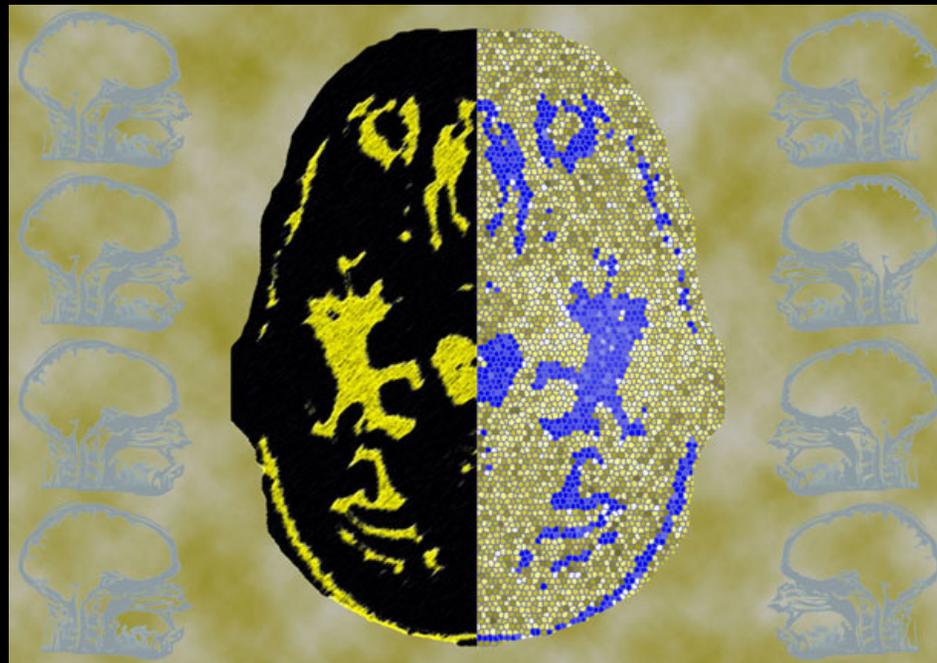


# Characterizing task-related interactions between brain regions with fMRI

*Functional and effective connectivity approaches*



