural images do not have the same signal intensity in the same areas.
 - They cannot be subtracted.
 - Their shapes may differ!
- Use at least an affine transformation (12 DOF).



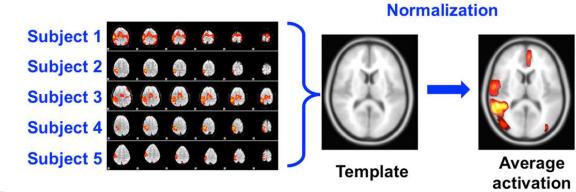
Normalization

- Normalization allows one to stretch, squeeze and warp each brain so that it is the same as some standard brain.
- The structural MR image used in the co-registration procedure is warped onto a template image.



Normalization

- Pros
 - Spatial locations can be reported and interpreted in a consistent manner.
 - Results can be generalized to larger population
 - Results can be compared across studies
 - Results can be averaged across subjects
- Cons
 - Reduces spatial resolution.
 - Introduces potential errors.



Brain Atlases

- Montreal Neurological Institute (MNI)
 - Combination of many MRI scans on normal controls (152 in current standard).
 - All right-handed subjects.
 - More representative of population.



Spatial and Temporal Filtering

- In fMRI it is common to spatially smooth the acquired data prior to statistical analysis.
- Can increase signal-to-noise, validate distributional assumptions and remove artifacts.





Pros

- May overcome limitations in the normalization by blurring any residual anatomical differences.
- Could increase the signal-to-noise ratio (SNR).
- May increase the validity of the statistical analysis.
- Cons
 - The image resolution is reduced.

Contents lists available at ScienceDirect



Journal of Neuroscience Methods

journal homepage: www.elsevier.com/locate/jneumeth

Efficient de-noising of high-resolution fMRI using local and sub-band information



NEUROSCIENCE Methods

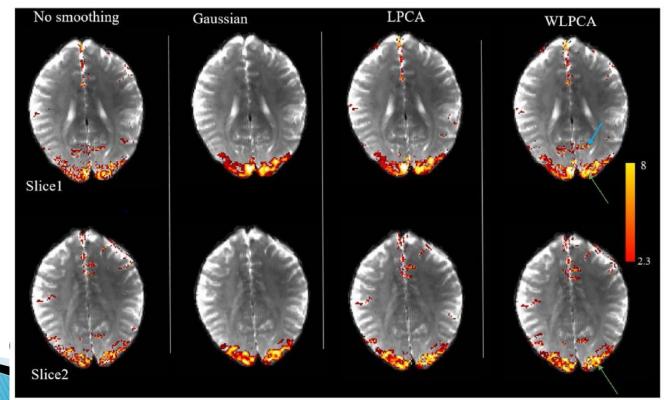
Vahid Malekian^{a,b}, Abbas Nasiraei-Moghaddam^{a,b,*}, Amir Akhavan^c, Gholam-Ali Hossein-Zadeh^{b,d}

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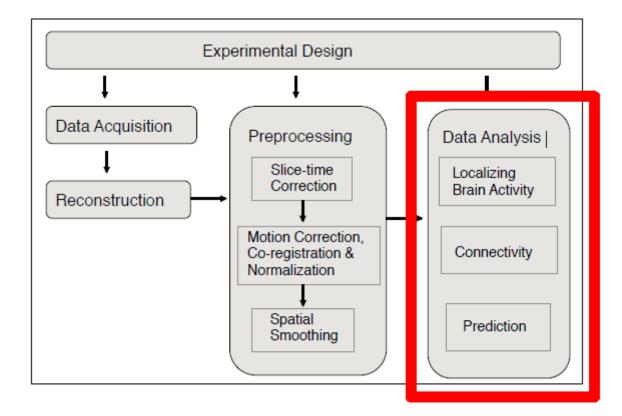
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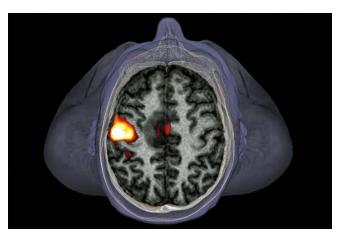


Data Analysis



Statistical Analysis

- There are multiple goals in the statistical analysis of fMRI data.
- They include:
 - localizing brain areas activated by the task;
 - determining networks corresponding to brain function;



Introducing The GLM

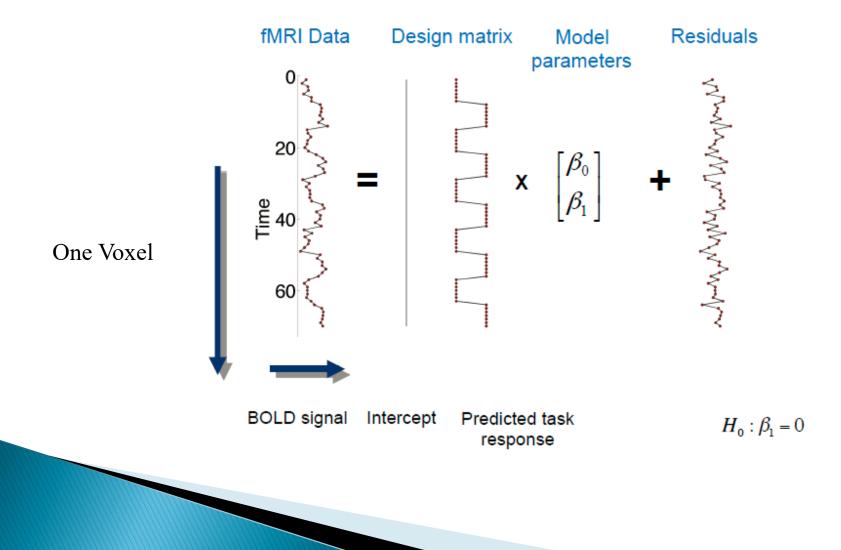
DATA = MODEL + ERROR DATA = KNOWN * UNKNOWN + ERROR

