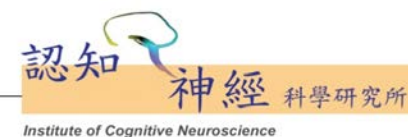


TMBIC 2017 資料分析助理研習營

# FMRI實驗設計與資料前處理

FMRI Experimental Designs and Data Preprocessing

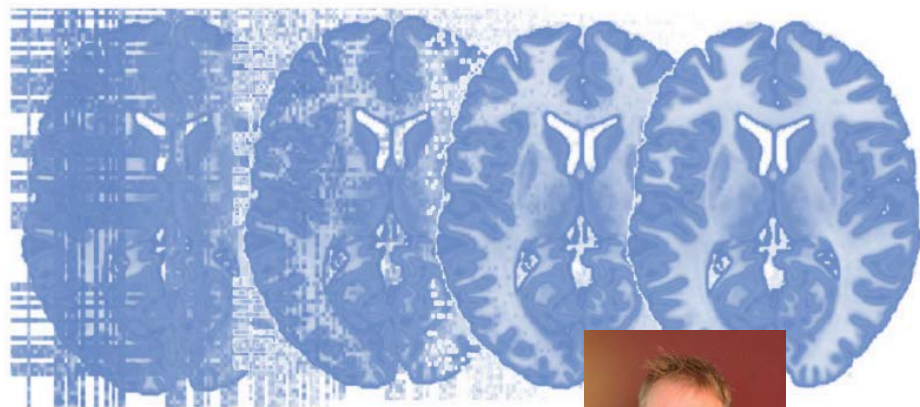
張智宏副教授 中央大學認知神經科學研究所



National Central University, Taiwan

# 參考書籍

## Principles of fMRI

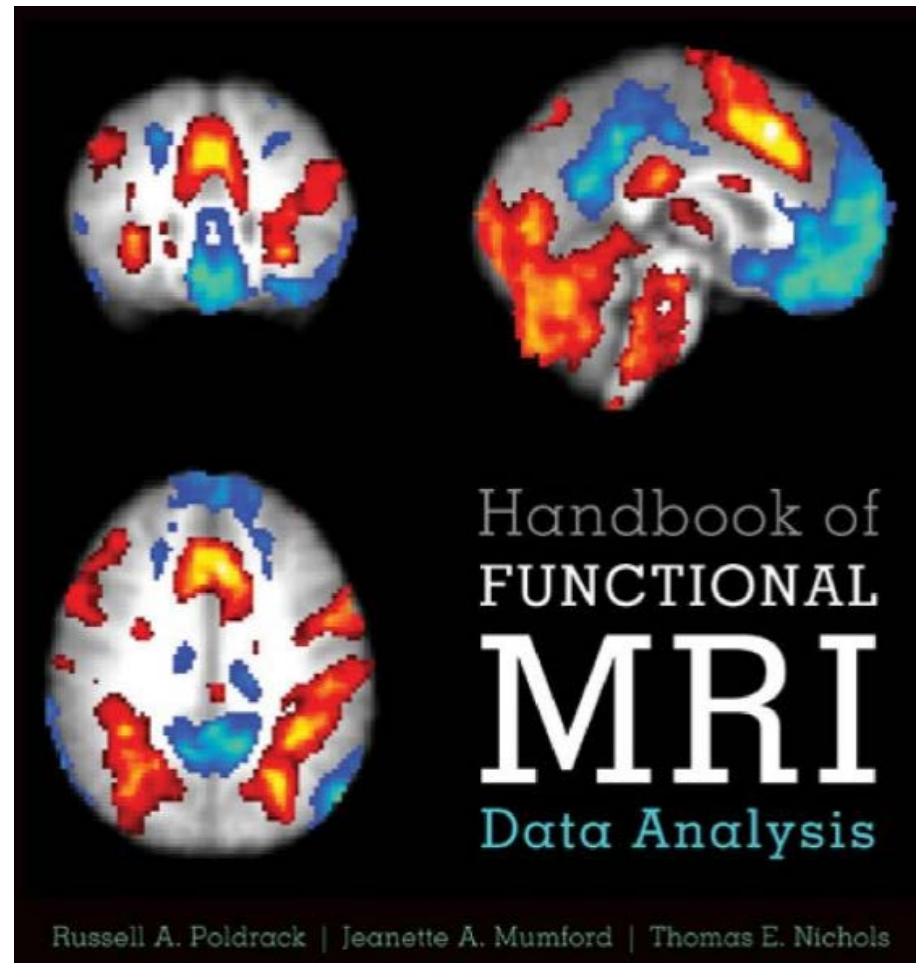


Martin A. Lindquist, Ph.D.



Tor D. Wager, Ph.D.

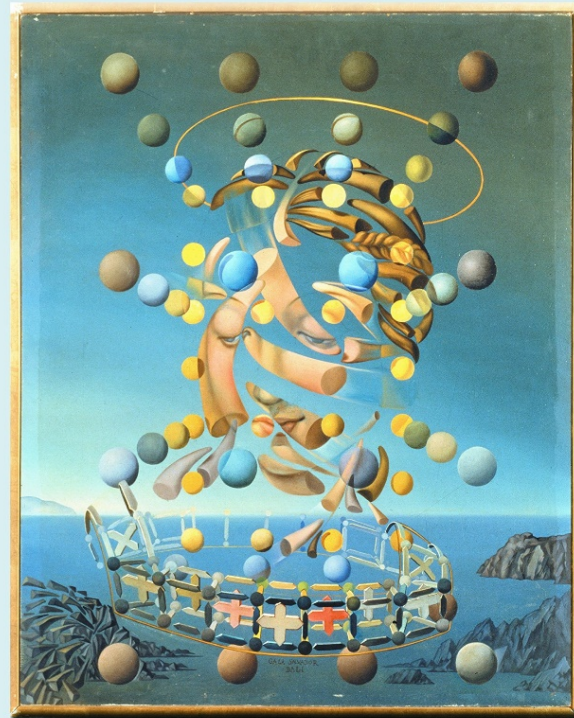
<https://leanpub.com/principlesoffmri>



<http://www.fmri-data-analysis.org>

# FUNCTIONAL Magnetic Resonance Imaging

Third Edition

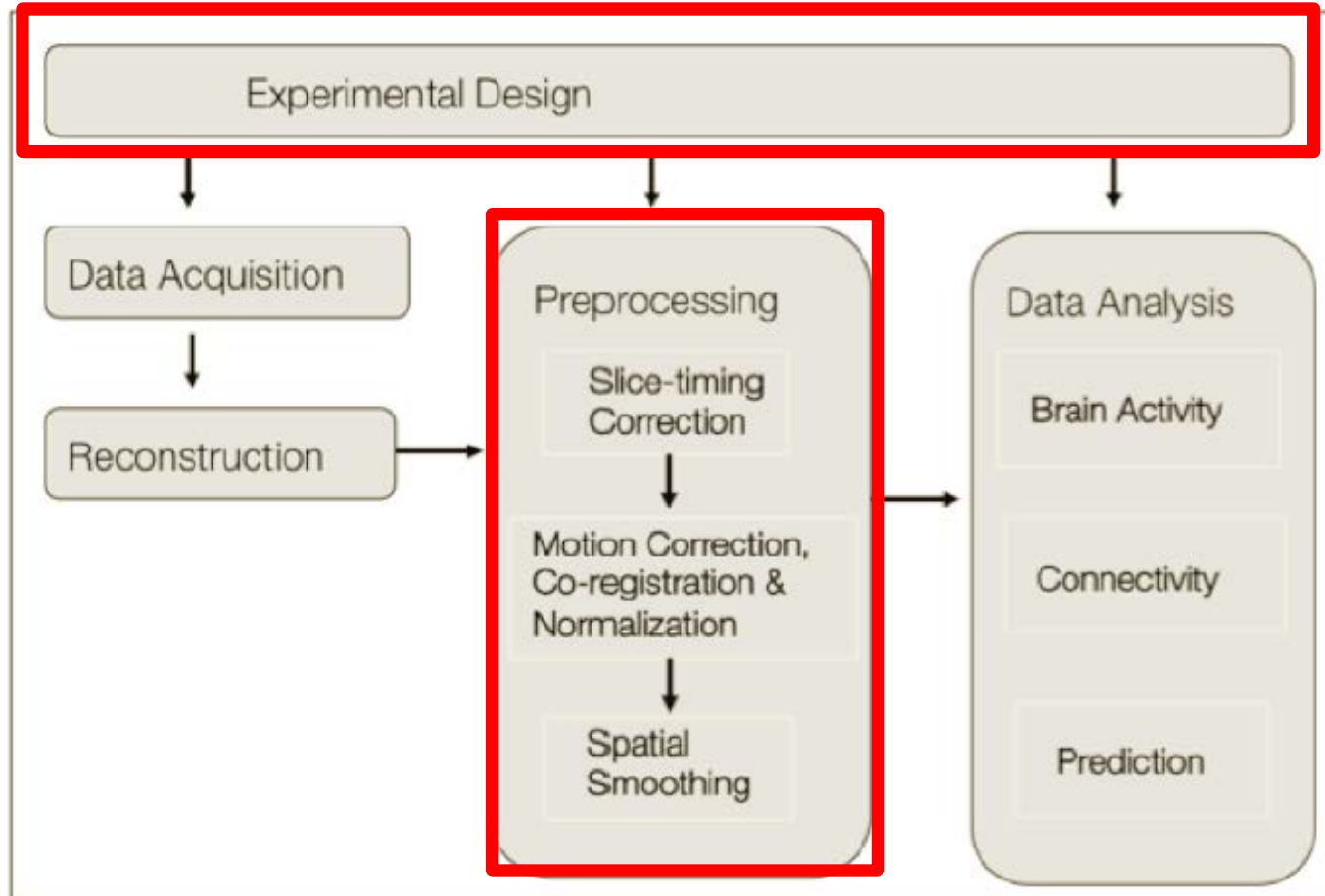


Scott A. Huettel • Allen W. Song • Gregory McCarthy



<http://www.sinauer.com/functional-magnetic-resonance-imaging-737.html>

# The Big Picture



Source: Lindquist & Tor (2015)

# 實驗基本概念

- 受控制的觀察

- 獨變項

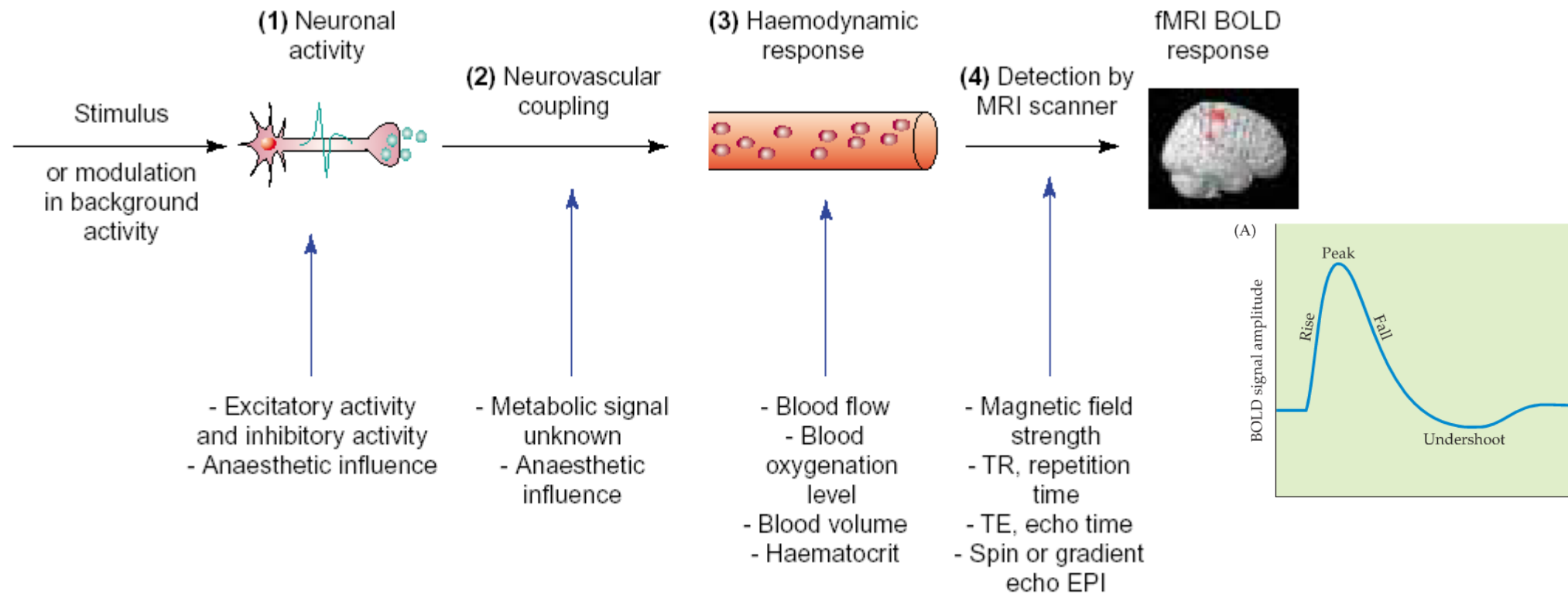
- ( Independent Variables; IV )

- 由研究者操弄
      - 至少兩個水準
      - 例：運動強度、心理壓力程度、圖形顏色、圖形類別

- 依變項(Dependent Variables, DV)

- 量化指標
  - 可受獨變項影響而改變
  - 例：心跳、壓力荷爾蒙、反應時間、**BOLD**

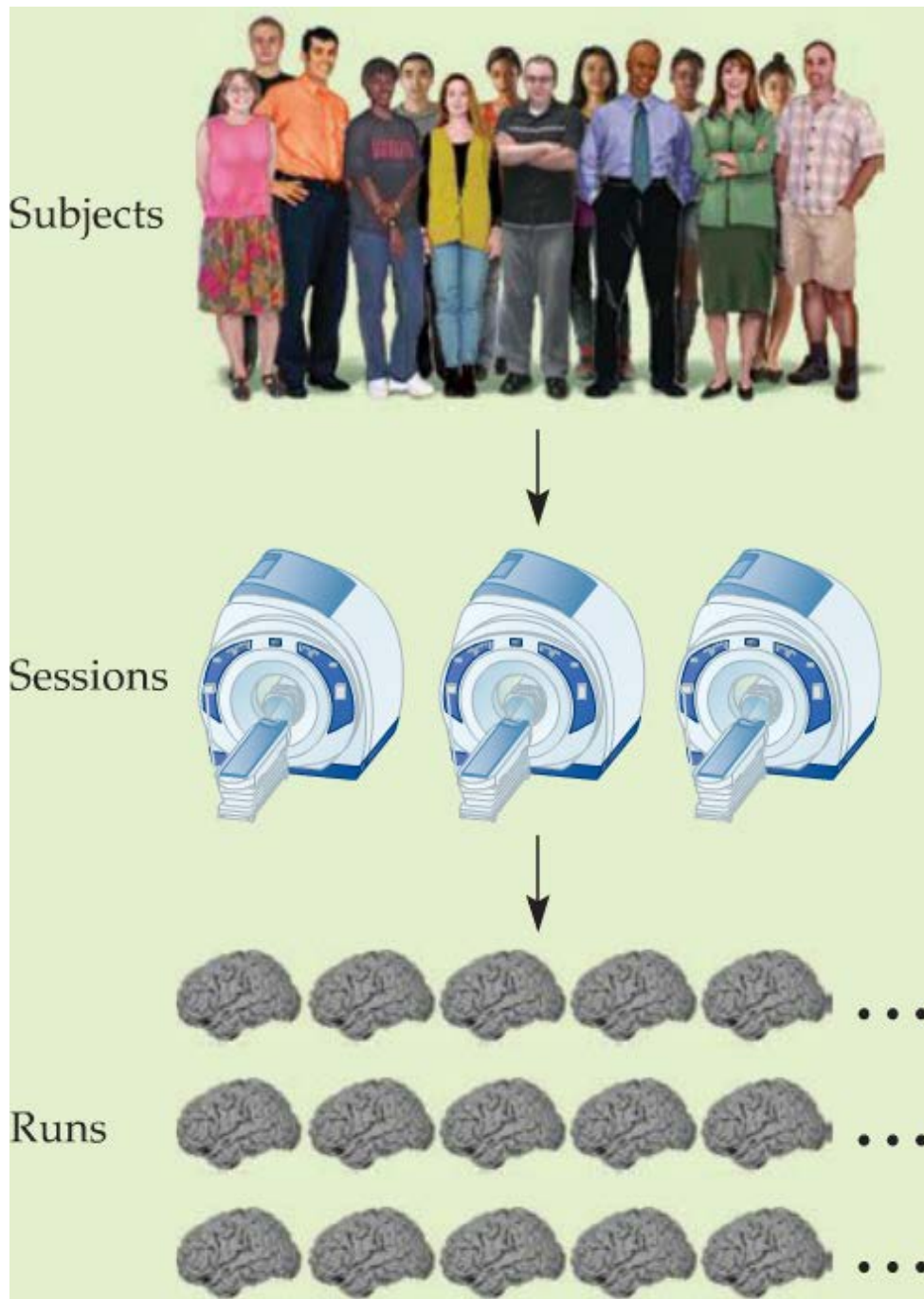
# 關於fMRI，我們觀察的是...



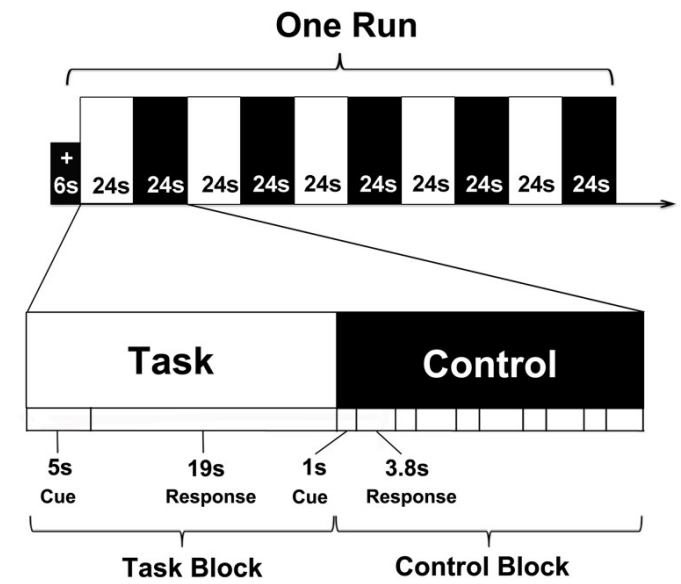
TRENDS in Neurosciences

Source: Arthurs & Boniface, 2002, *Trends in Neurosciences*

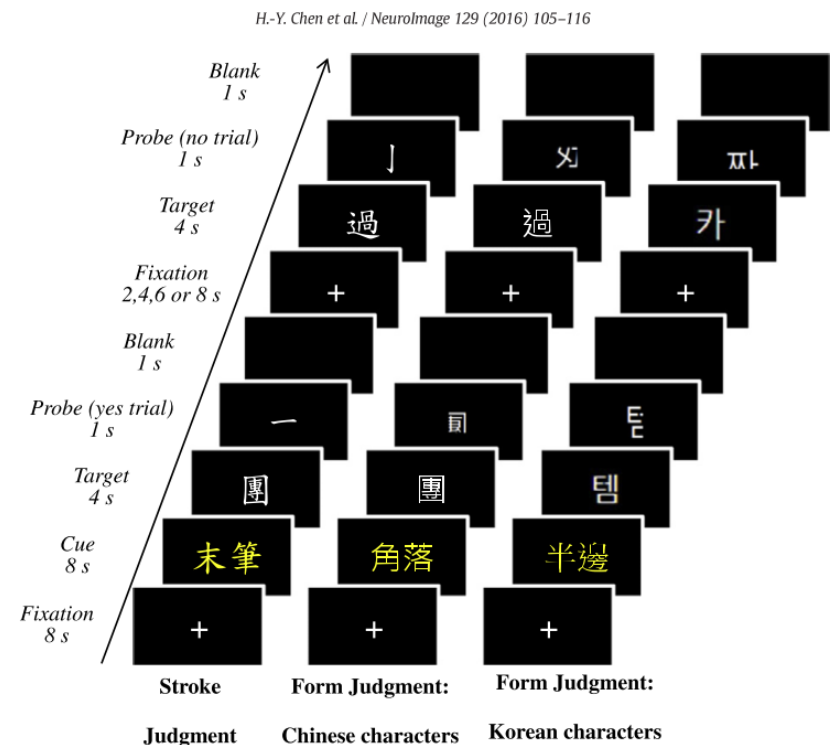




- Blocks
- Conditions



- Trials
- Events



(A)