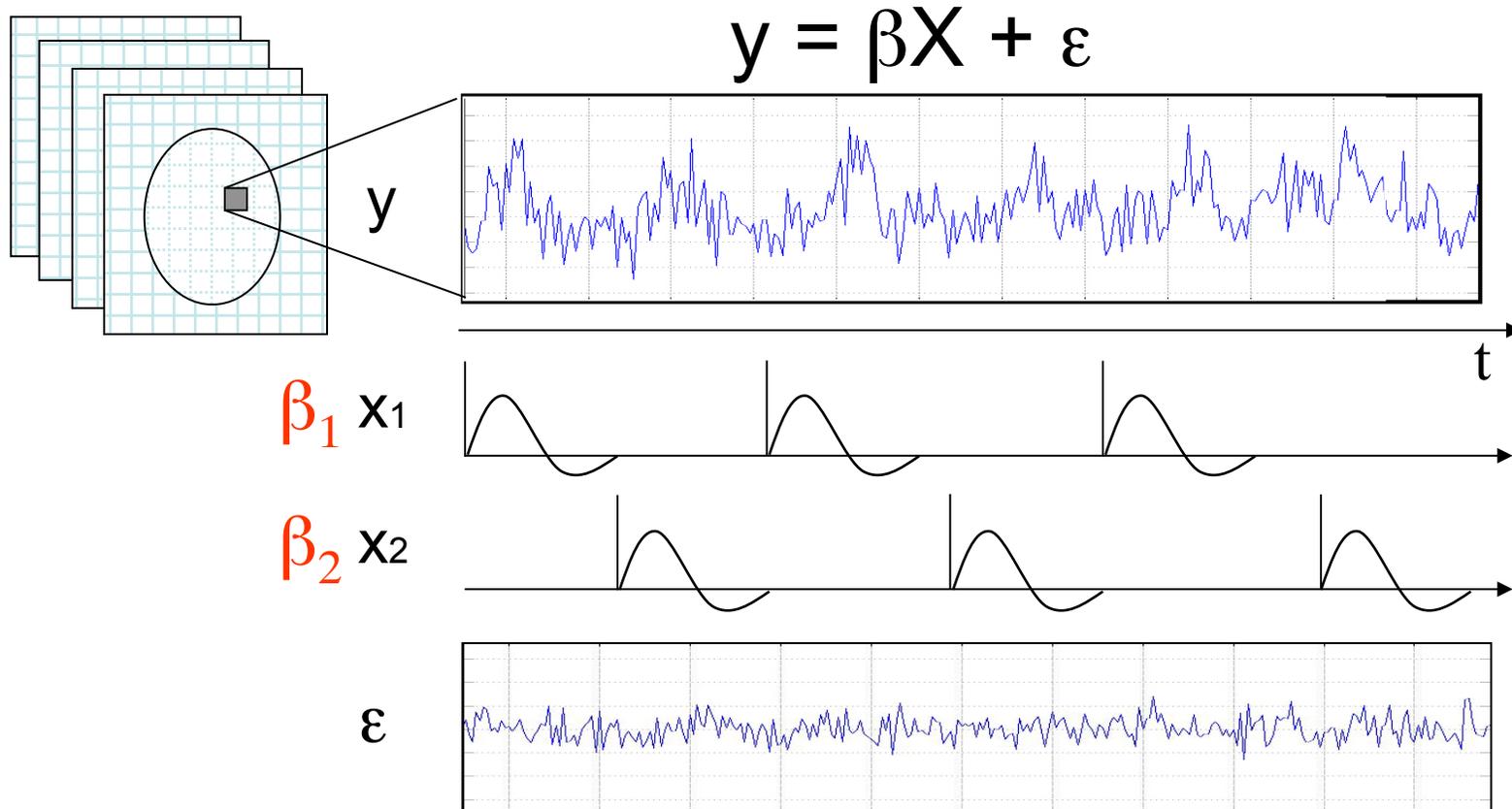
**Jesse Rissman, Ph.D.**

Dept. of Psychology

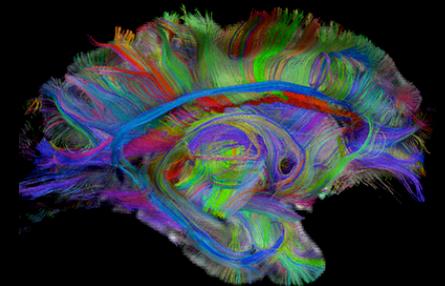
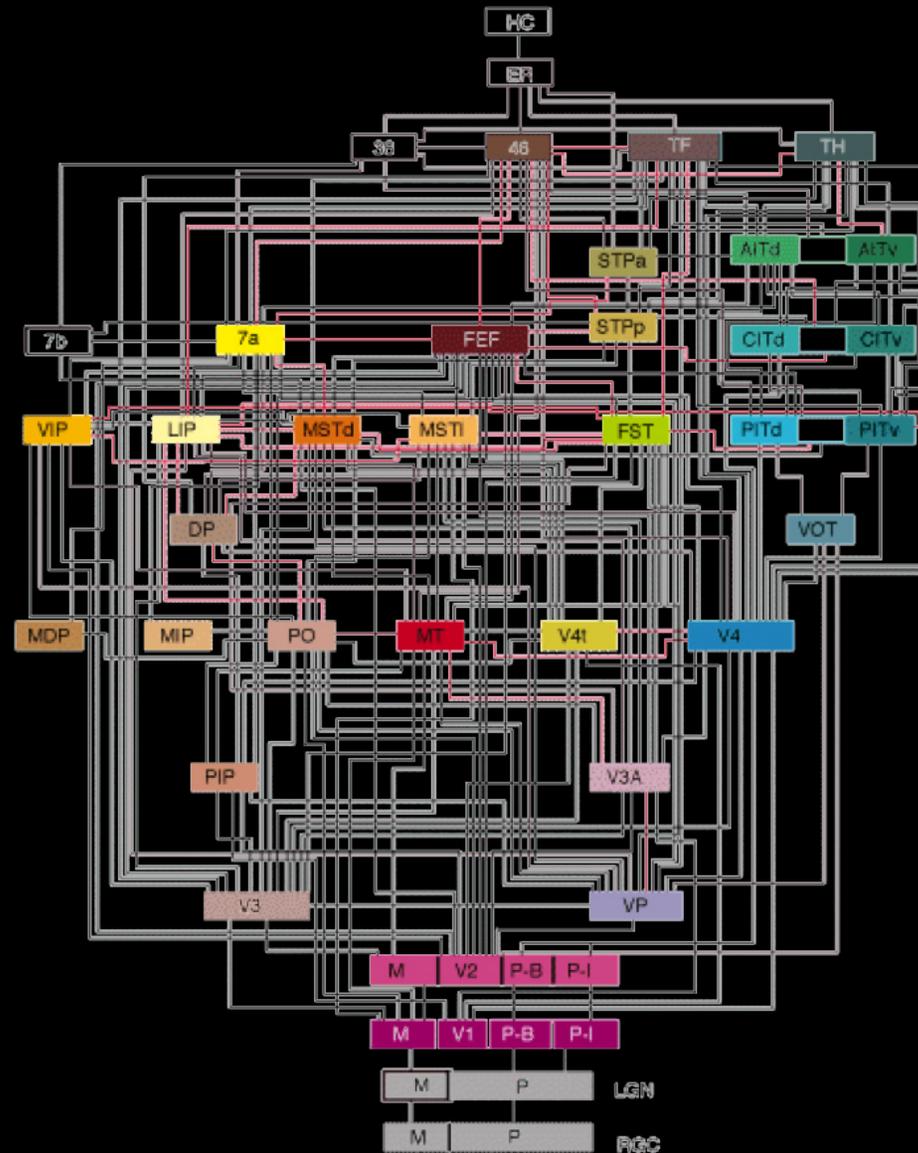
UCLA