- Where Y is a matrix with a series of observed measurements
- Where X is a matrix that might be a design matrix
- Where **b** is a matrix containing parameters to be estimated
- And ε is a matrix containing error or noise

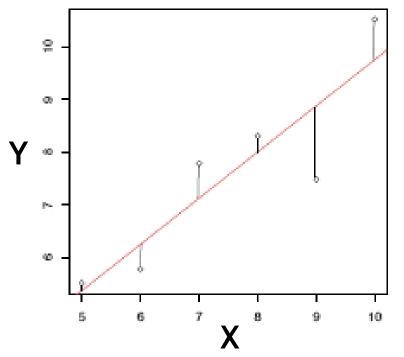
The model functions are assumed to have known shapes, but their amplitudes are unknown and need to be estimated.

GLM

 $Y = X.\beta + \varepsilon$



GLM: Simple Linear Regression



$\mathbf{Y} = \mathbf{\beta}_0 + \mathbf{X}_1 \mathbf{\beta}_1 + \mathbf{\varepsilon}$

 $\beta_{0:}$ is the Y axis intercept

 $\beta_{1:}$ is the gradient of slope

Y: the black circles

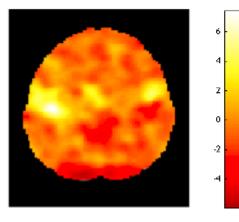
E: diff between Y and observed Y

• This is done by choosing β_0 and β_1 so that the sum of the squares of the estimated errors $\sum \epsilon_i^2$ is as small as possible. Method of Least Squares

Statistical Images

- Calculate β for each voxel time-series.
- Perform a statistical test to determine whether task related activation is present in the voxel.

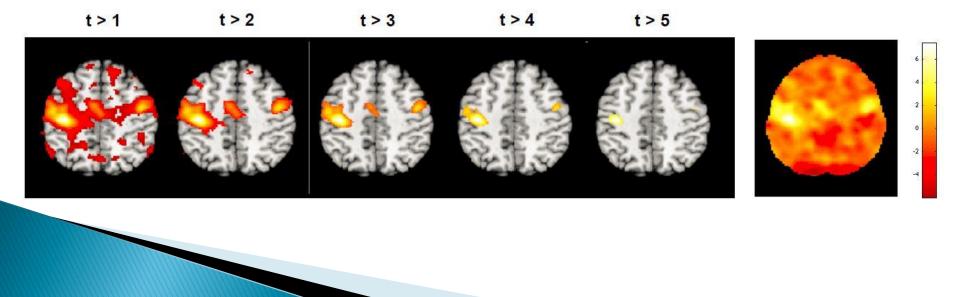
$$H_0: \mathbf{c}^T \boldsymbol{\beta} = \mathbf{0}$$



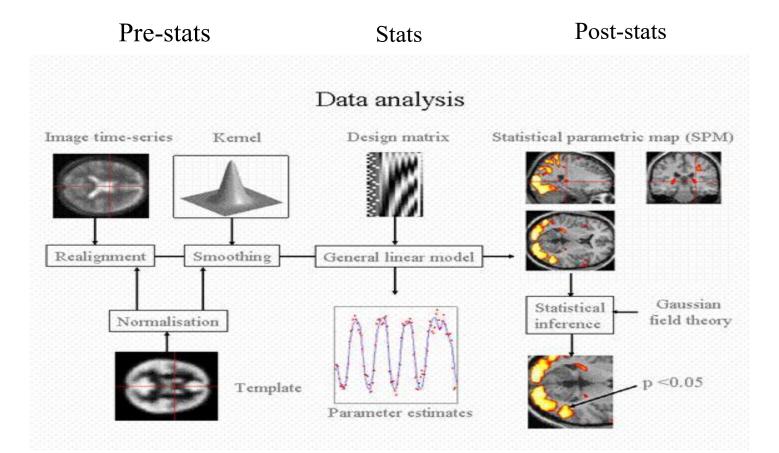
Statistical image: Map of t-tests across all voxels (a.k.a t-map).

Thresholding

- Choose an appropriate threshold for determining statistical significance.
- Choosing a threshold is a balance between sensitivity (true positive rate) and specificity (true negative rate).



So Far ...



Software...

- FSL (Oxford)
- > SPM (UCL)



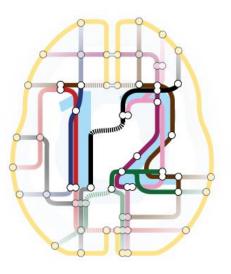


- Free-surfer (MGH)
- AFNI (NIH)





- CONN
- Brain Voyager, Brain Suite and ...



Take home message

- Be aware of your data acquisition protocol. Learning MRI could help!
- Look at your data before start!
- Motion is WORST: When in doubt, throw it out!
- Choosing a threshold is a balance. Take care of it!
- Pre and Post processing are not always the "one size fits all" practice in fMRI studies!

A journey of a thousand miles must begin with a single step.

Lao Zi

Thank you ...