B1

"Carrot" "Mailbox" "Knife" "Tiger" "Sweater" "Teapot" "Auto" "Doorbell" "Spider" "Parsley"

B2

"Plant" "Handbag" "Pebble" "Chess" "Book" "Phone" "Anger" "Watch" "Window" "Night"

(B) **Alternating Design**

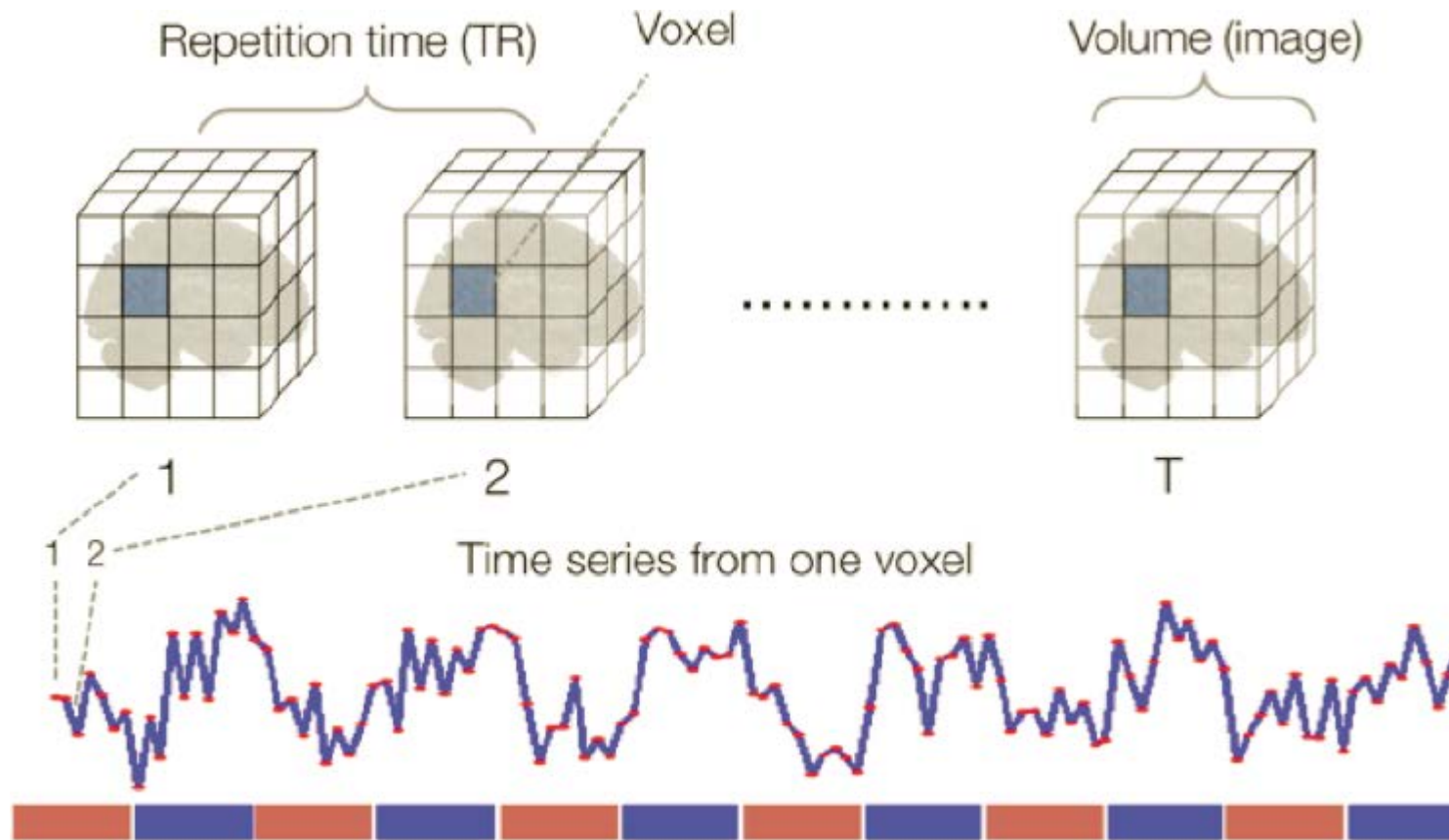


(C) **Interleaving null-task blocks**





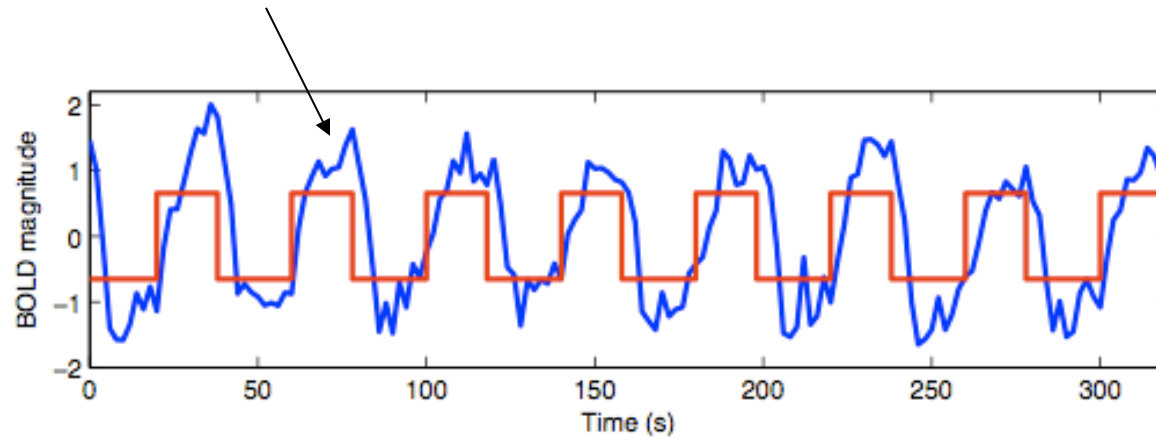
# FMRI Time Series



Source: Lindquist & Tor (2015)

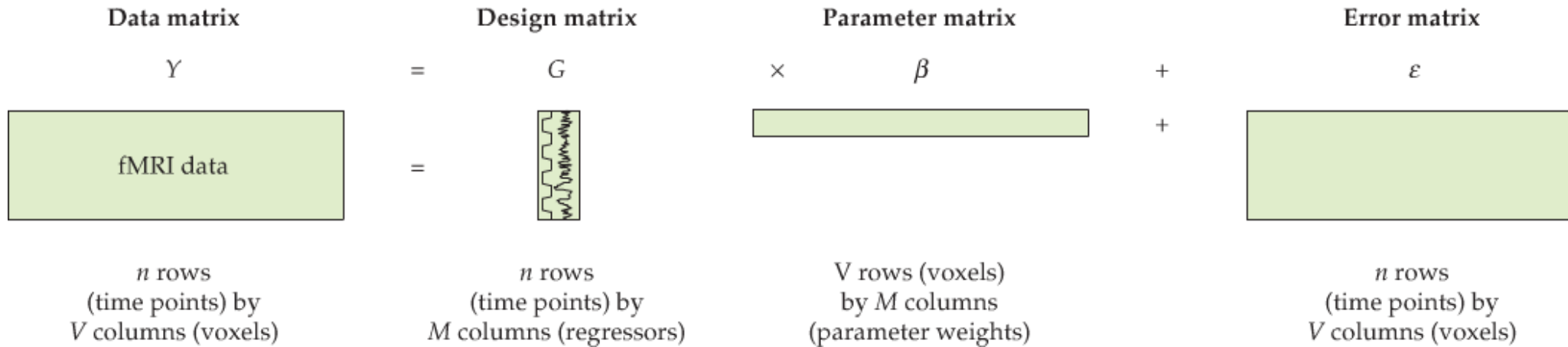
# IV and DV from the Perspective of Data Analysis

- Observed BOLD signal (example)



- BOLD signal did not look exactly like the predicted neural activity (in red)

# General Linear Model



# FMRI Experimental Designs

- Blocked designs
- Event-related designs
- Mixed designs

**Block  
Design**



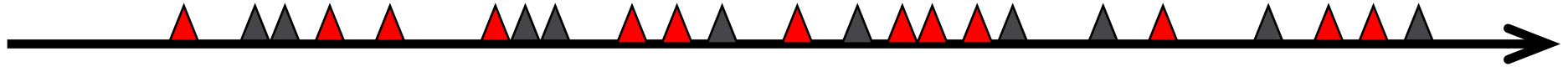
**Slow ER  
Design**



**Rapid  
Counterbalanced  
ER Design**



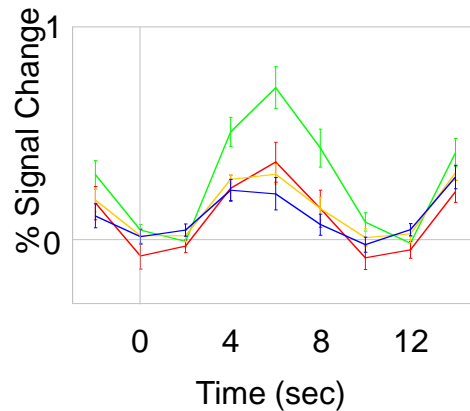
**Rapid  
Jittered ER  
Design**



**Mixed  
Design**



# Detection vs. Estimation



- Detection: determination of whether activity of a given voxel (or region) changes in response to the experimental manipulation
- “which voxel?”
  - Estimation: measurement of the time course within an active voxel in response to the experimental manipulation
  - “How does signal change in a voxel?”

*Definitions modified from: Huettel, Song & McCarthy, 2004, Functional Magnetic Resonance Imaging*

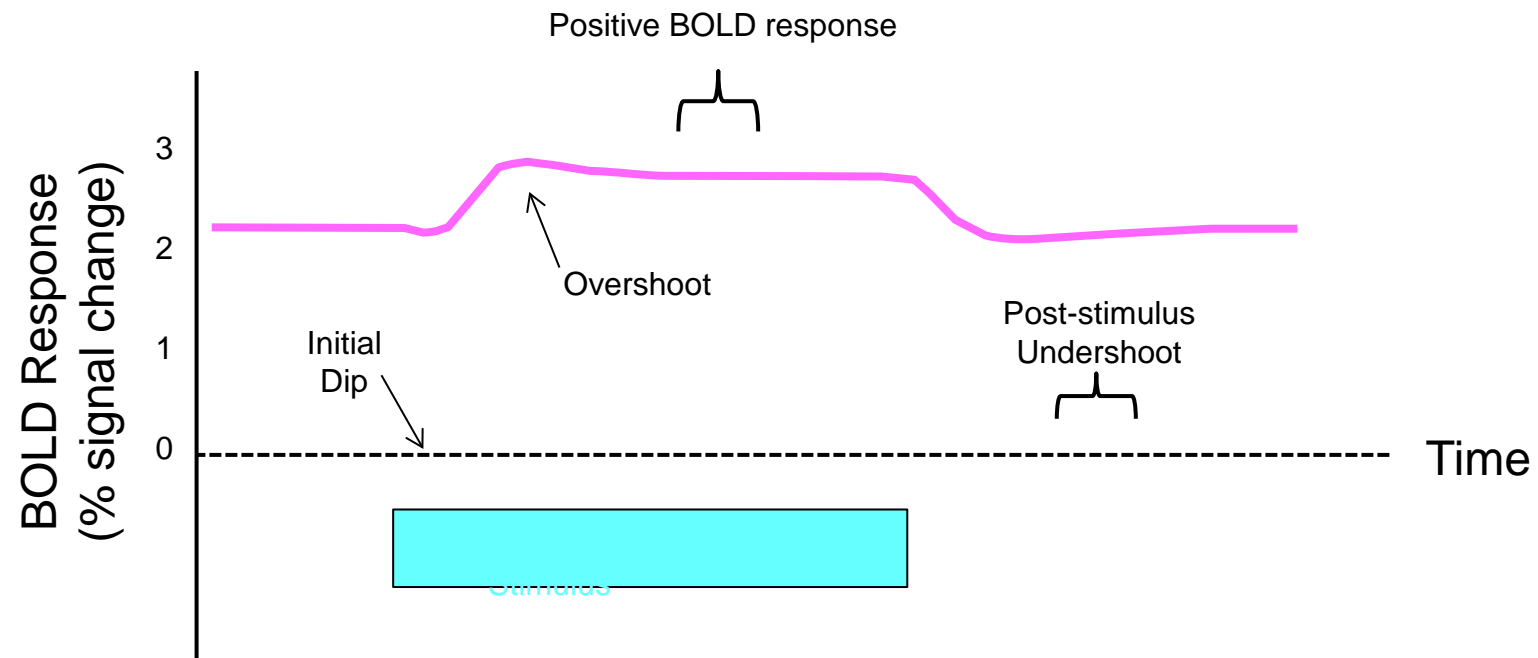


# Block Designs

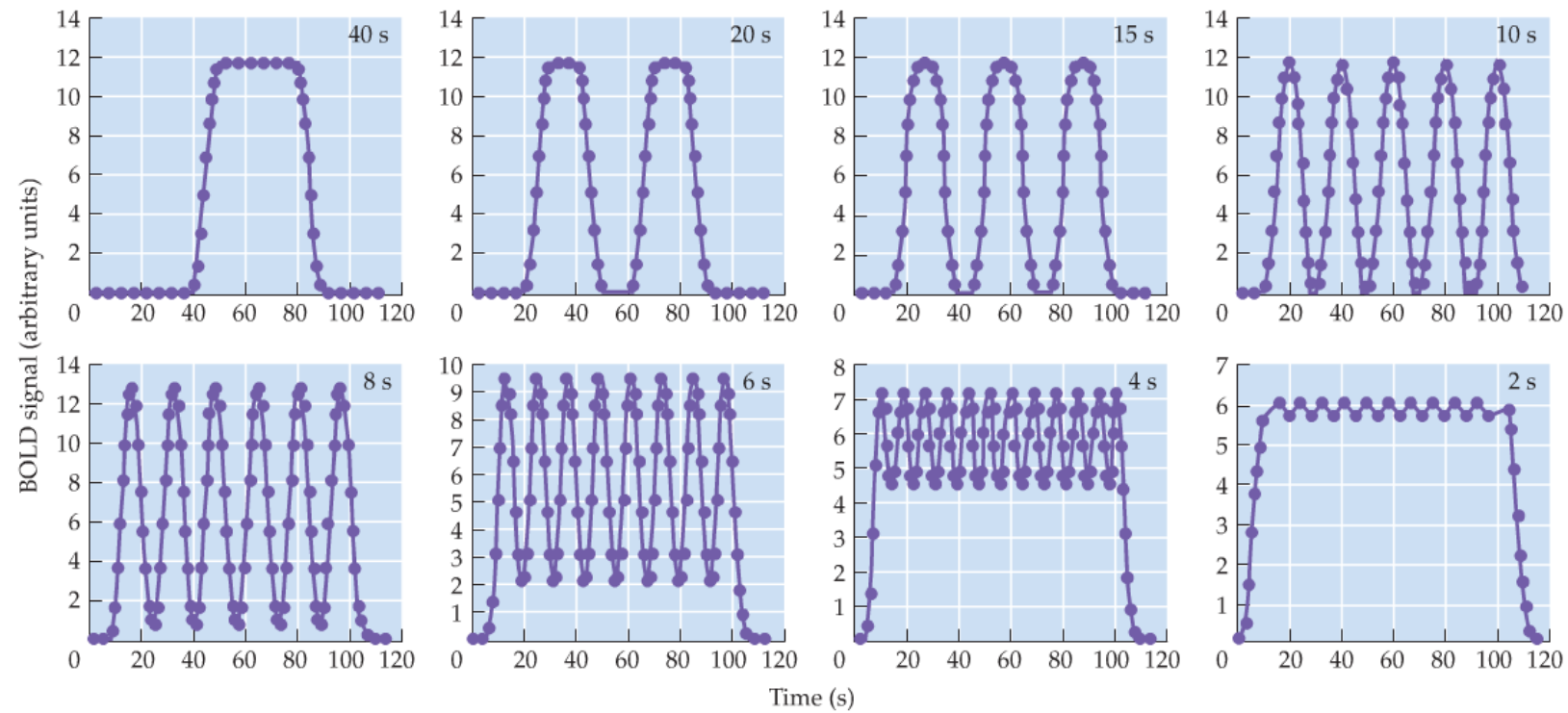
▲ = trial of one type (e.g., face image)    ▲ = trial of another type (e.g., place image)



Early Assumption: Because the hemodynamic response delays and blurs the response to activation, the temporal resolution of fMRI is limited.



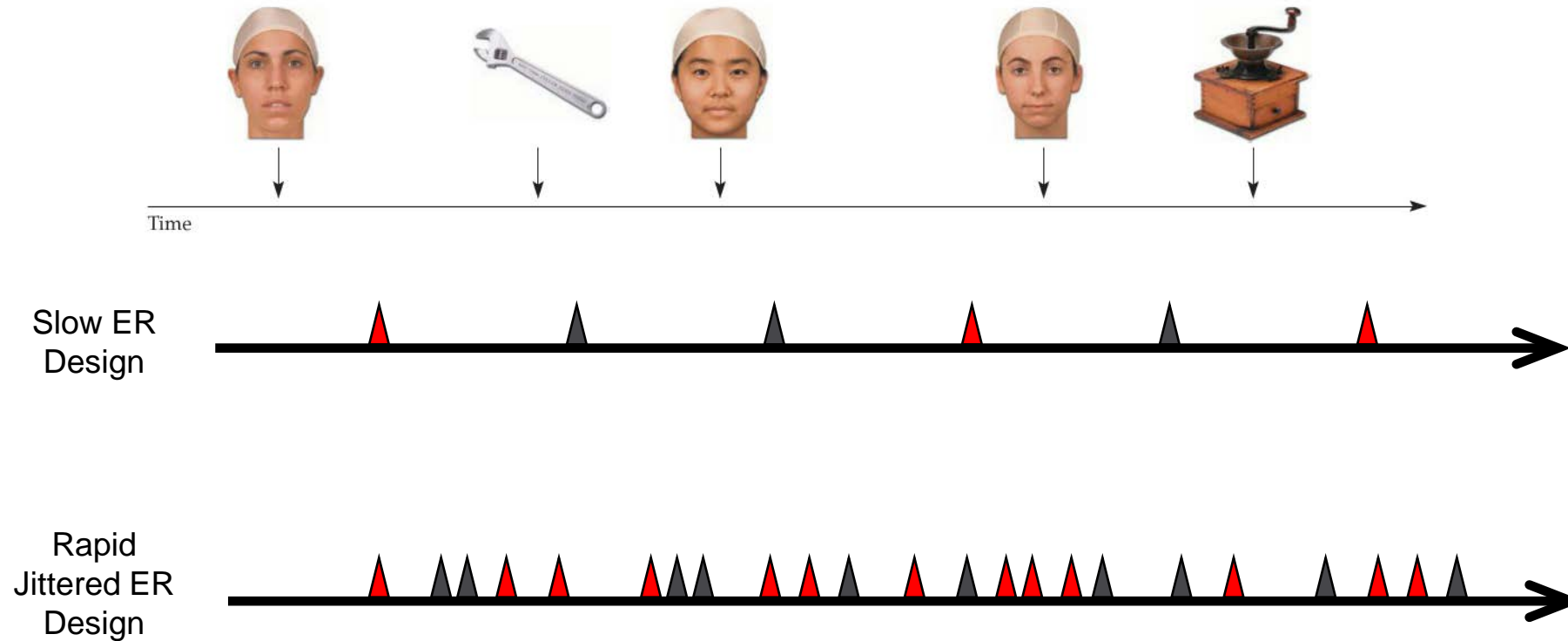
# Effect of Block Interval on FMRI HRF



# Recommendations for Using Blocked Design

- Length of a block
  - Minimally 10s and optimally 16s (Liu, 2004).
  - Equivalent for conditions or combination of conditions to be compared
- Evoking the same mental process throughout a block