# Mapping brain function with fMRI:

*Traditional univariate GLM approach to functional localization*



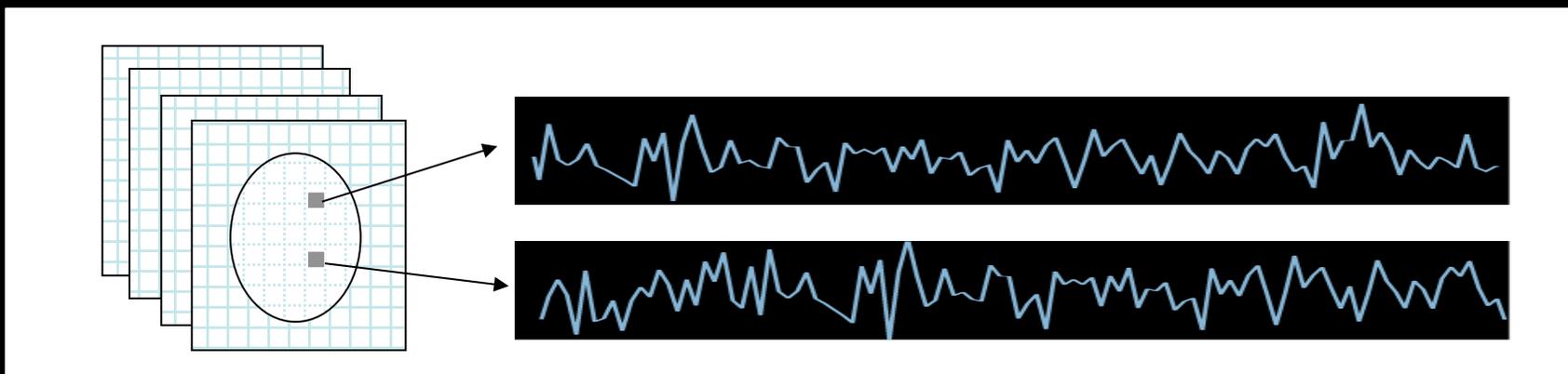
But brain regions do not act in isolation...



Felleman & Van Essen, 1991

# Mapping brain function with fMRI:

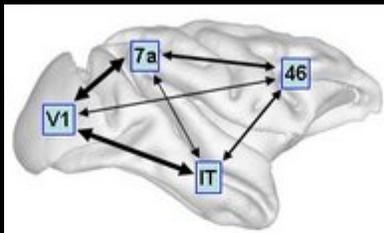
## *Functional connectivity approach*



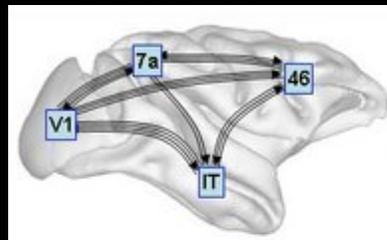
- Early functional connectivity analyses examined temporal correlations during the resting state (e.g., Biswal et al., 1995)
  - This approach continues to an extremely useful assay of inter-regional coupling
- Later studies began to examine task-dependent modulation of inter-regional coupling
  - Attending to motion vs. not attending to motion (Friston et al., 1997)
  - 2-back vs. 0-back working memory task (Lowe et al., 2000)
  - Listening to continuous speech vs. resting (Hampson et al., 2002)
- A growing collection of methods have been developed to extract information about functional interactions from fMRI data

# Types of connectivity

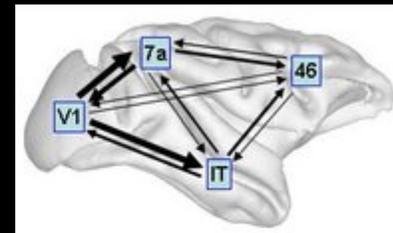
## Functional Connectivity



## Structural Connectivity



## Effective Connectivity



**Documenting the correlation between spatially remote neurophysiological events**

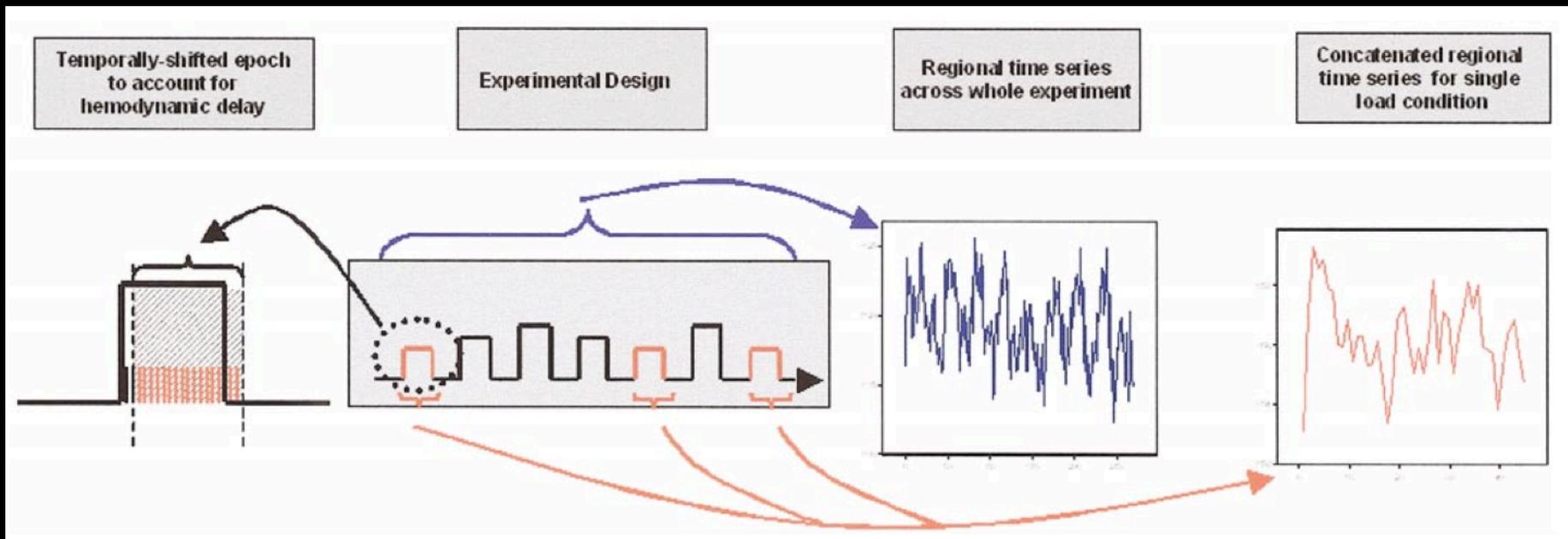
**Characterizing the influence that one neuronal system exerts over another**

- Data-driven / exploratory
  - Bivariate approaches (seed-based or ROI-based)
    - Timeseries correlations
    - Beta series correlations
  - Multivariate approaches
    - PCA, ICA, PLS
- Hypothesis-driven / model-based
  - Psychophysiological interactions (PPI)
  - Structural equation modeling (SEM)
  - Dynamic causal modeling (DCM)
  - Granger causality

# Functional connectivity

## *Time series correlations*

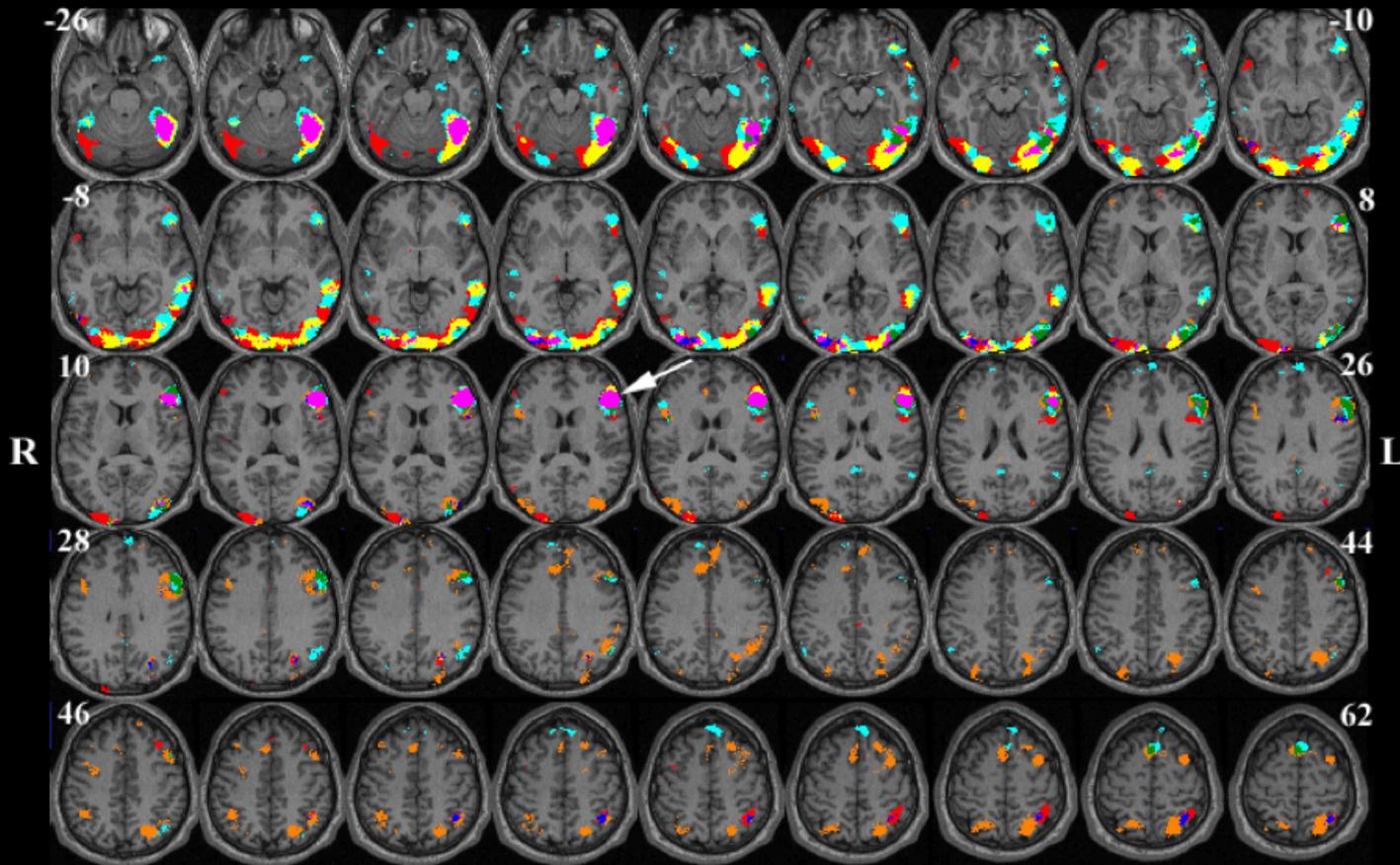
- Similar to resting state analyses, except that correlations are computed on data collected during task performance
  - *Requires that experiment utilizes a block design*