# Event-related Designs



# Slow Event-Related Designs

Slow ER  
Design

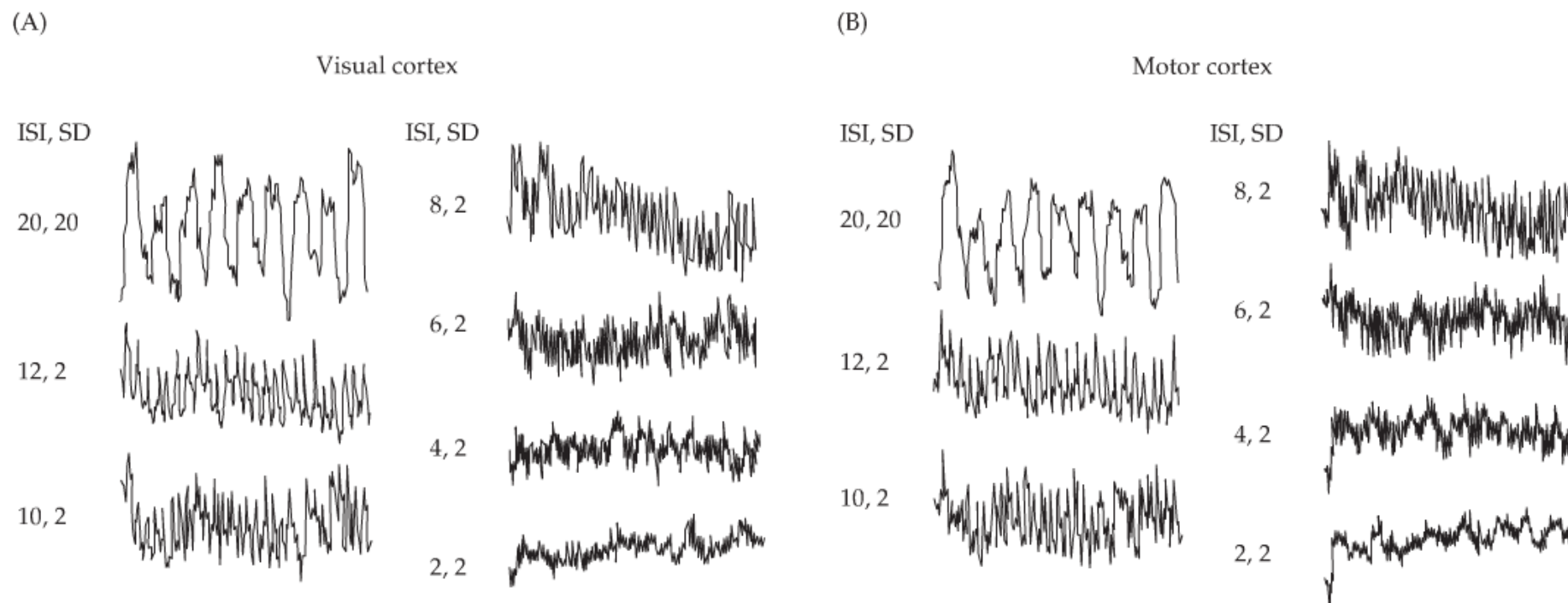


# Periodic (Slow) ER Design

- Fixed and long ISI
  - Usually  $> 15s$
  - Each event evokes a complete HR, and corresponding BOLD are selectively averaged.
  - Inefficient

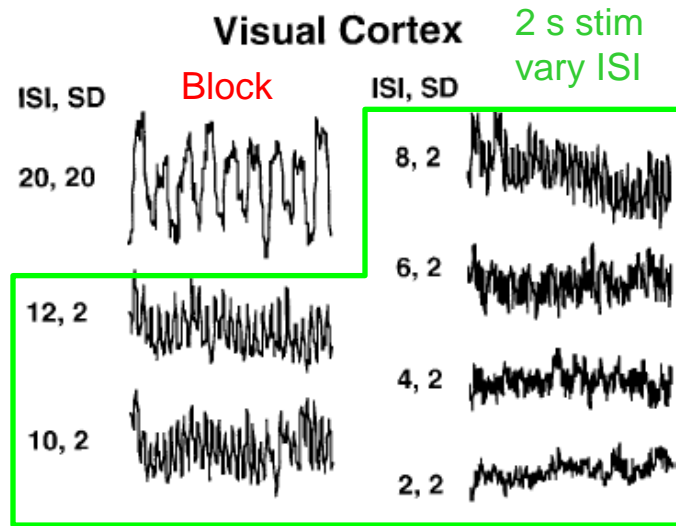


# Effects of ISI on ER-FMRI Activation



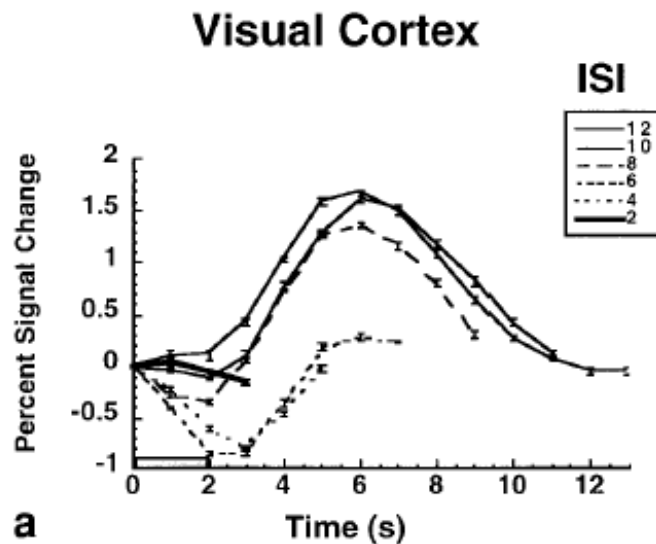
Bandettinni & Cox (2000)

# Slow Event-Related Design: Constant ITI



Bandettini et al. (2000)

What is the optimal trial spacing (duration + intertrial interval, ITI) for a Spaced Mixed Trial design with constant stimulus duration?

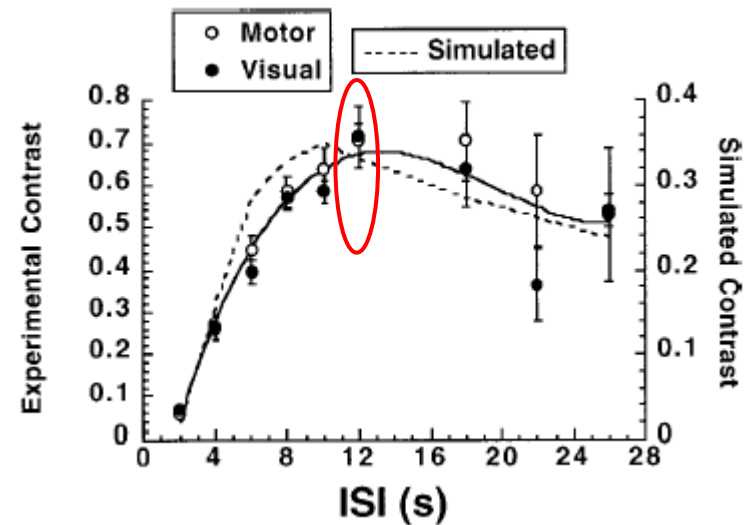


Event-related average

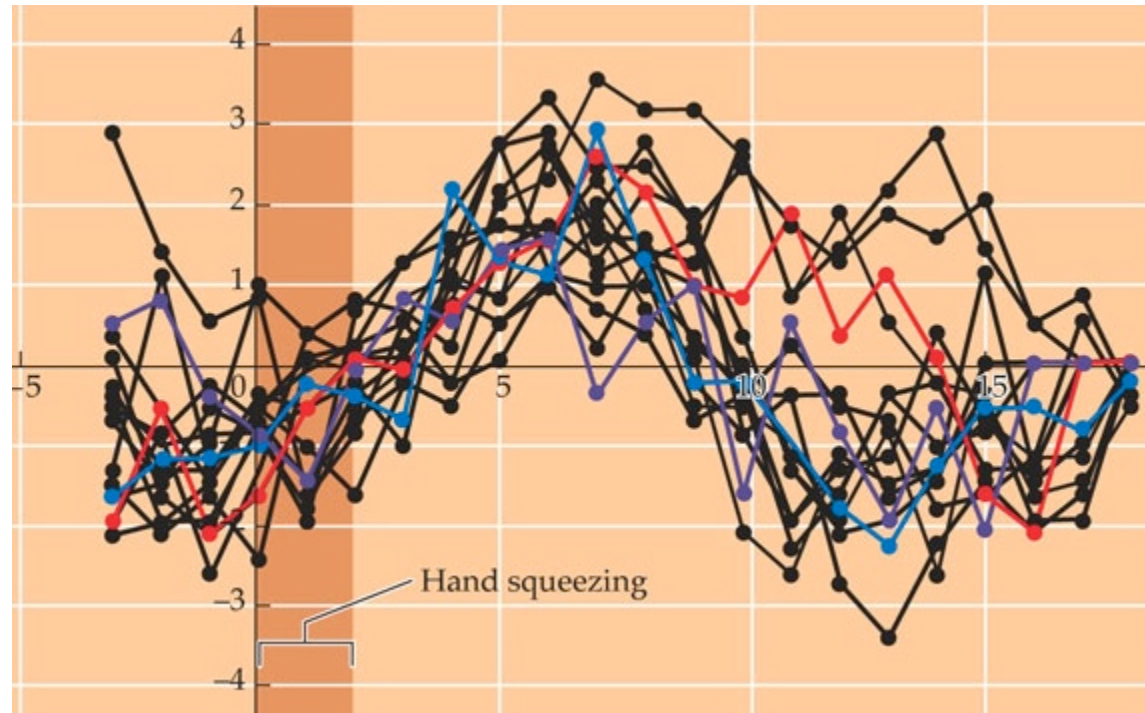
Source: Bandettini et al., 2000

# Optimal Constant ITI

- Brief (< 2 sec) stimuli:
  - optimal trial spacing = 12 sec
- For longer stimuli:
  - optimal trial spacing =  $8 + 2 * \text{stimulus duration}$
- Effective loss in power of event related design:
  - = -35%
  - i.e., for 6 minutes of block design, run ~9 min ER design

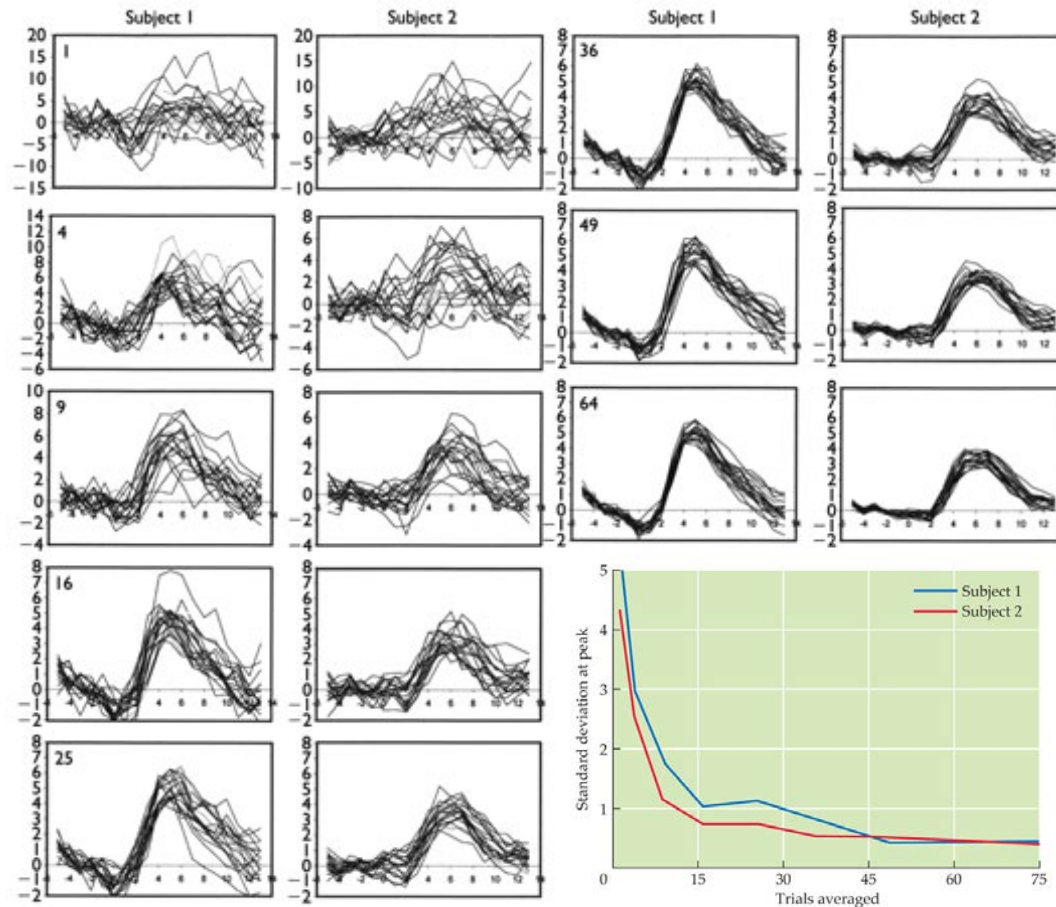


# Trial to Trial Variability



*Huettel, Song & McCarthy, 2004,  
Functional Magnetic Resonance Imaging*

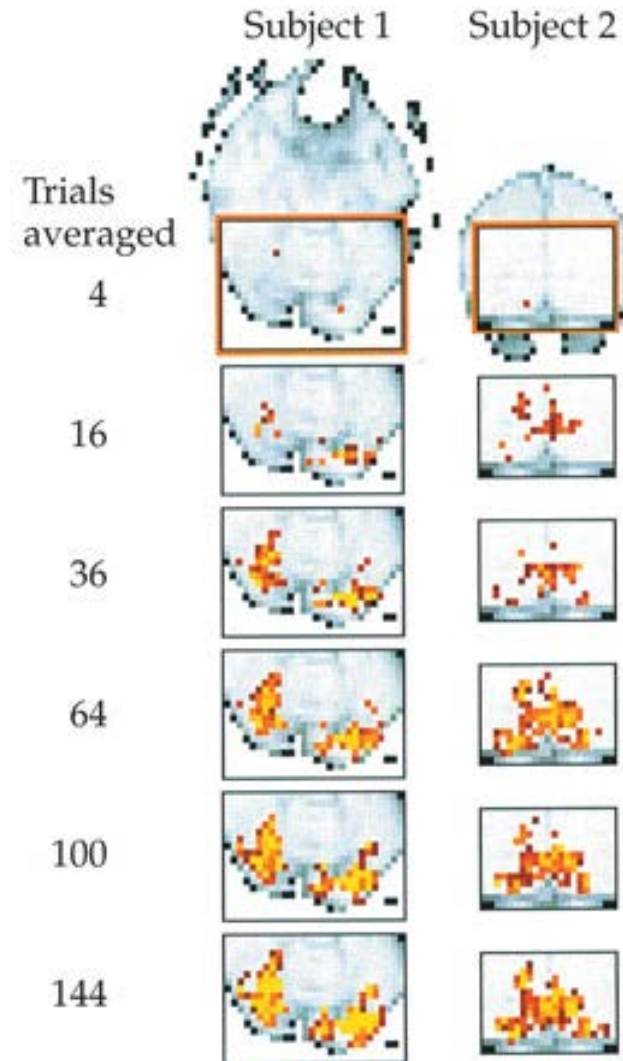
# How Many Trials Do You Need?



*Huettel, Song & McCarthy, 2004, Functional Magnetic Resonance Imaging*

- standard error of the mean varies with square root of number of trials
- Number of trials needed will vary with effect size
- Function begins to asymptote around 15 trials

# Effect of Adding Trials



*Huettel, Song & McCarthy, 2004, Functional Magnetic Resonance Imaging*



# Pros & Cons of Slow ER Designs

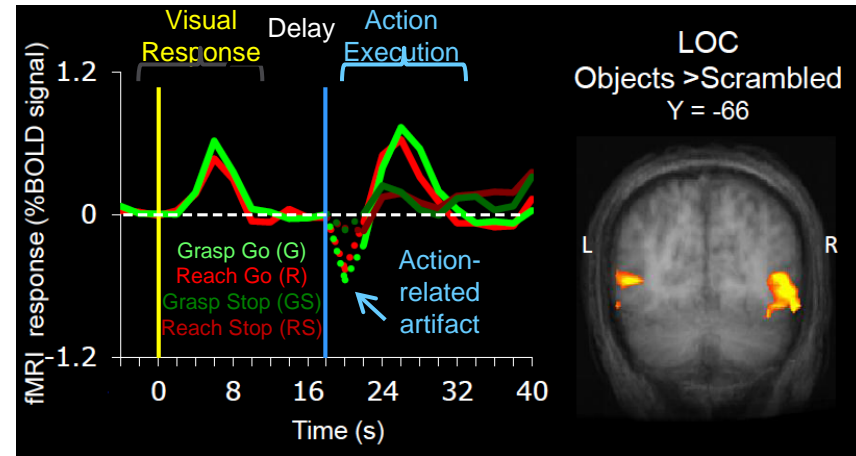
## Pros

- excellent estimation
- useful for studies with delay periods
- very useful for designs with motion artifacts (grasping, swallowing, speech) because you can tease out artifacts
- analysis is straightforward

## Cons

- poor detection power because you get very few trials per condition by spending most of your sampling power on estimating the baseline
- subjects can get VERY bored and sleepy with long inter-trial intervals

Example: Delayed Hand Actions  
(Singhal et al., under revision)



Really long delay:  
18 s



Effect of this design  
on our subject

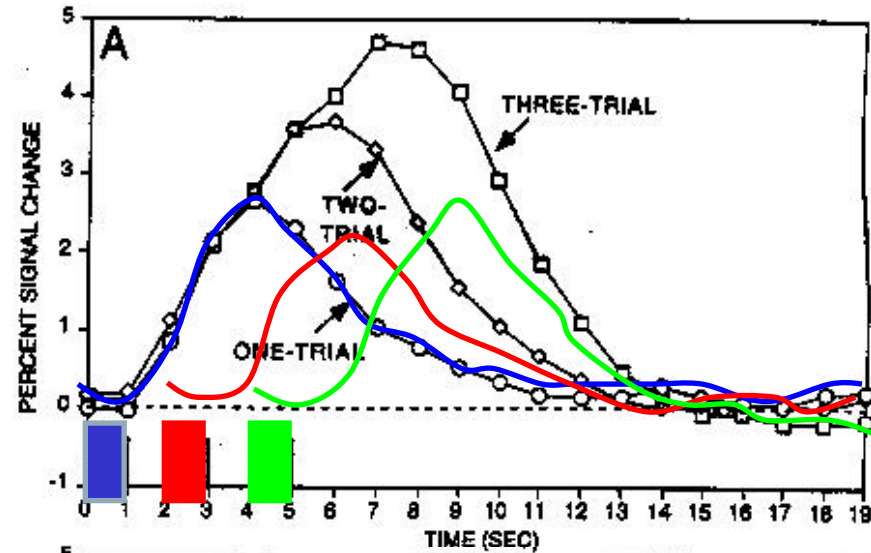
# 要不要再快一點？

- Yes, but we have to test assumptions regarding linearity of BOLD signal first

Rapid  
Jittered ER  
Design



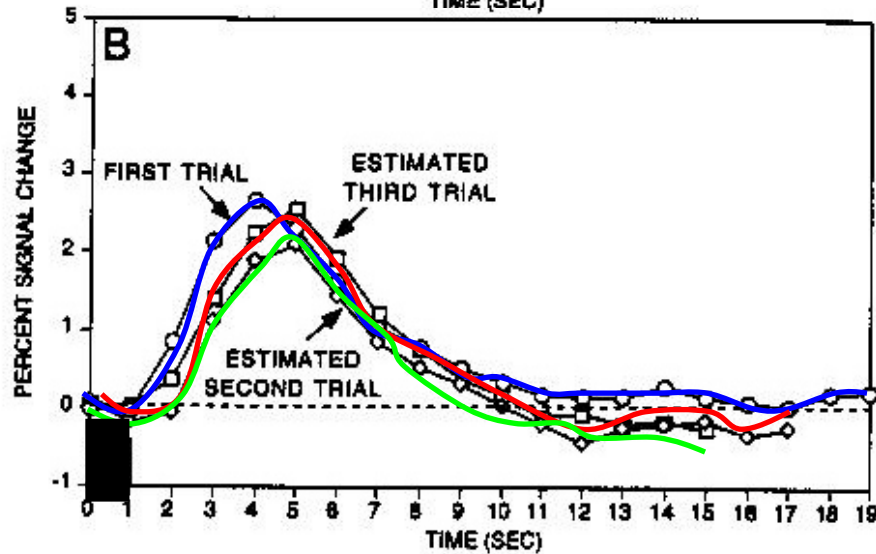
# Linearity of BOLD response



Linearity:  
“Do things add up?”

red = 2 - 1

green = 3 - 2

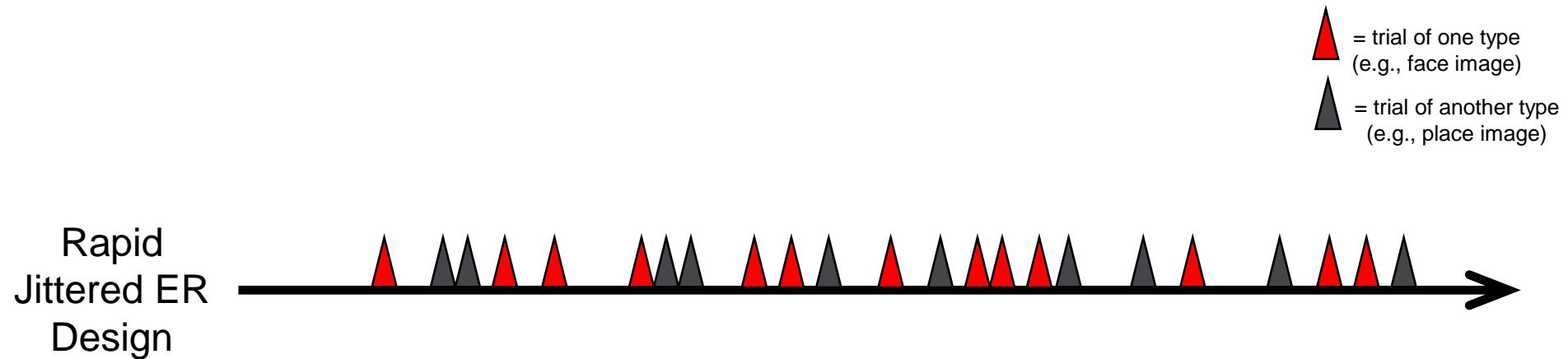


Sync each trial response  
to start of trial

Not quite linear  
but good enough!