# Functional connectivity

## Time series correlations

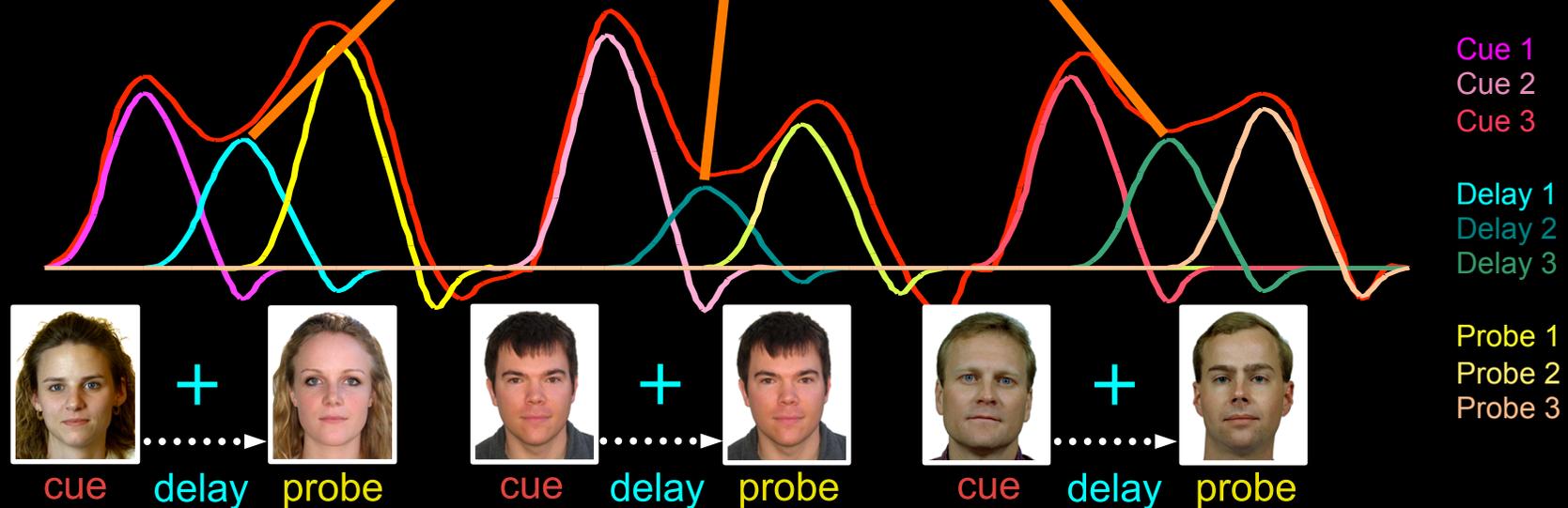
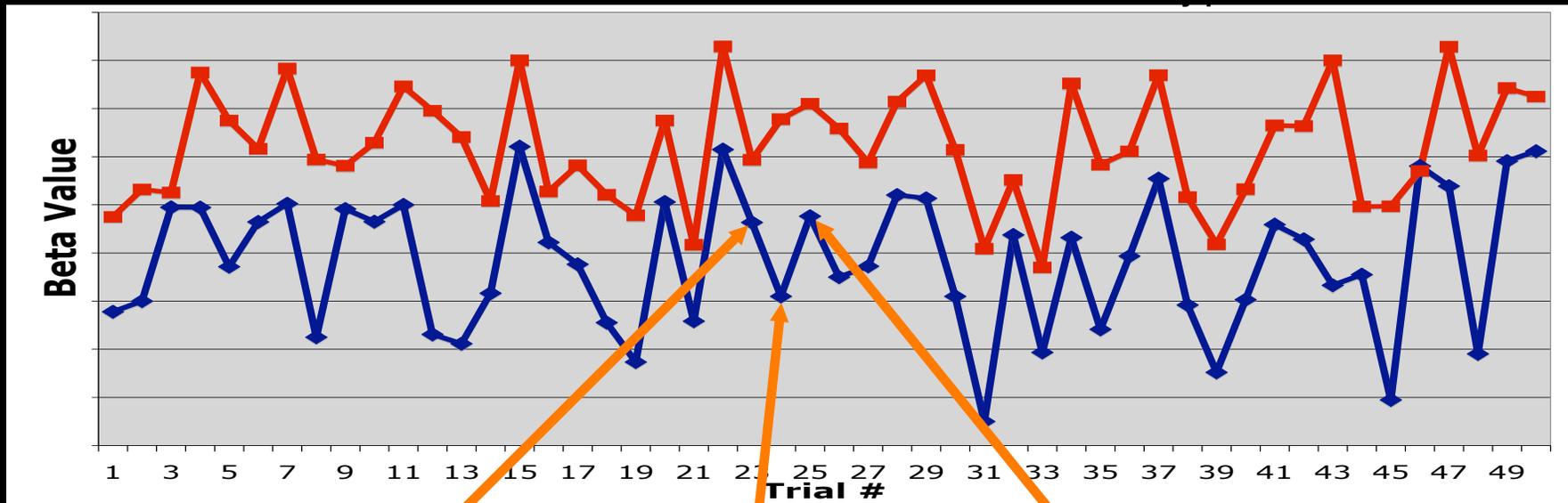
Functional Connectivity Map - Reference Voxel in BA44/45 (-50 28 16)