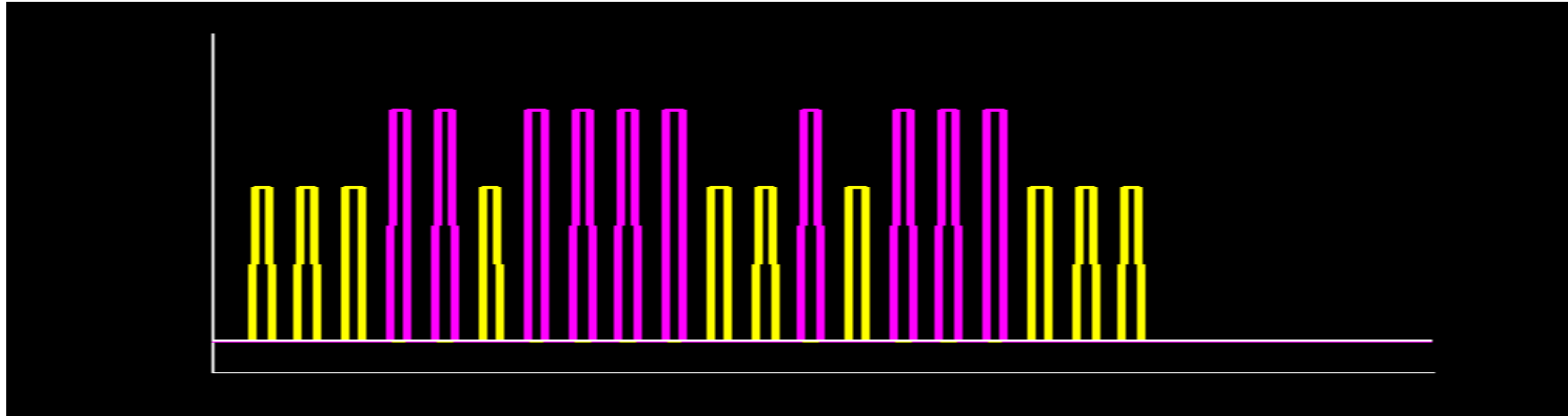
Source: Dale & Buckner, 1997

# Rapid Jittered ER Design

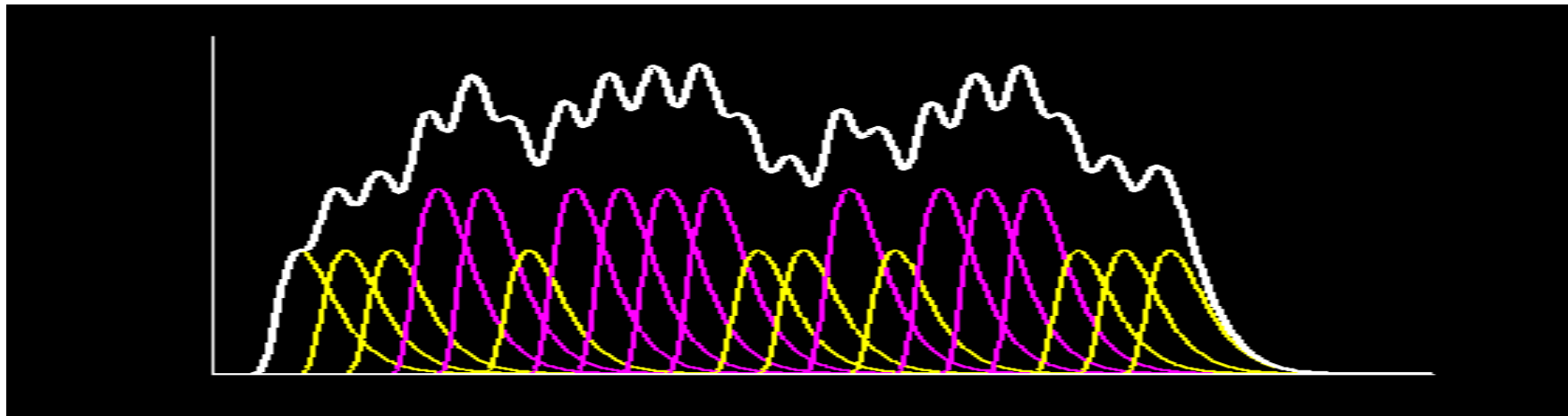


A popular choice is to use 'jittered' designs with inter-stimulus intervals of at least 4s and with exponentially decreasing delay frequencies up to 16s.

# BOLD Overlap With Regular Trial Spacing



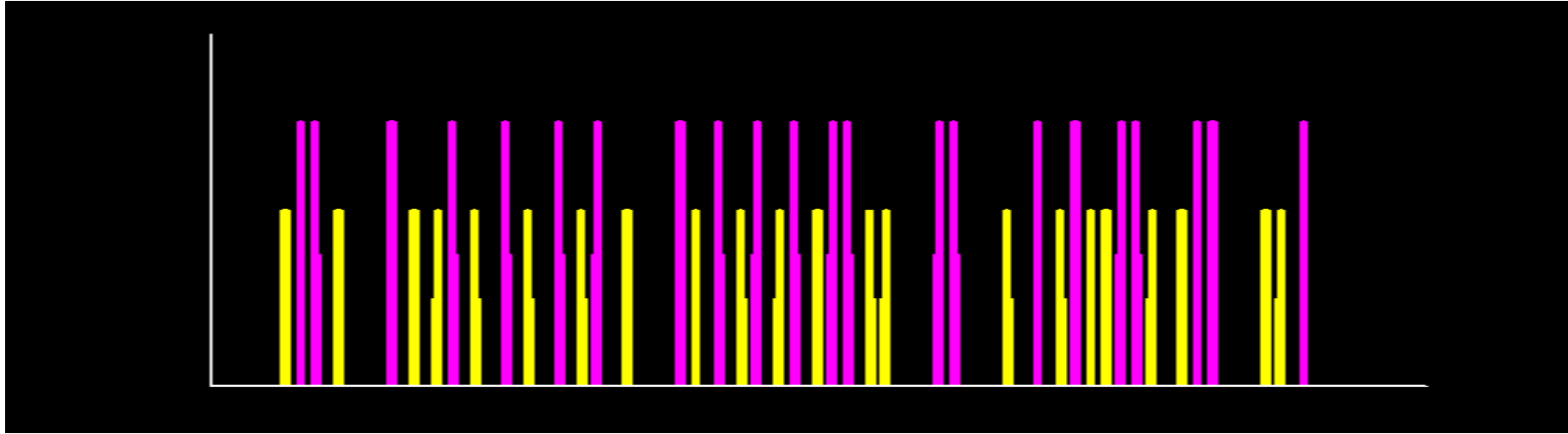
Neuronal activity from TWO event types with constant ITI



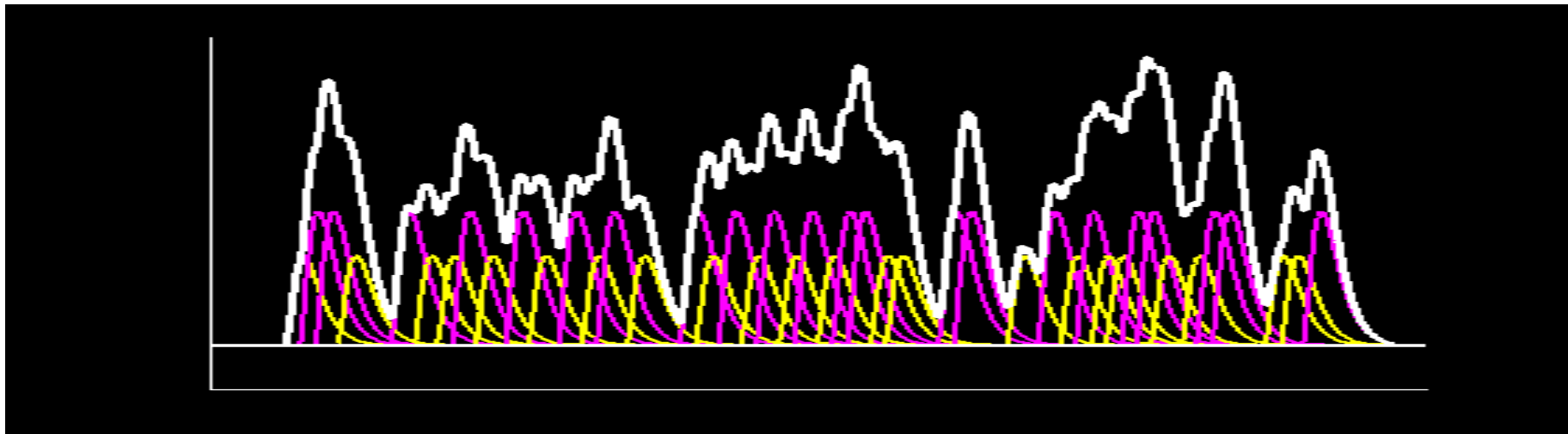
Partial tetanus BOLD activity from two event types

*Slide from Matt Brown*

# BOLD Overlap with Jittering



Neuronal activity from closely-spaced, jittered events



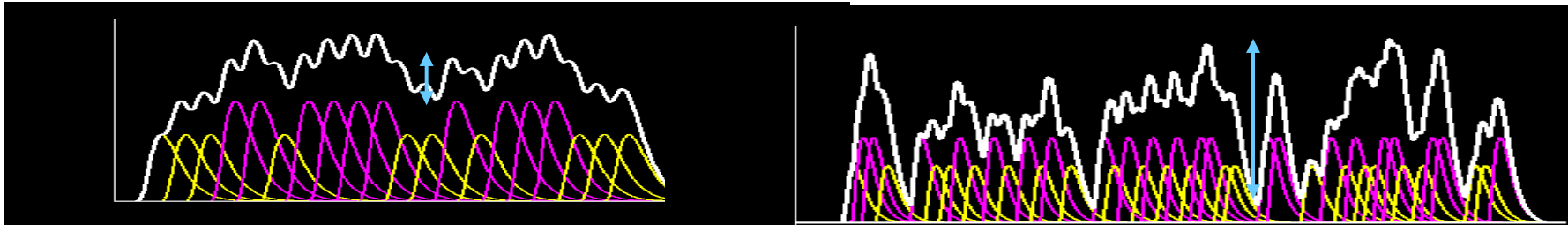
BOLD activity from closely-spaced, jittered events

*Slide from Matt Brown*



# Why jitter?

- Yields larger **fluctuations** in signal

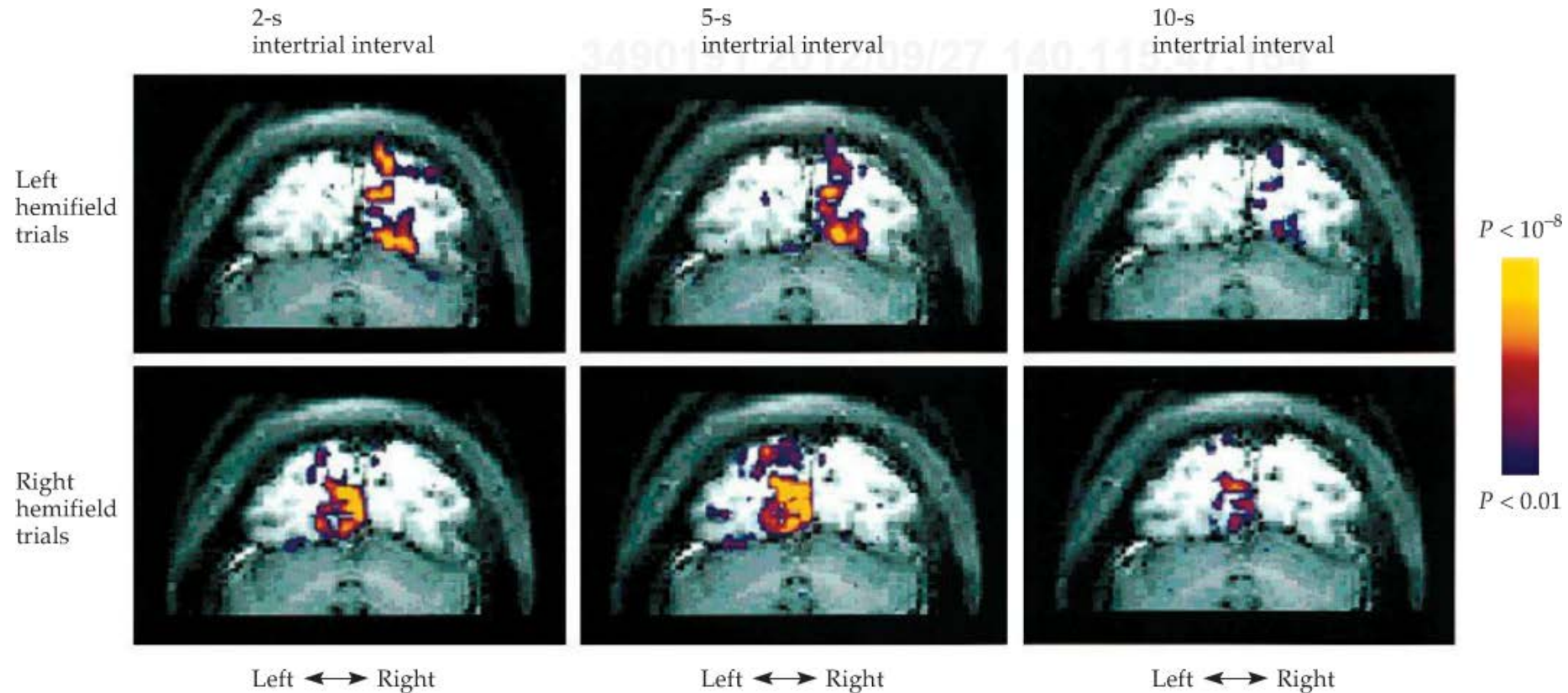


When pink is on, yellow is off  
→ pink and yellow are anticorrelated

Includes cases when both pink and yellow are off  
→ less anticorrelation

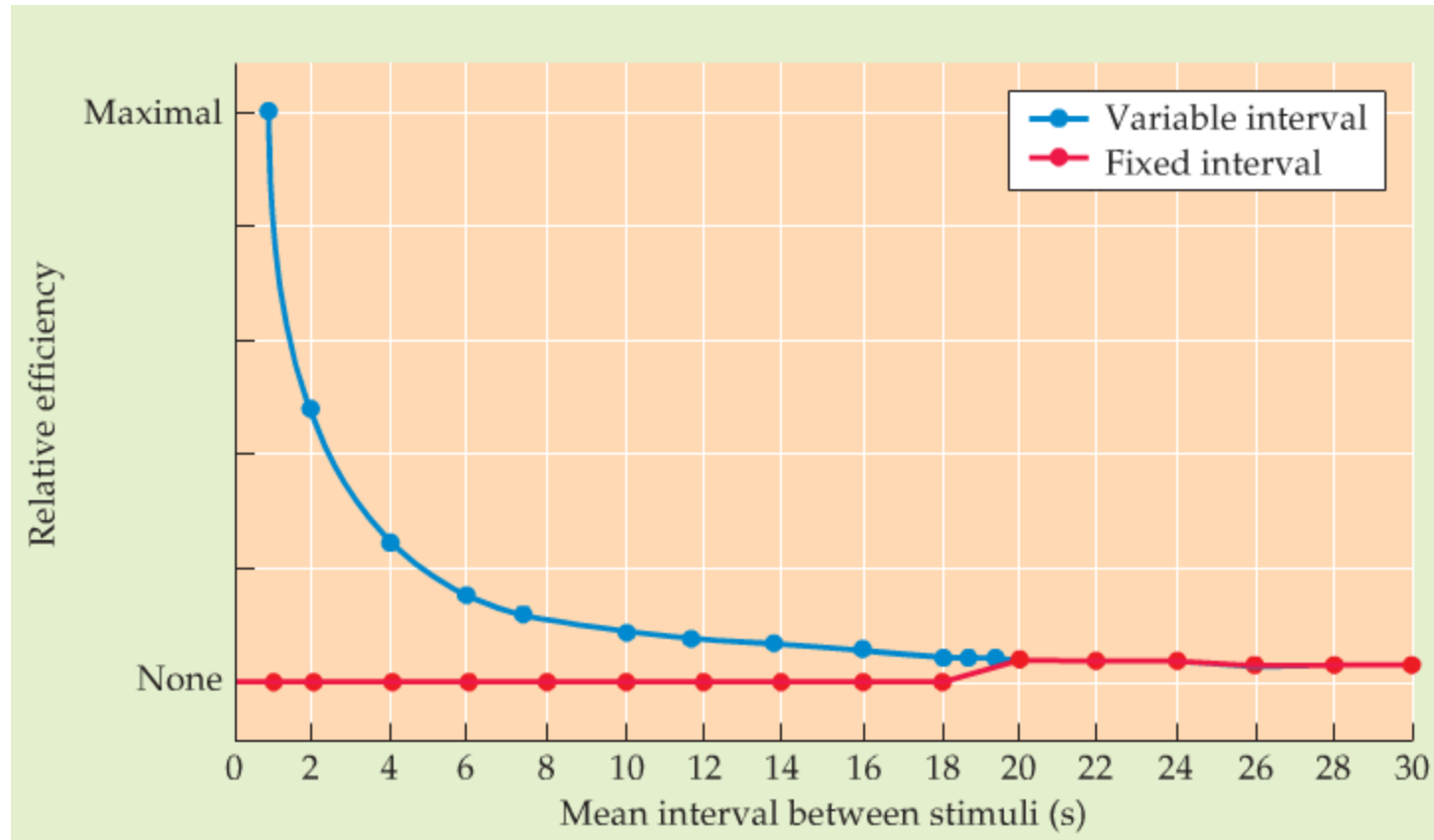
- Without jittering predictors from different trial types are strongly **anticorrelated**
  - As we know, the GLM doesn't do so well when predictors are correlated (or anticorrelated)

# Rapid ER-FMRI with Randomized Stimulus Presentation



Short randomized ITI enhances detection power.

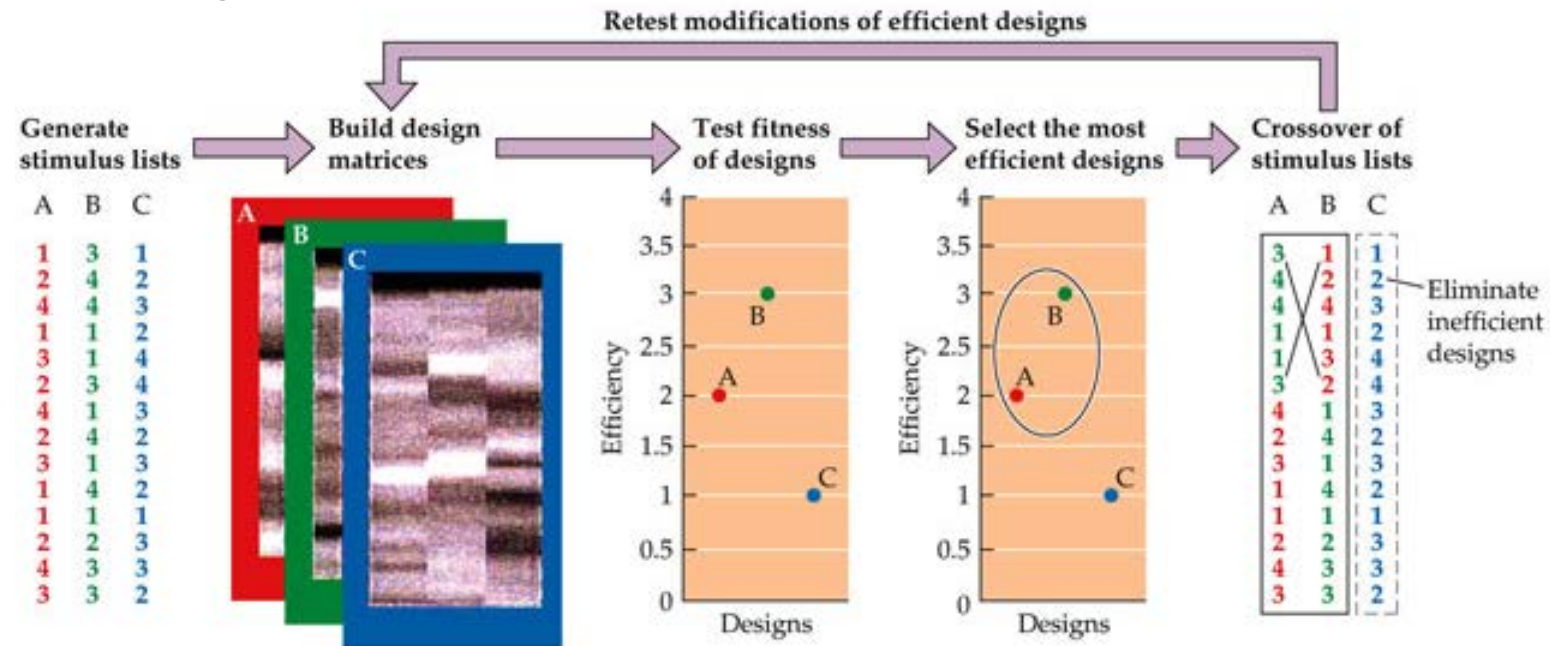
# Variable vs. Fixed Intervals



After Dale (1999)

# Algorithms for Picking Efficient Designs

## Genetic Algorithms



<http://wagerlab.colorado.edu/tools>

# Pros & Cons of Applying Standard GLM to Rapid-ER Designs

## Pros

- Acceptable detection power
- trials can be put in unpredictable order
- subjects don't get so bored

## Cons and Caveats

- reduced detection compared to block designs
- requires stronger assumptions about linearity
  - BOLD is non-linear with inter-event intervals  $< 6$  sec.
  - Nonlinearity becomes severe under 2 sec.
- errors in HRF model can introduce errors in activation estimates

# Good Practices in fMRI

- Evoke the cognitive processes of interest
- Maximize data collection from each subject
- Maximize sample size
- Choose conditions and timings that maximize evoked changes in the process of interests
- Minimize correlation between BOLDs of successive events
- Compute correlation between behavioral performance and activation

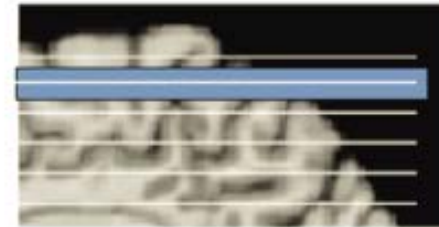
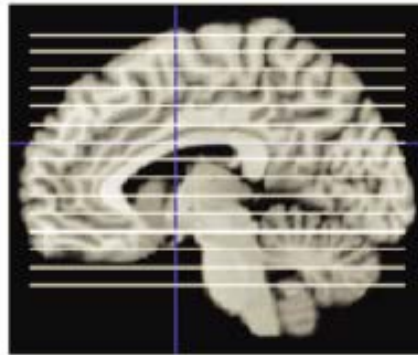
腦造影 vs. 腦照影

# FMRI資料前處理

Preprocessing

# Images: Basic Terminology

Field of View (FOV)  
(e.g. 192 mm)

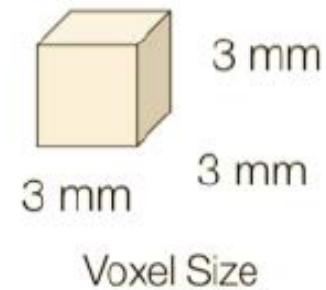


Slice thickness  
(e.g., 3 mm)

Matrix Size  
(e.g., 64 x 64)



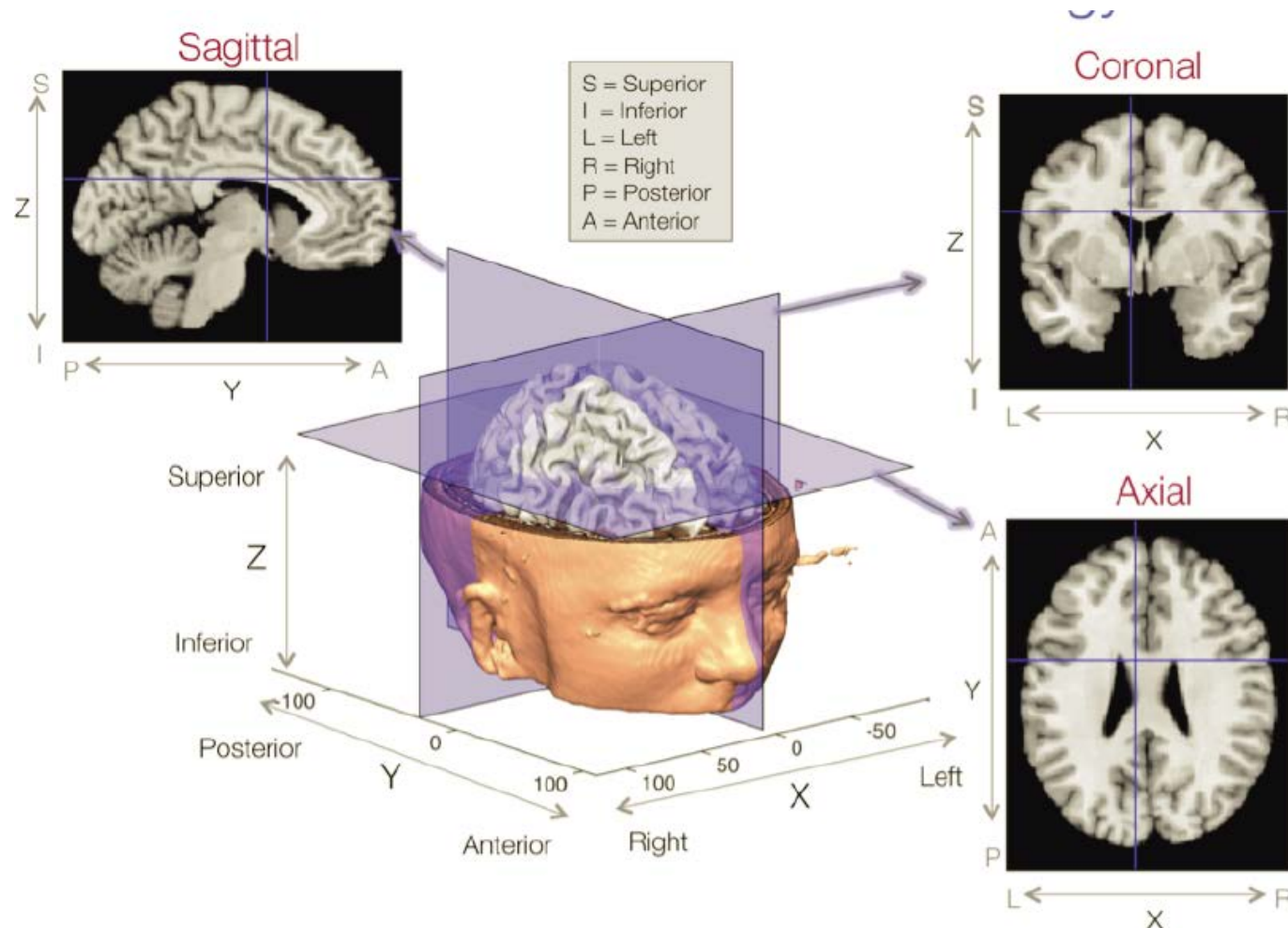
In-plane resolution  
 $192 \text{ mm} / 64 = 3 \text{ mm}$



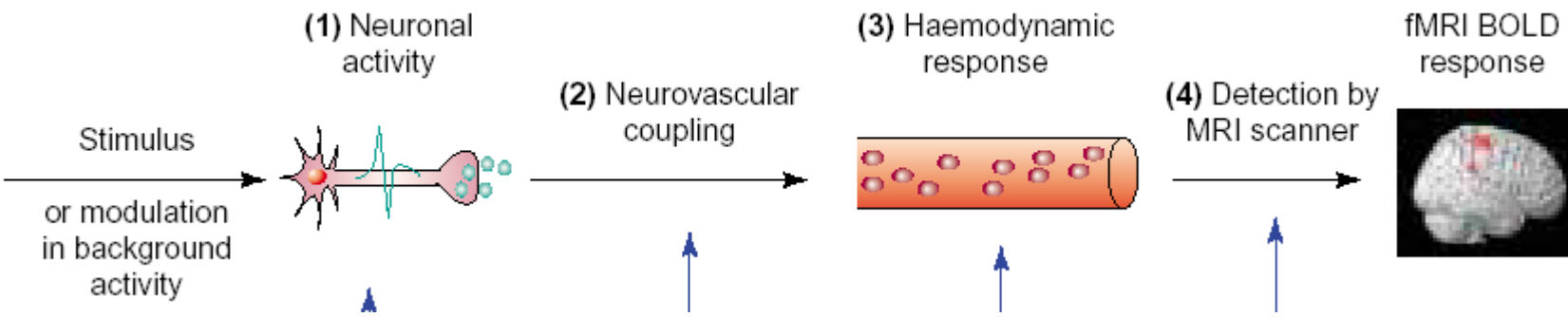
Source: Lindquist & Tor (2015)



# Brain Dimensions and Terminology



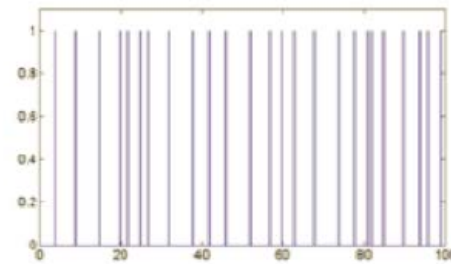
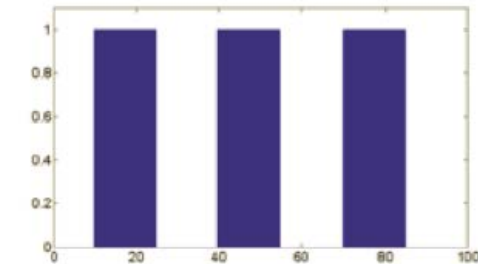
Source: Lindquist & Tor (2015)



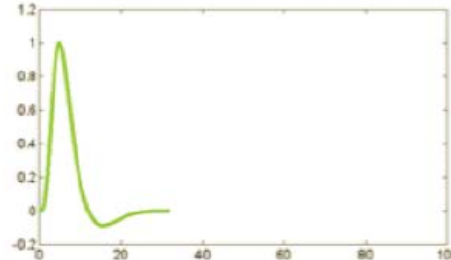
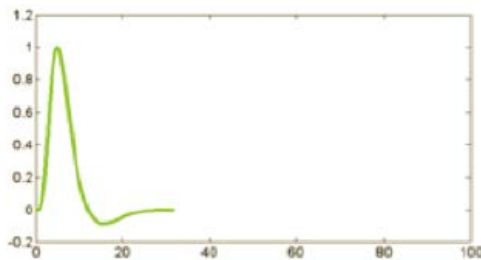
Block Design

Event-Related

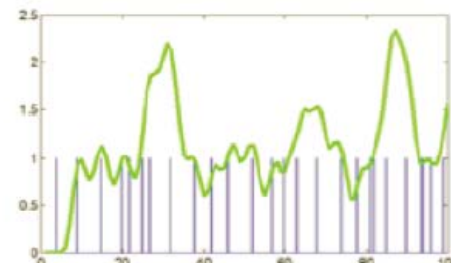
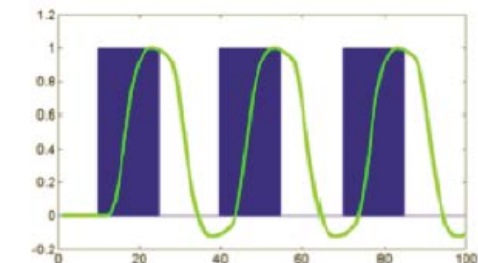
Experimental  
Stimulus Function



Hemodynamic  
Response  
Function



Predicted  
Response

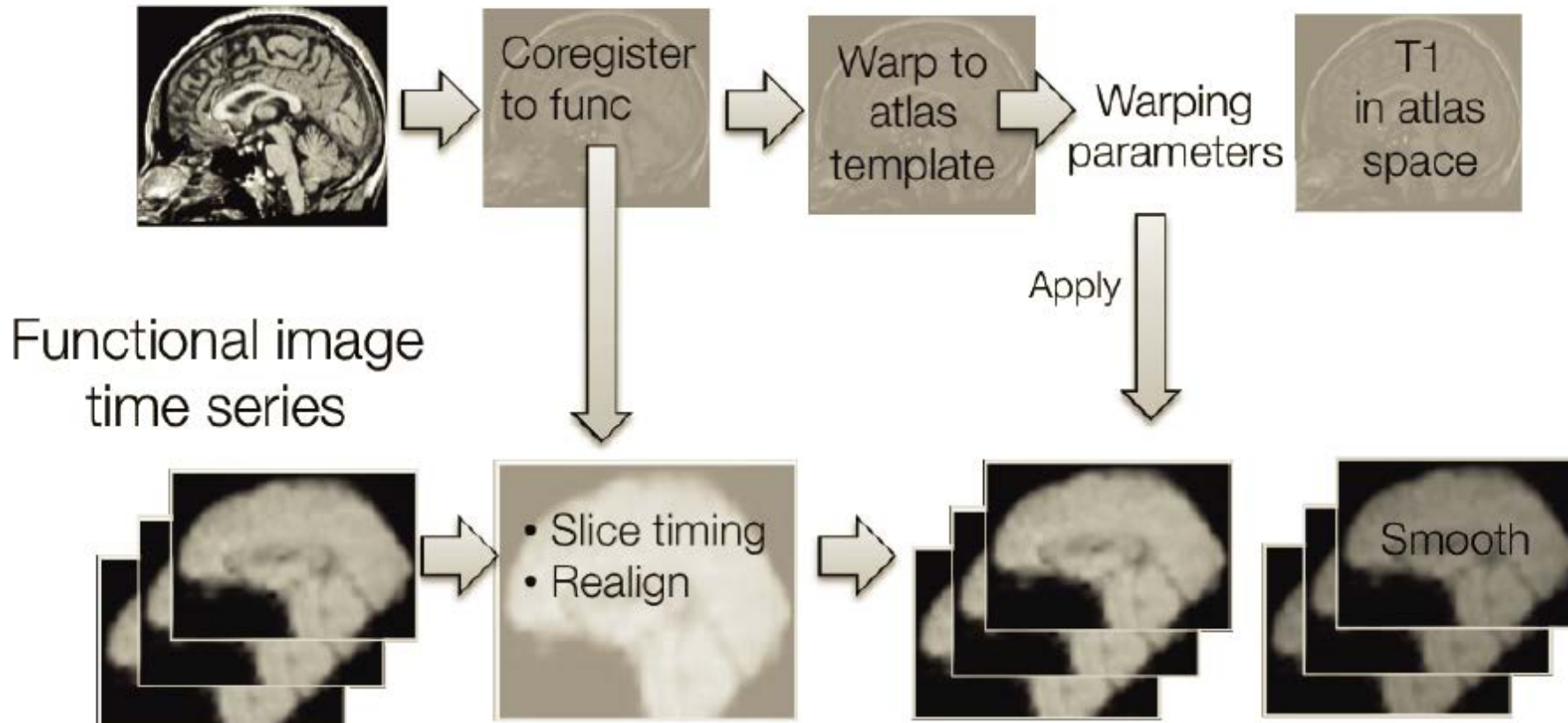


Two-gamma function

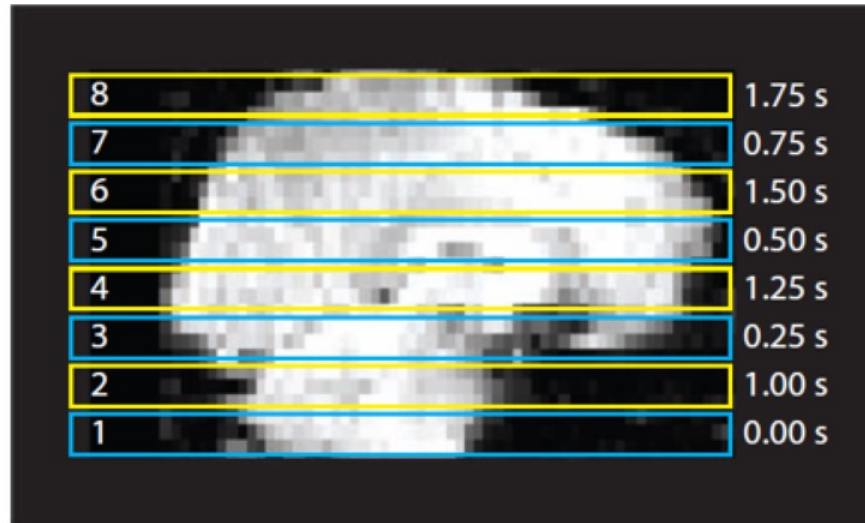
$$h(t) = \frac{t^{\alpha_1-1} \beta_1^{\alpha_1} e^{-\beta_1 t}}{\Gamma(\alpha_1)} - c \frac{t^{\alpha_2-1} \beta_2^{\alpha_2} e^{-\beta_2 t}}{\Gamma(\alpha_2)}$$

# Overview of Preprocessing

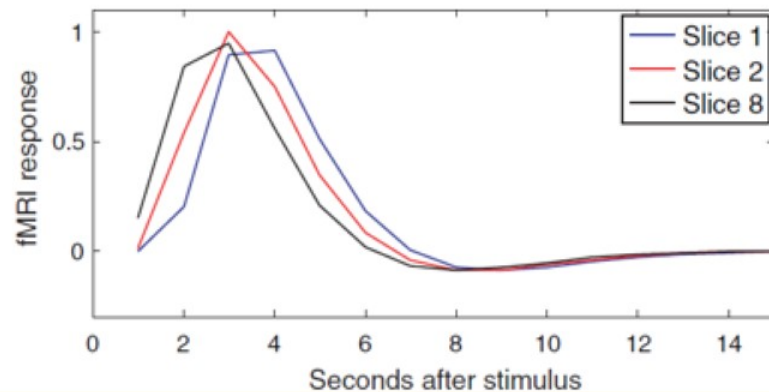
Structural (T1)



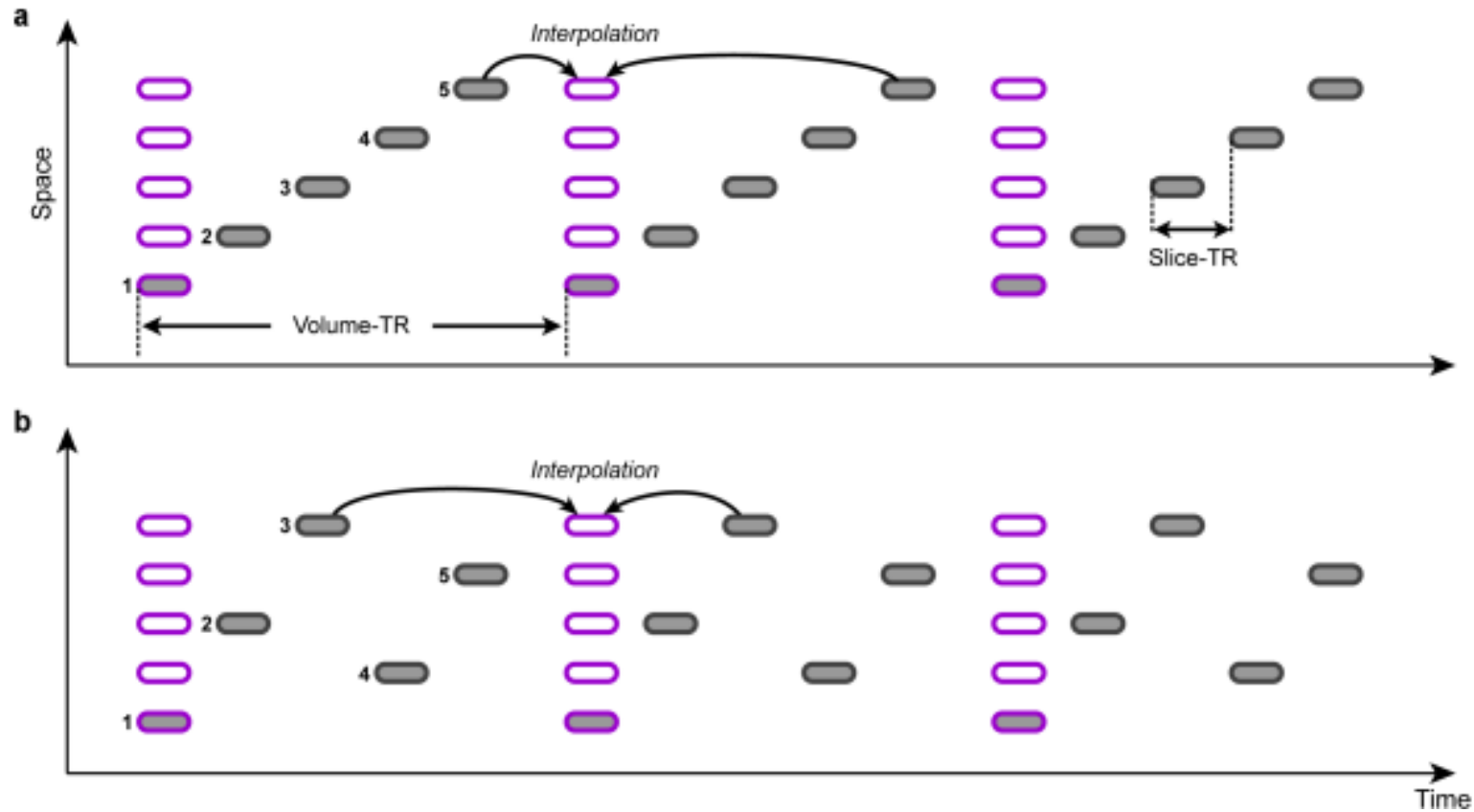
# Slice Time Problem



Not accounting for the timing differences between slices may lead to problematic time course differences between voxels on different slices.



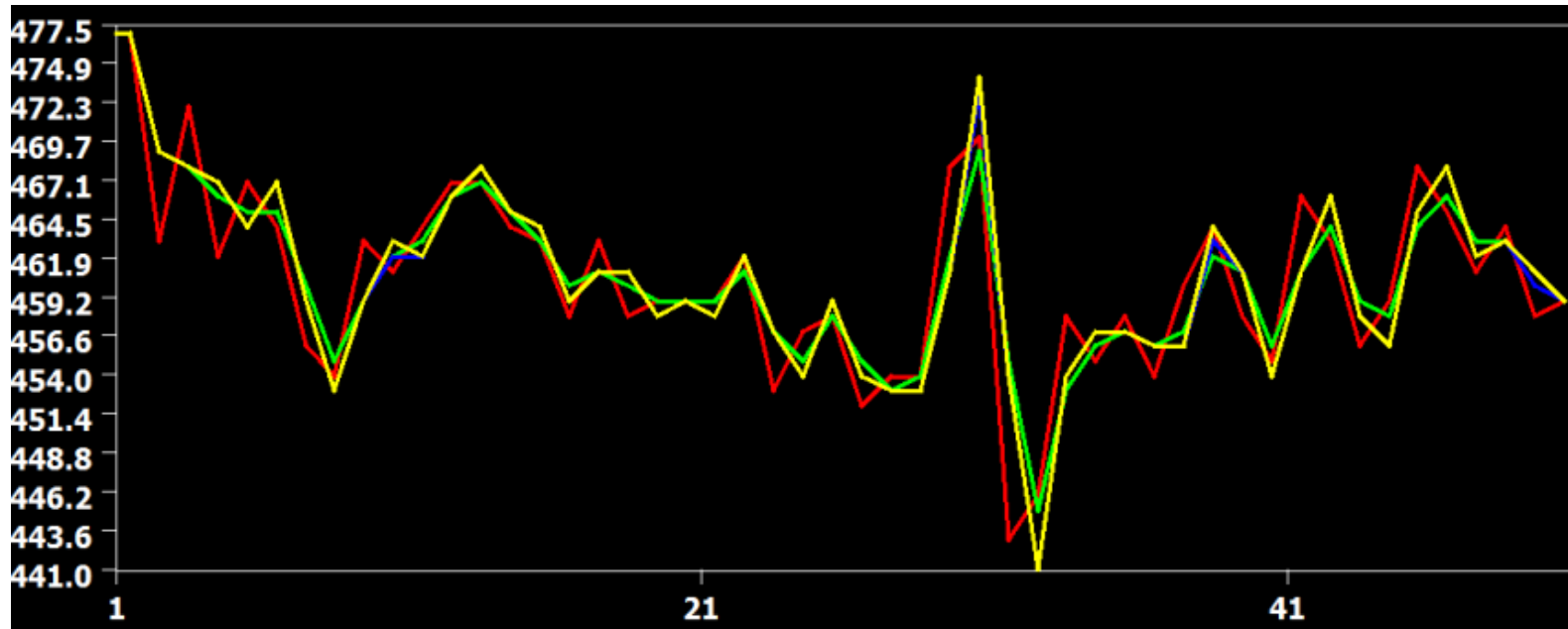
# More on Slice Time Correction



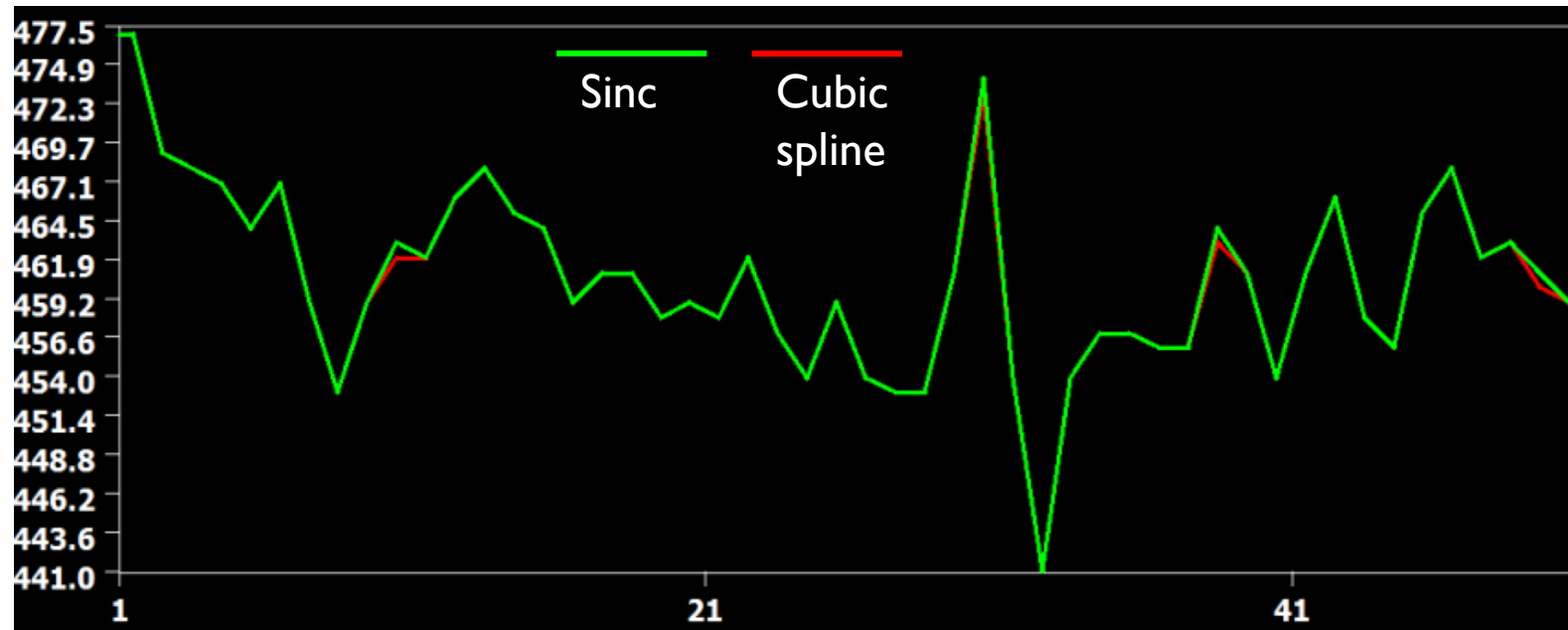
# Raw from the Last Slice



# Raw Data from Middle Slice



# Cubic Spline vs. sinc

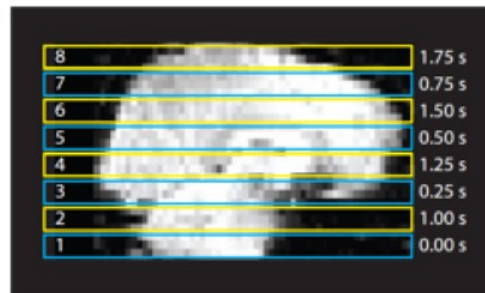




# Reasons Not to Correct for Slice Time

---

Propagation of artifacts



With short TR and interleaved acquisition, slice-timing problems is minimal

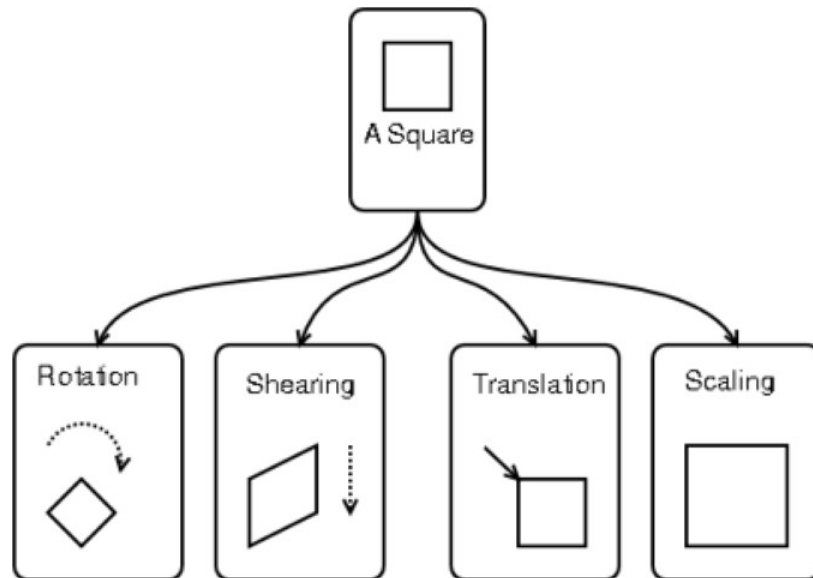
- Particularly after spatial smoothing

Temporal derivatives absorbs the impacts

# Spatial Transformation

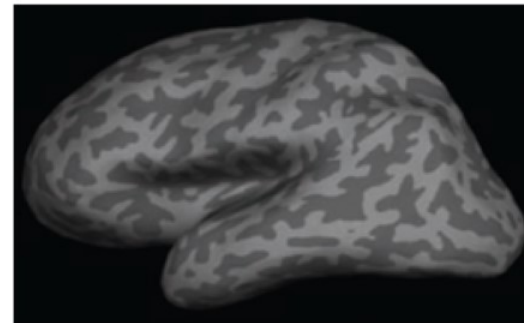
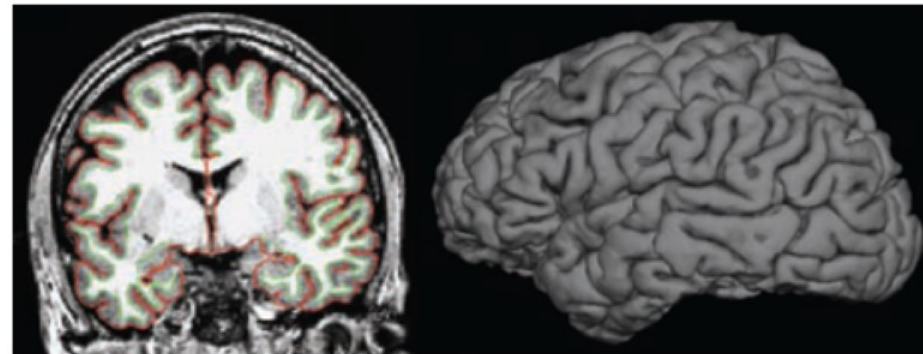
## Volume-based transformations

- Changes to 3D volume of data



## Surface-based registration

- Changes to surface data



# Models for Spatial Transformations

---

## Affine transformation

- Translation, rotation, scaling, shearing
- Rigid-body transformation
  - No scaling and shearing

## Piecewise linear transformation

- Divide the image into sub-regions and transform each of them respectively

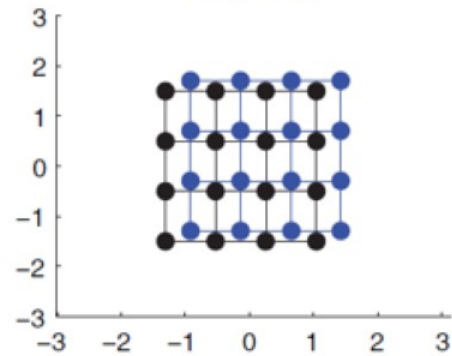
## Nonlinear transformation

- Transforming higher-dimensional representations of the image in a non-linear fashion

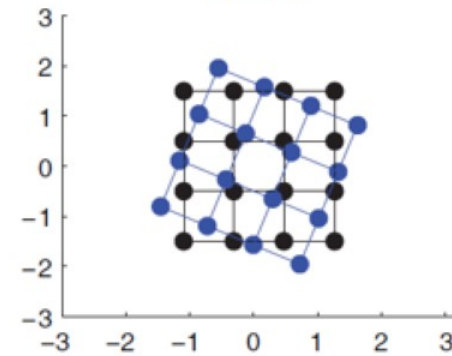
# Affine Transformations

---

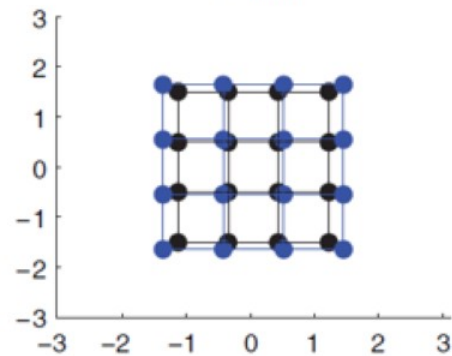
Translation



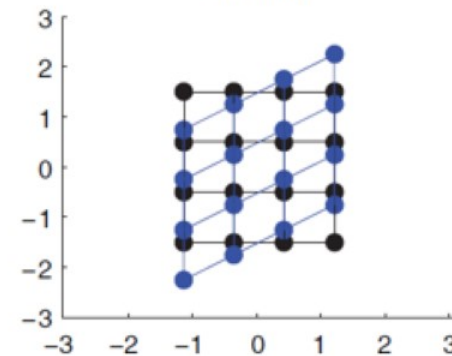
Rotation



Scaling



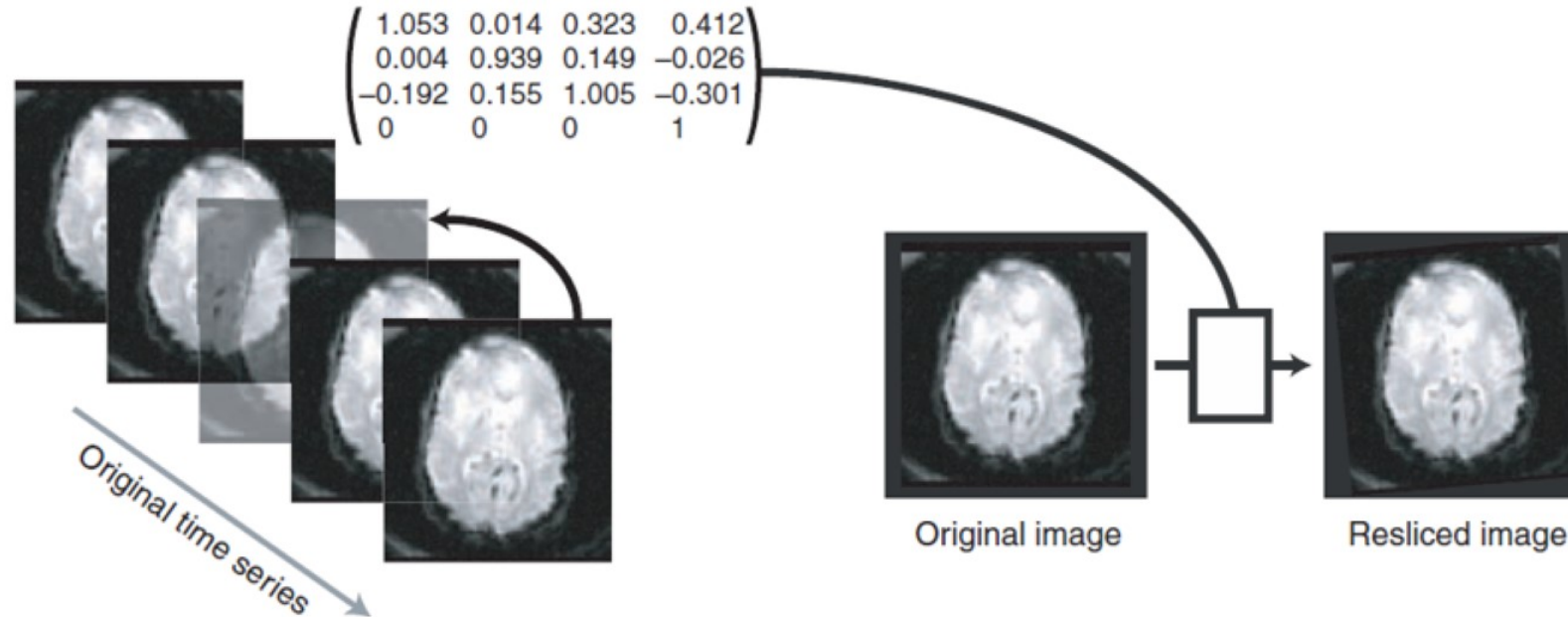
Shearing



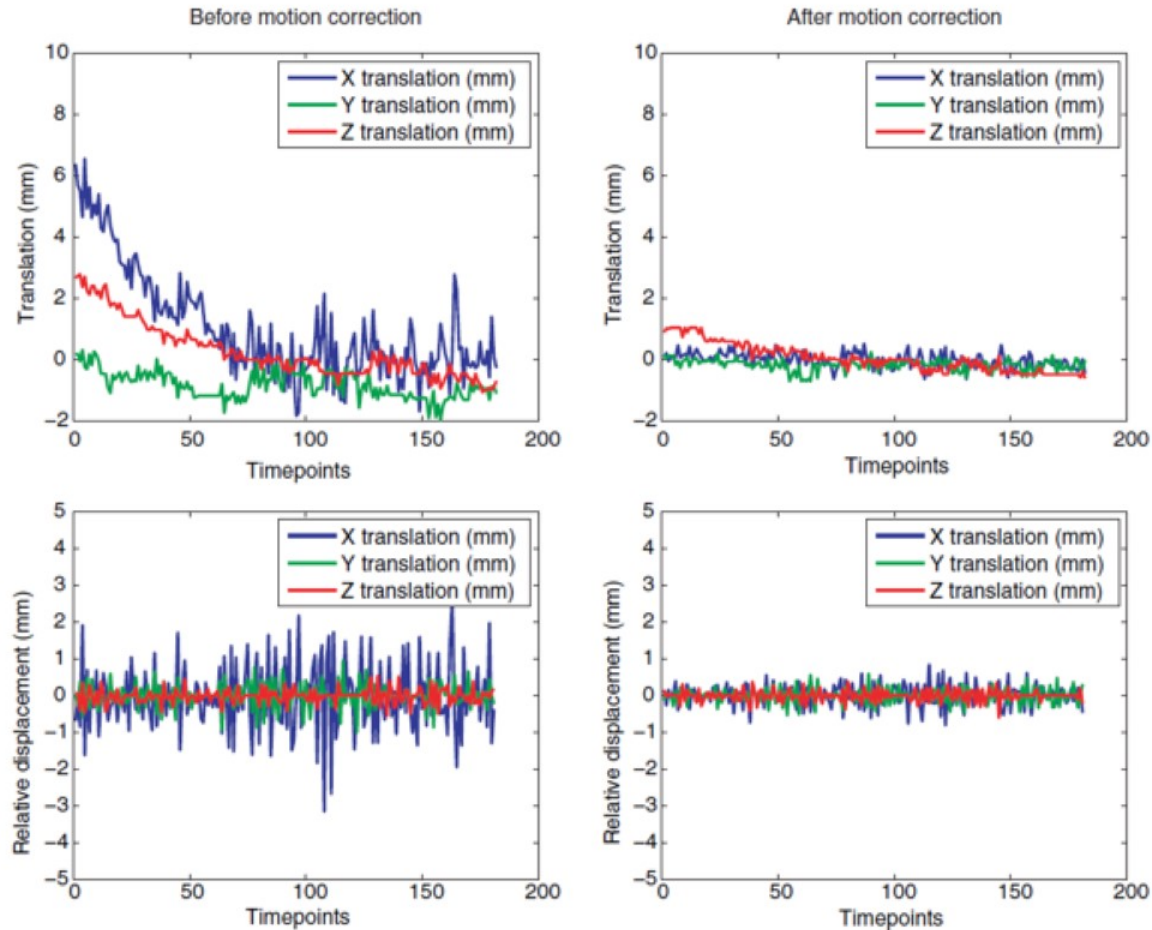
# Motion Correction (Realignment)

Step 1: Estimate parameters

Step 2: Reslice images



# Plots of Estimated Head Motion





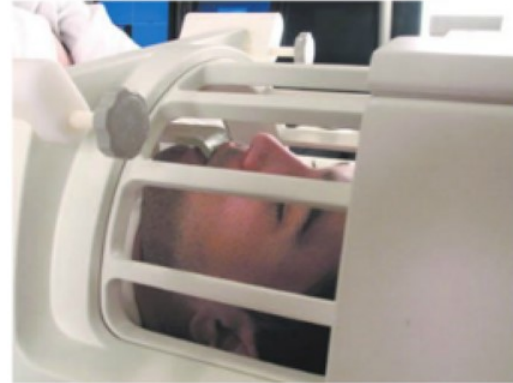
# Methods to Prevent Head Motion

---

(A)



(B)



(C)



(D)



# Spatial Smoothing

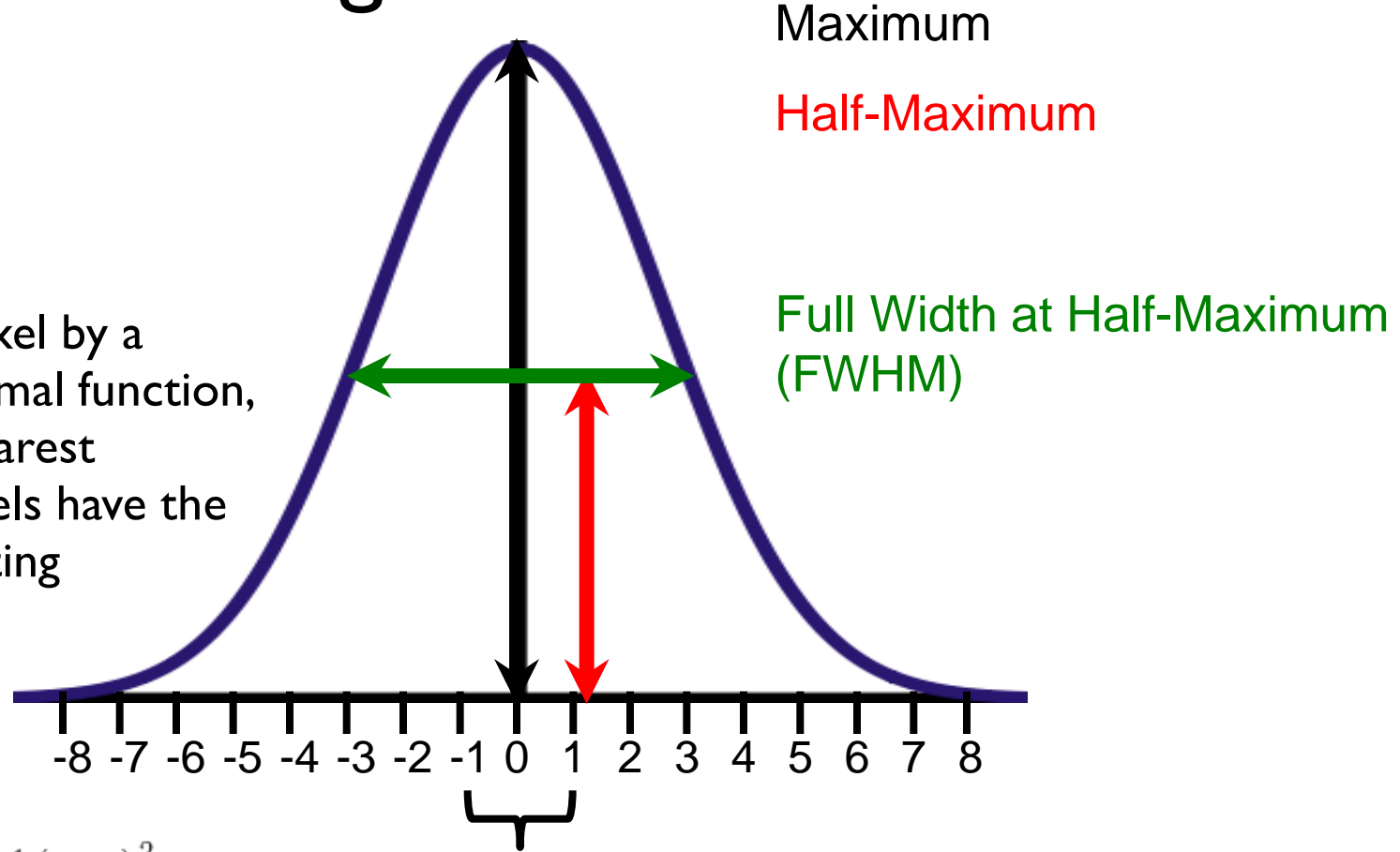
- Enhancing signal-to-noise ratio
  - By averaged out variation at smaller scale
- Enhancing cross-individual overlap
  - Sacrifice spatial resolution for power
- Fulfilling assumption of data analysis
  - Gaussian random fields



# Spatial Smoothing

Gaussian kernel

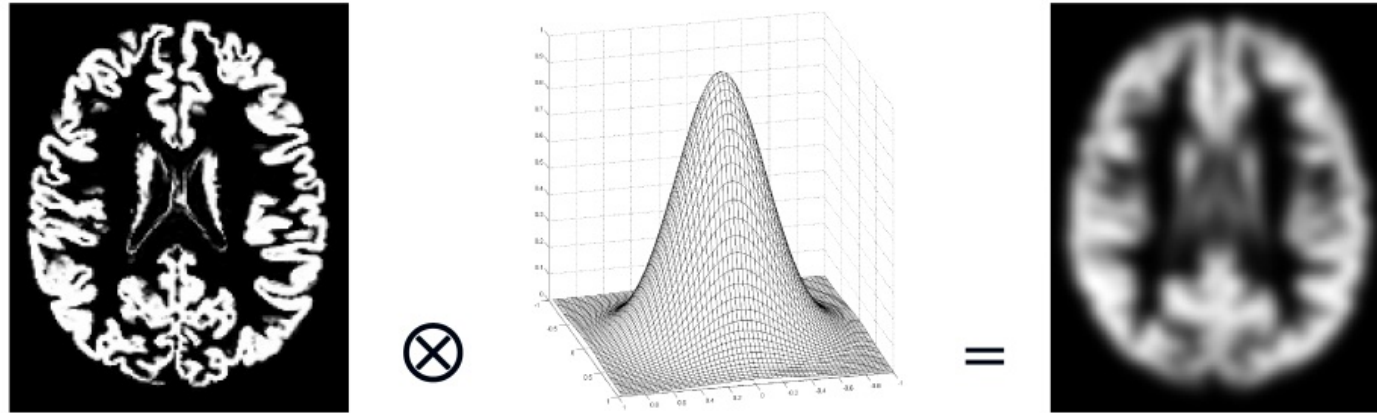
smooth each voxel by a Gaussian or normal function, such that the nearest neighboring voxels have the strongest weighting



$$g(x) = \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{1}{2}\left(\frac{x-\mu}{\sigma}\right)^2}$$

$$FWHM = 2\sigma\sqrt{2\ln(2)}$$

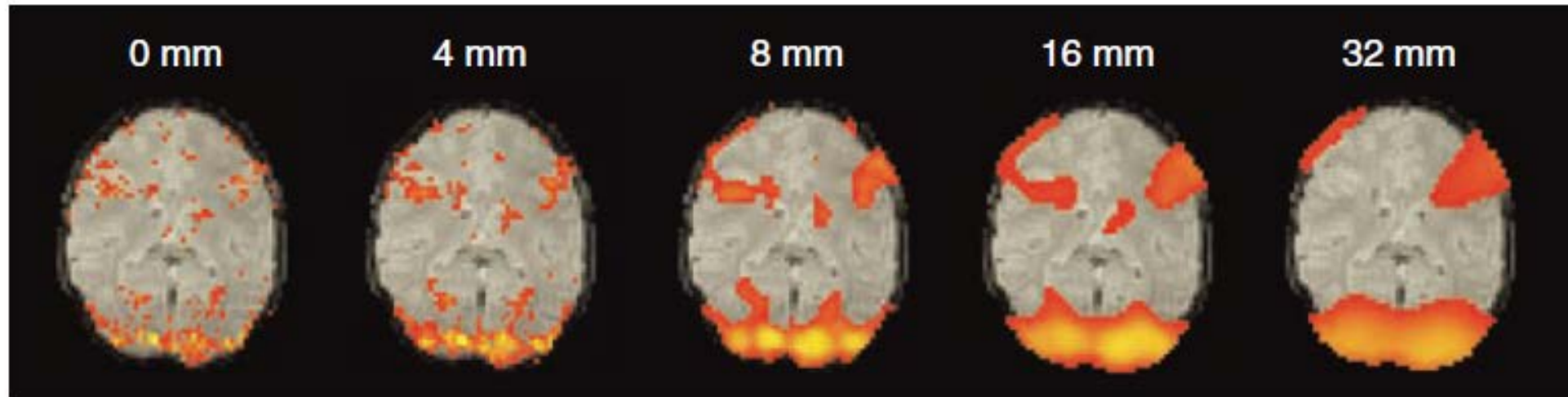
# 2D Spatial Smoothing



$$f(x, y) = A \exp \left( - \left( \frac{(x - x_0)^2}{2\sigma_x^2} + \frac{(y - y_0)^2}{2\sigma_y^2} \right) \right).$$

$$FWHM = \sqrt{FWHM_{intrinsic}^2 + FWHM_{applied}^2}$$

# Effect of Smoothing on Activation



# Should you spatially smooth?

- Advantages

- Increases Signal to Noise Ratio (SNR)
  - **Matched Filter Theorem**: Maximum increase in SNR by filter with same shape/size as signal
- Reduces number of comparisons
  - Allows application of **Gaussian Field Theory**
- May improve comparisons across subjects
  - Signal may be spread widely across cortex, due to inter-subject variability

"Why would you spend \$4 million to buy an MRI scanner and then blur the data till it looked like PET?"

-- Ravi Menon

- Disadvantages

- Reduces spatial resolution
- Challenging to smooth accurately if size/shape of signal is not known

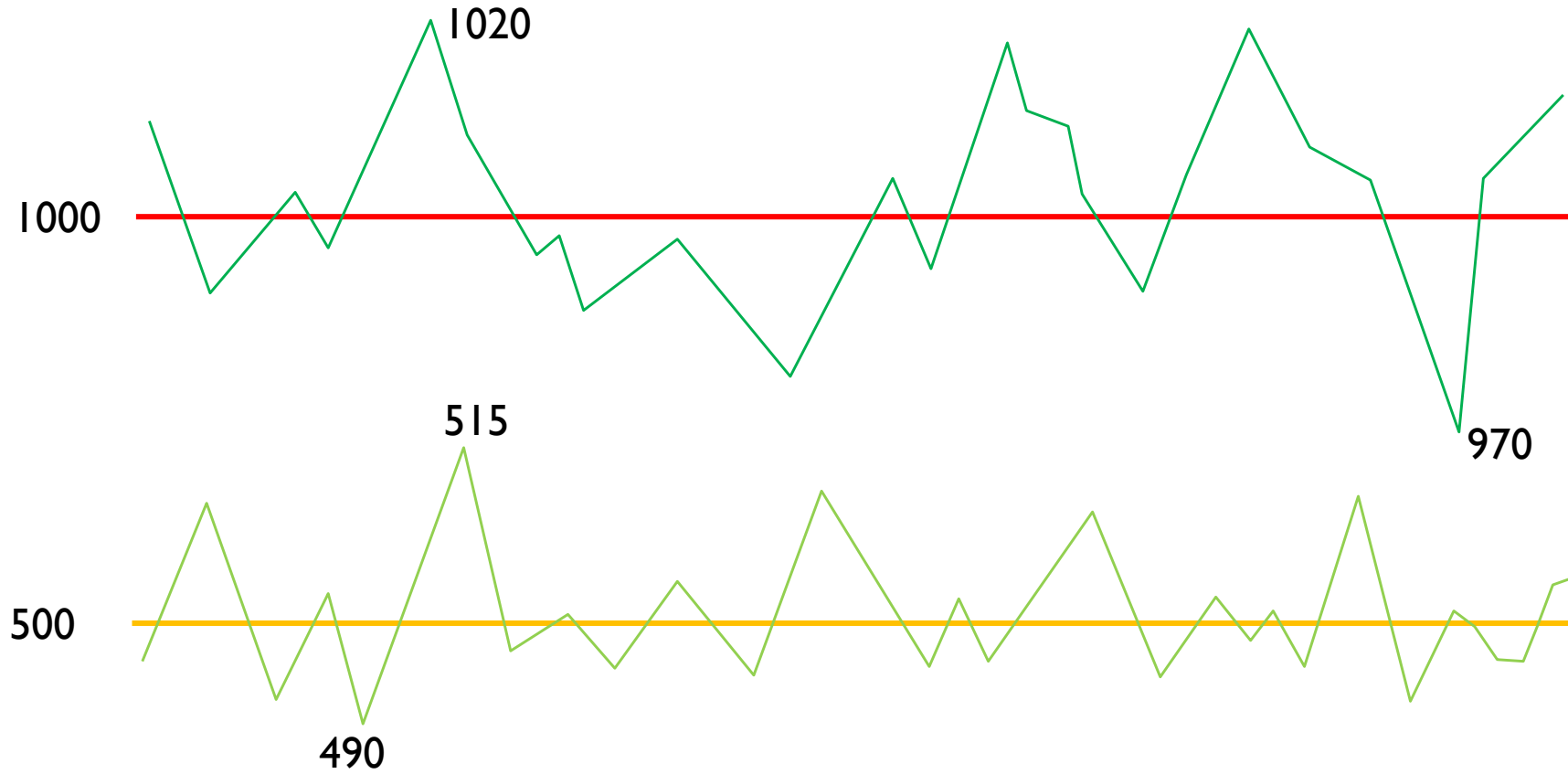
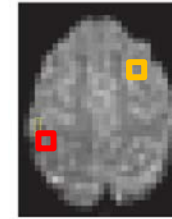
# Recommendation for Smoothing

- Noise reduction
  - Filter smaller than expected extent of activation
- Reducing structural variability
  - Variability in the population
  - Efficiency of normalization
- Gaussian random fields assumption
  - FWHM twice the voxel size

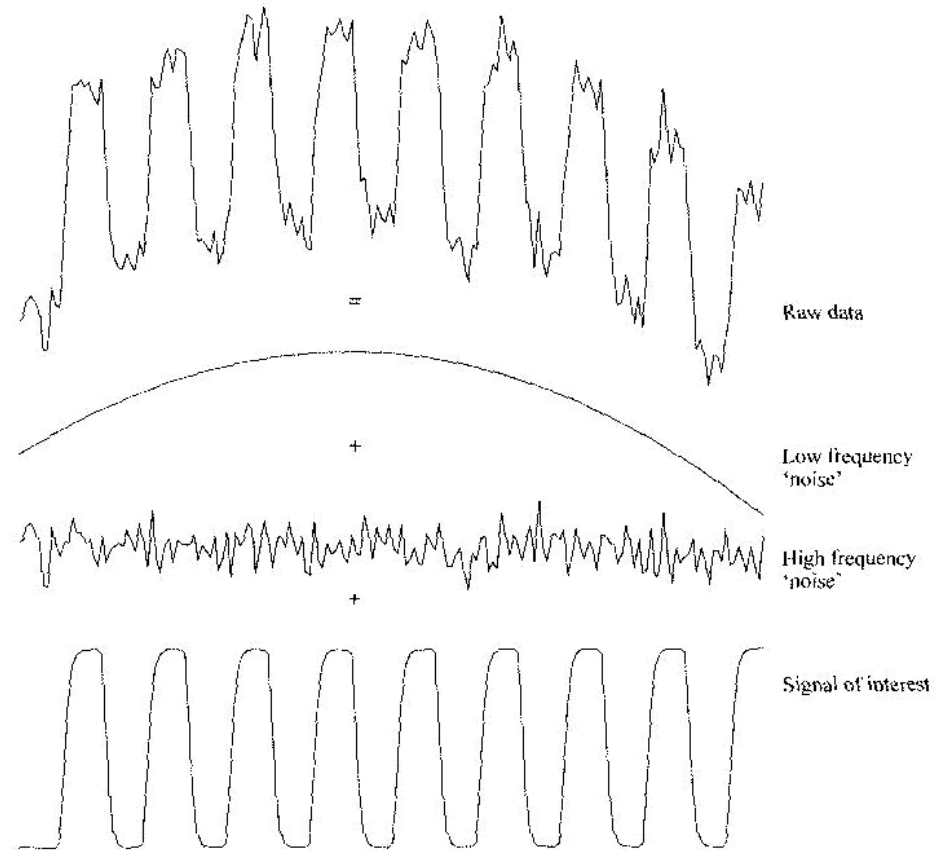
# Scaling

- Magnitude of BOLD fluctuation  $\propto$  baseline level
- Scaling time series of each voxel and for each run by the mean of all TRs of that run
  - Percentage
  - Z-score
- For proper comparison of statistics between runs

# Voxelwise Scaling

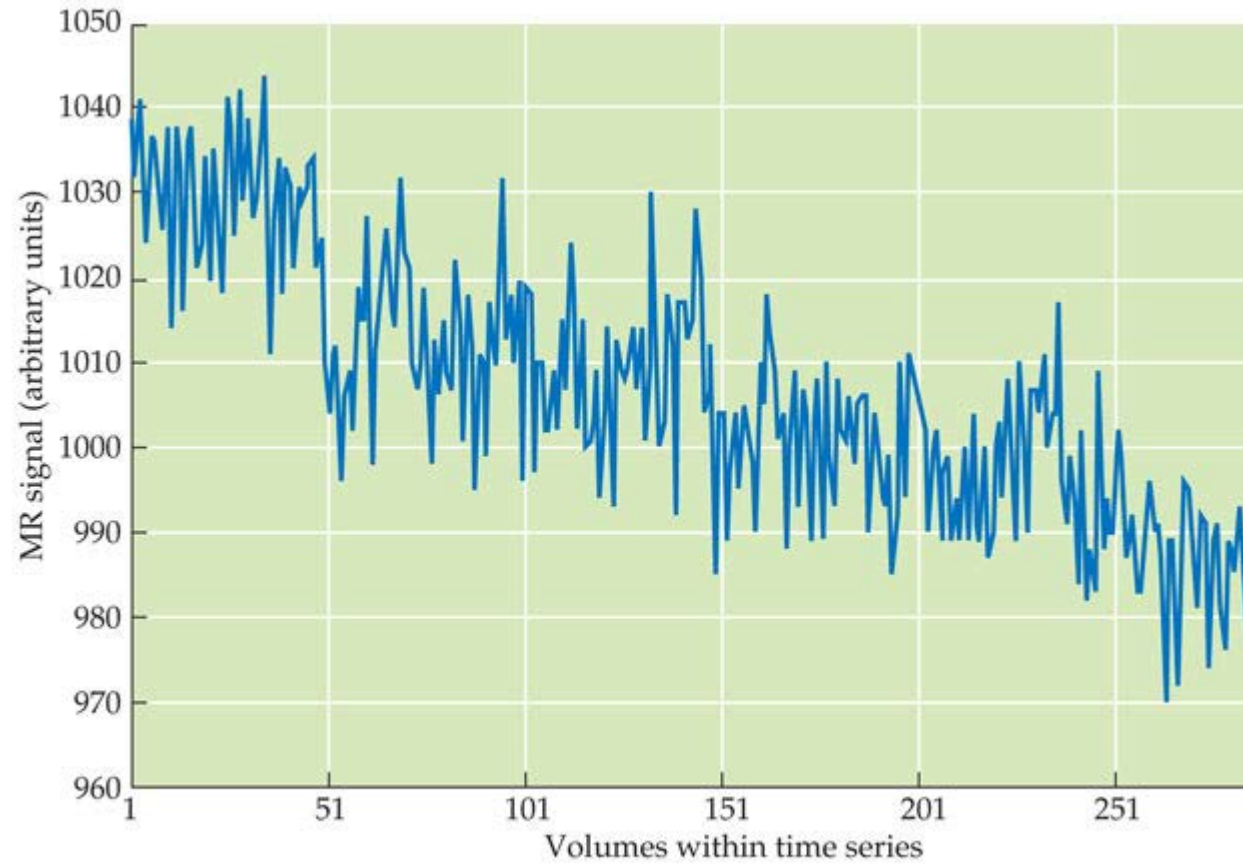


# Components of Time Course Data





# Linear Drift



FUNCTIONAL MAGNETIC RESONANCE IMAGING, Figure 9.7 © 2004 Sinauer Associates, Inc.

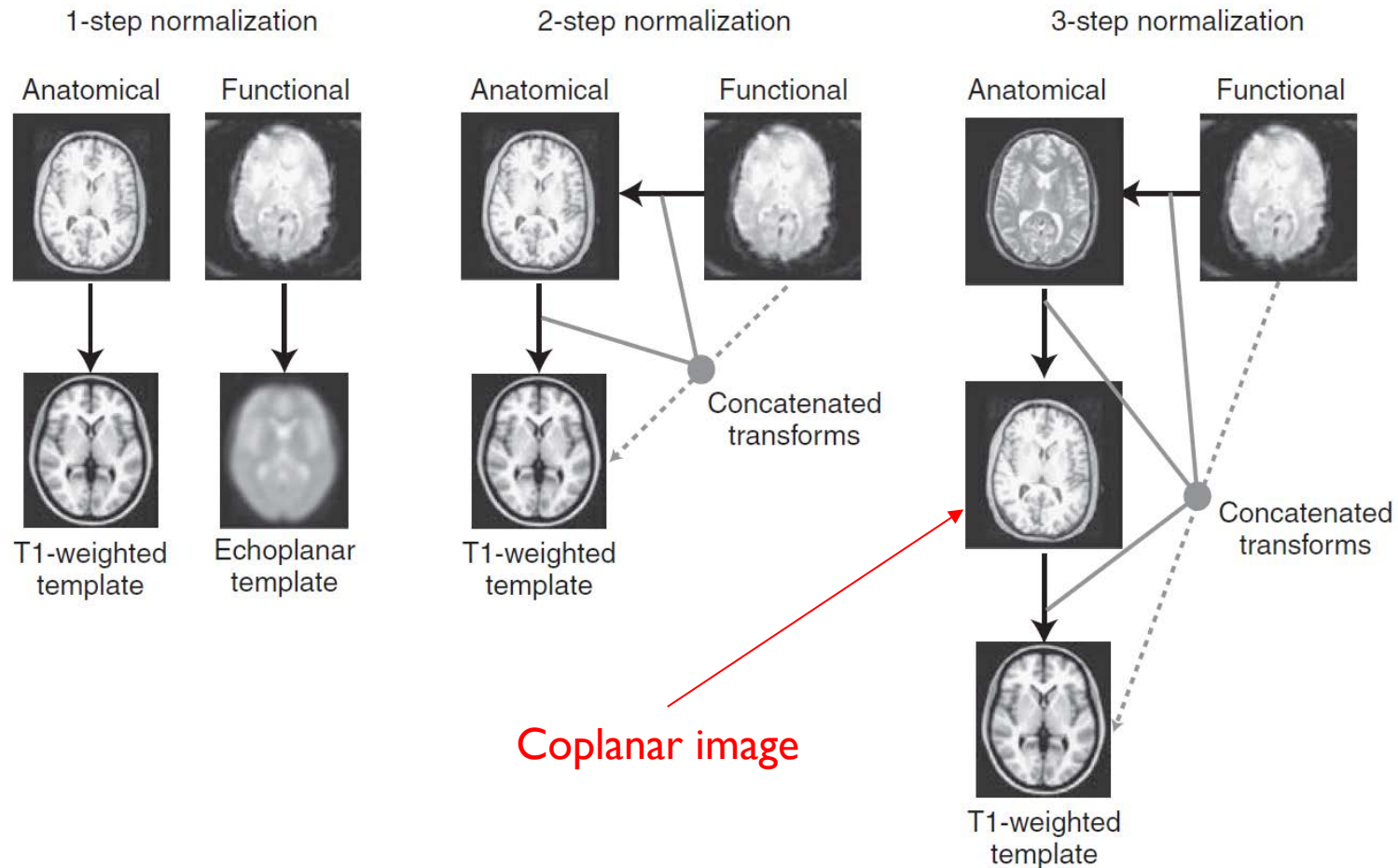
# Spatial Normalization

- Process of spatially transforming data into a common space for analysis
  - Aka. Intersubject registration
- Necessary for integrating results from multiple individuals
  - generalization

# Prestatistics Approach

- Compute the 1<sup>st</sup> level GLM for all subjects
  - Preprocessing → GLM
- Spatial normalization of 1<sup>st</sup> level statistical outputs

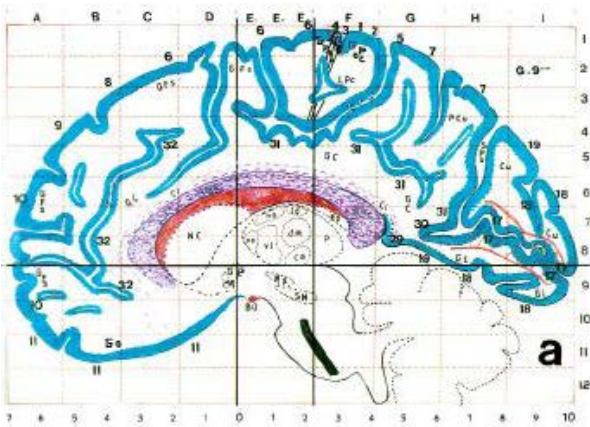
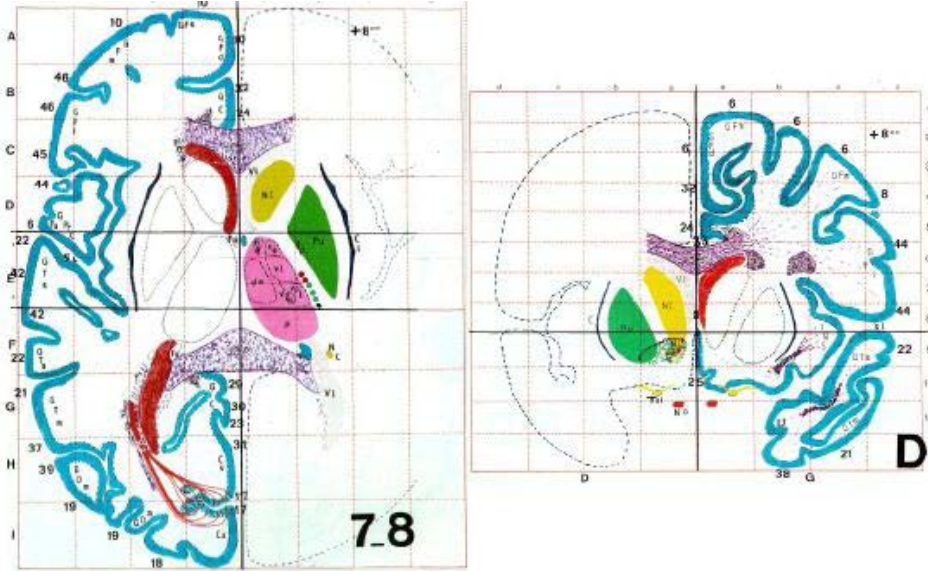
# Approaches in Operation of Spatial Normalization



# Atlas and Templates

- Atlas
  - Providing a guide to the location of anatomical features in a coordinate space
  - E.g., Talairach atlas
- Template
  - An image representative of the atlas
  - Providing a target where individual image can be aligned
  - E.g., MNI305, ICBM-152, ... etc

# Talairach Atlas



Talairach Client

File Edit Help

Database Search Options

Single Point  Nearest Gray Matter  Cube Range

Coordinates

Single Coordinate Search:

X:  Y:  Z:

From file:  Show Results

Input: <no file selected>

Output: <no file selected>

Search Clear

Welcome to the Talairach Client.

Gray Matter nearest to (0, 0, 0):  
Right Cerebrum, Limbic Lobe, Anterior Cingulate, Gray Matter, Brodmann area 25, Range=4

# Spatial Normalization Methods

- Landmark-based

- E.g., Talaraich Landmarks

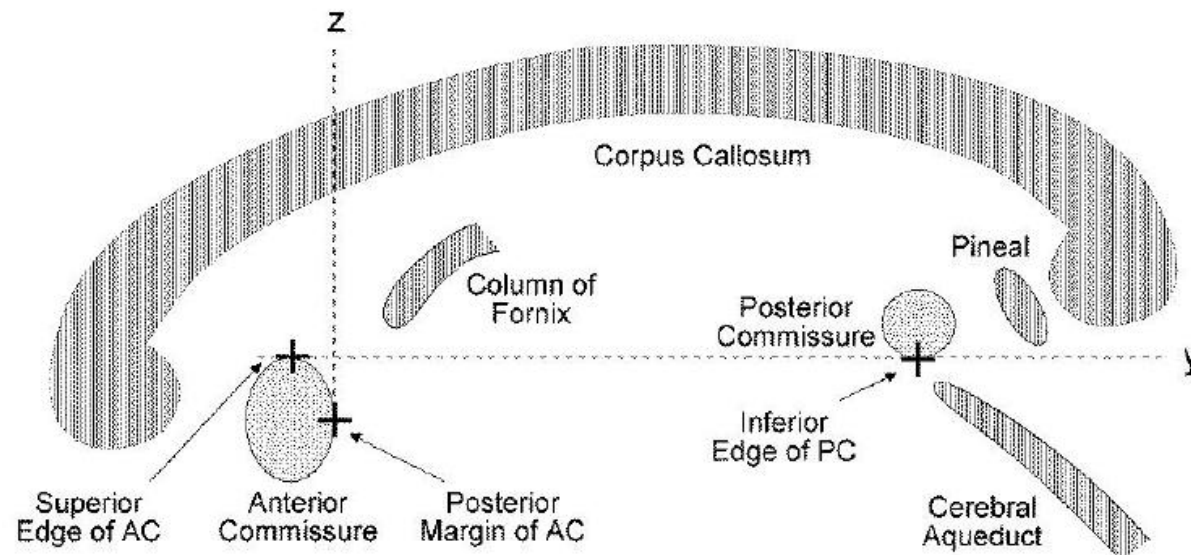
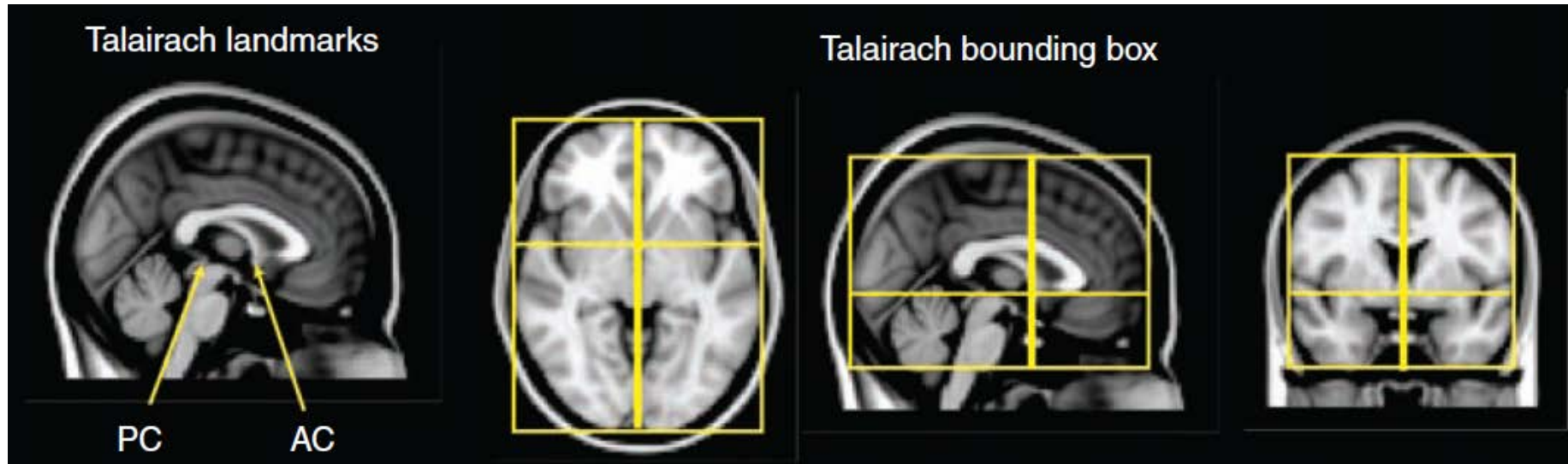
- Anterior and posterior commissures, midline sagittal plane, and the exterior boundaries of the brain in each direction

- Volume-based

- Surface-based



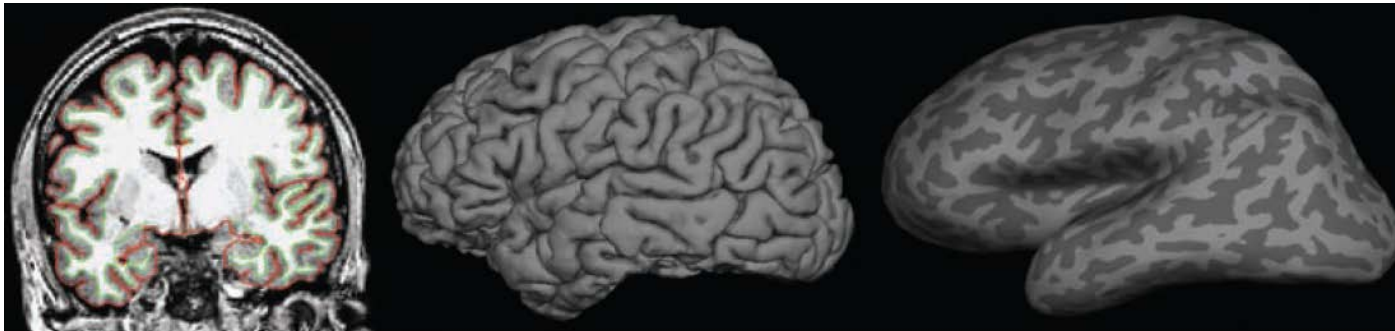
# Landmark Based





# Surface based

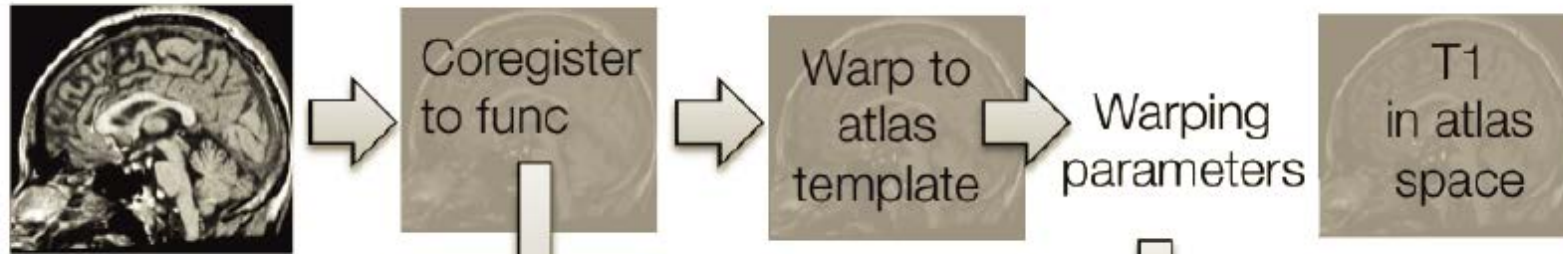
- Extraction of cortical surface



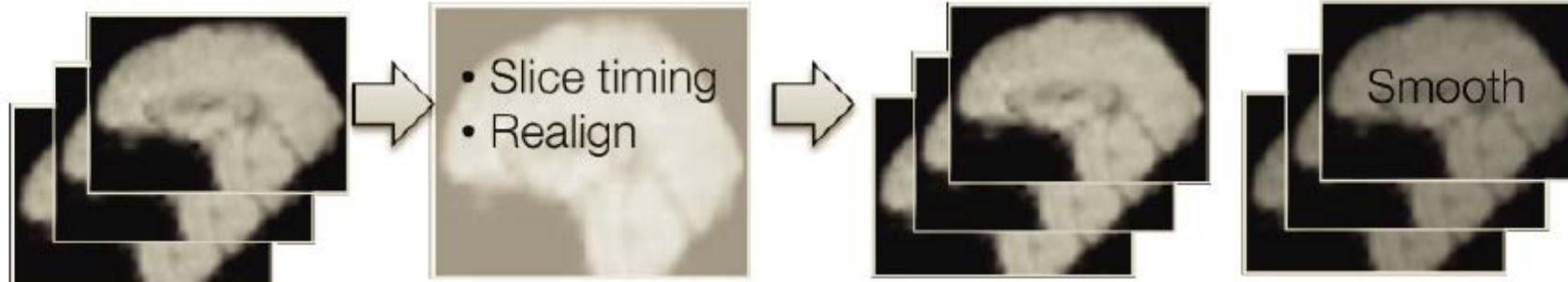
- Registration to surface atlas
  - More accurate registration of cortical features
  - Not ready for subcortical structures yet

# Summary

Structural (T1)



Functional image time series



Apply

# Questions?

# Matrix Expression of GLM

$$Y = X \cdot \beta + \varepsilon$$

- Write out equation for each observation of variable Y from 1 to J:

$$Y_1 = X_{11}\beta_1 + \dots + X_{1l}\beta_l + \dots + X_{1L}\beta_L + \varepsilon_1$$

$$Y_j = X_{j1}\beta_1 + \dots + X_{jl}\beta_l + \dots + X_{jL}\beta_L + \varepsilon_j$$

$$Y_j = X_{j1}\beta_1 + \dots + X_{jl}\beta_l + \dots + X_{jL}\beta_L + \varepsilon_j$$

Can turn these simultaneous equations into matrix form to get a single equation:

$$\begin{pmatrix} Y_1 \\ Y_j \\ Y_J \end{pmatrix} = \begin{pmatrix} X_{11} & \dots & X_{1l} & \dots & X_{1L} \\ X_{j1} & \dots & X_{jl} & \dots & X_{jL} \\ X_{J1} & \dots & X_{Jl} & \dots & X_{JL} \end{pmatrix} \begin{pmatrix} \beta_1 \\ \beta_j \\ \beta_J \end{pmatrix} + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_j \\ \varepsilon_J \end{pmatrix}$$

$$Y = X \cdot \beta + \varepsilon$$

Observed data

Design Matrix

Parameters

Residuals/Error

# Solution to the Equation

$$X'Y = X'X\beta \qquad \hat{\sigma}^2 = \frac{e'e}{T-(p+1)}$$

Any  $\beta$  satisfies the normal equation minimizes the sum of the squares of residuals ( $e'e$ )

$$\hat{\beta} = (X'X)^{-1}X'Y$$

 Assuming this is invertible

# Hypothesis Testing: Contrast t-test

$$c\hat{\beta} \sim N(0, c(X'X)^{-1}c'\sigma^2)$$

$$t = \frac{c\hat{\beta}}{\sqrt{c(X'X)^{-1}c'\hat{\sigma}^2}}$$

$$df: T - (p + 1)$$

$$H_A : c\beta > 0$$

$$P(T_{T-(p+1)} \geq t)$$

$$H_A : c\beta \neq 0$$

$$P(T_{T-(p+1)} \geq |t|)$$

$$t = \frac{c\hat{\beta}}{\sqrt{c(X'X)^{-1}c'\hat{\sigma}^2}}$$

Design matrix & Contrast Vector;  
depending on your experimental  
design

Residual error unaccounted for  
by your design; depending on  
the quality of data