Color Legend: words, pseudowords & letter-strings  
words & pseudowords  
words & letter-strings  
pseudowords & letter-strings

words  
pseudowords  
letter-strings

# Measuring functional connectivity during distinct task stages: *Beta series correlation analysis*



Rissman, Gazzaley, and D'Esposito (2004), *NeuroImage*

# Beta series correlations:

## *Methodological validation*

- Beta series correlation analysis method applied to simple bimanual motor task.
- In the **Right-then-Left** condition, subjects played a sequence of 4 keystrokes with their right hand and then played a different sequence with their left.
- In the **Interleaved** condition, subjects played 8 keystrokes alternating between hands – a task requiring increased bimanual coordination.
- Hypothesis: The Interleaved condition should induce more inter-hemispheric cross-talk between motor regions.

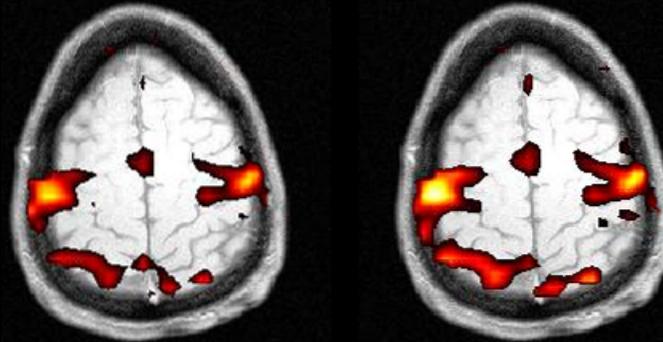
# Beta series correlations:

*A meaningful metric of inter-regional coupling?*

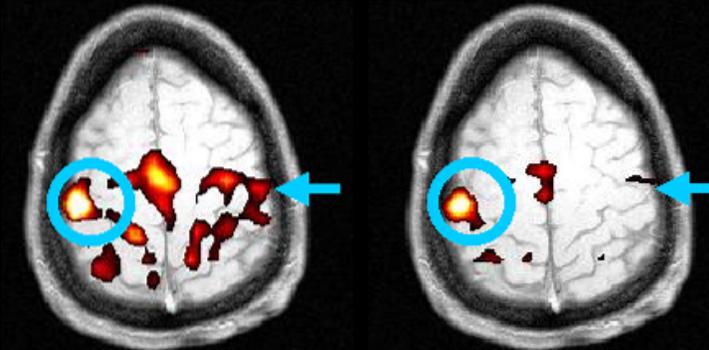
**Bimanual  
coordination**

**One hand  
at a time**

**Univariate**

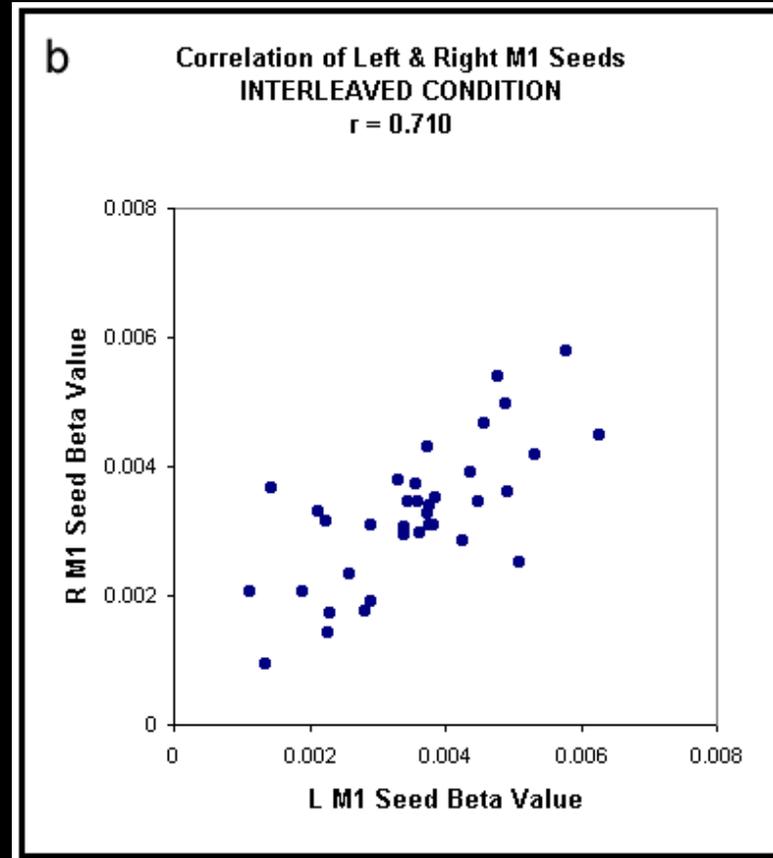
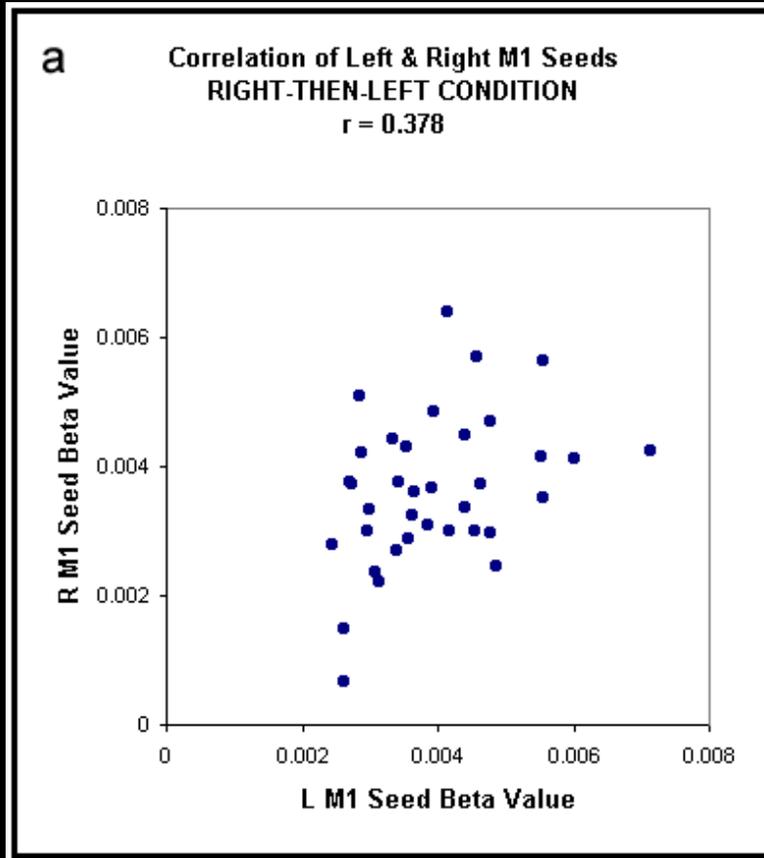
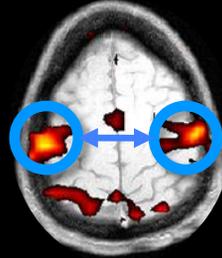


**Correlation**



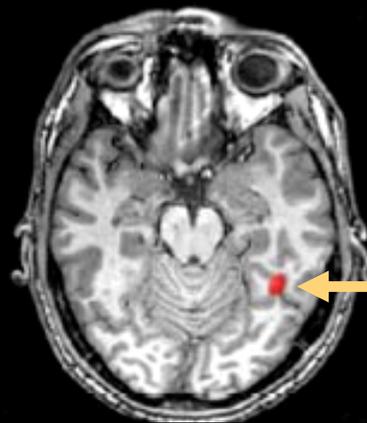
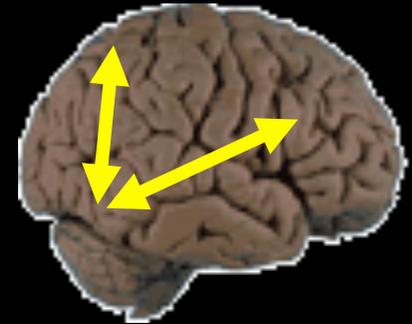
# Beta series correlations:

*A meaningful metric of inter-regional coupling?*



# Beta series correlation analysis applied to a basic visual working memory task

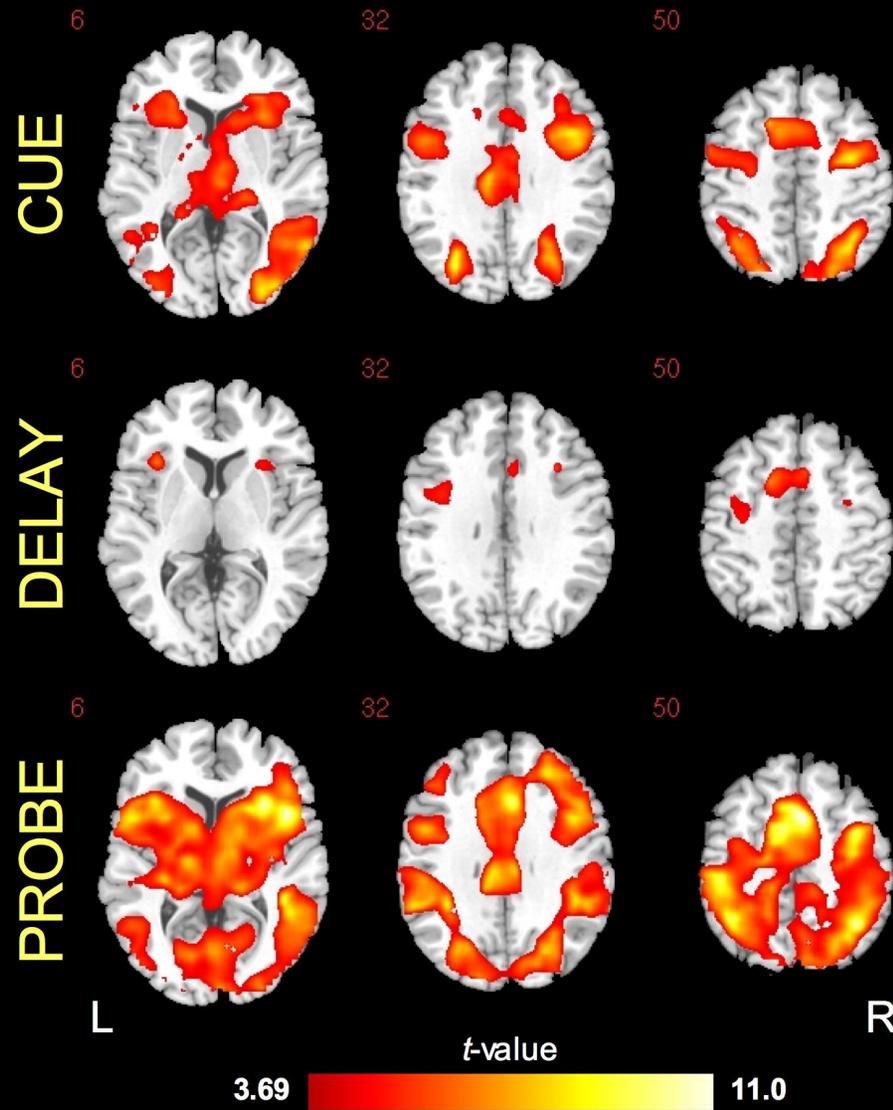
- Hypothesis: Frontoparietal regions interact with neural ensembles in inferotemporal cortex to keep behaviorally-relevant visual representations active
- Analysis performed on fMRI data from 17 subjects
- Task: maintain a single face across a 7-8 sec delay period



**right fusiform face area (FFA) “seed”**

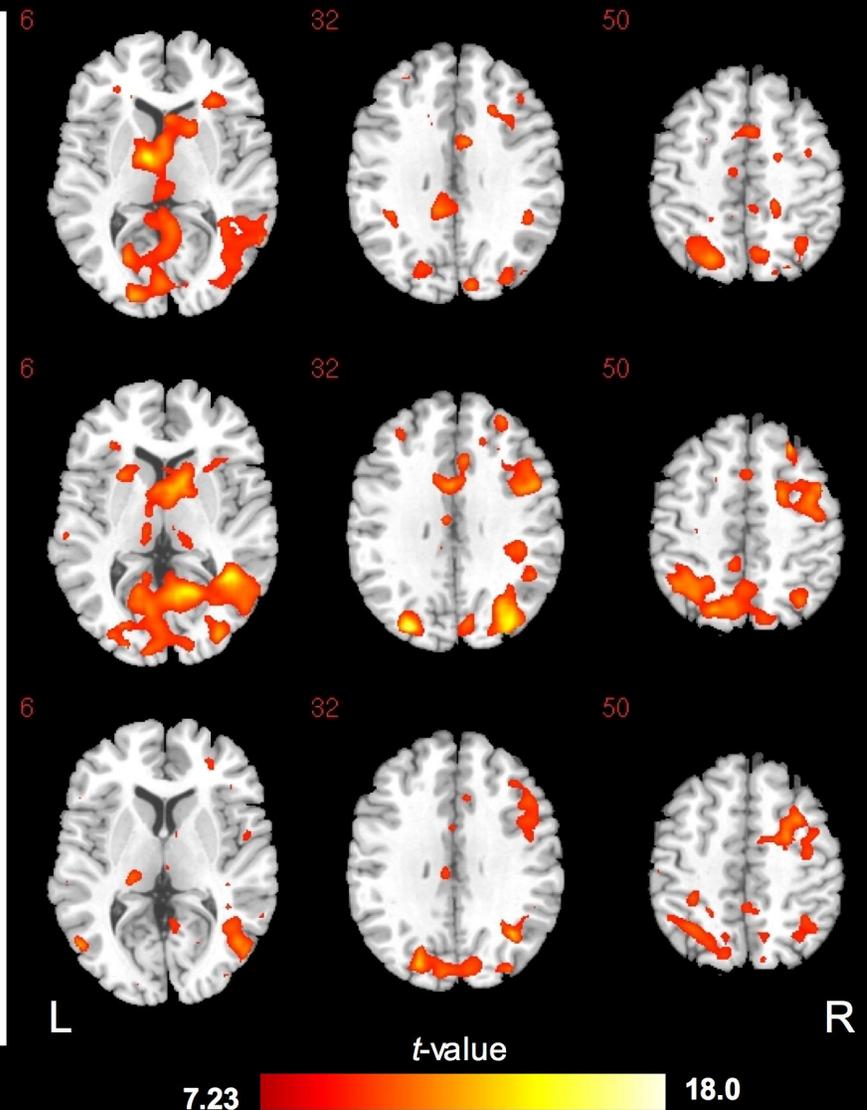
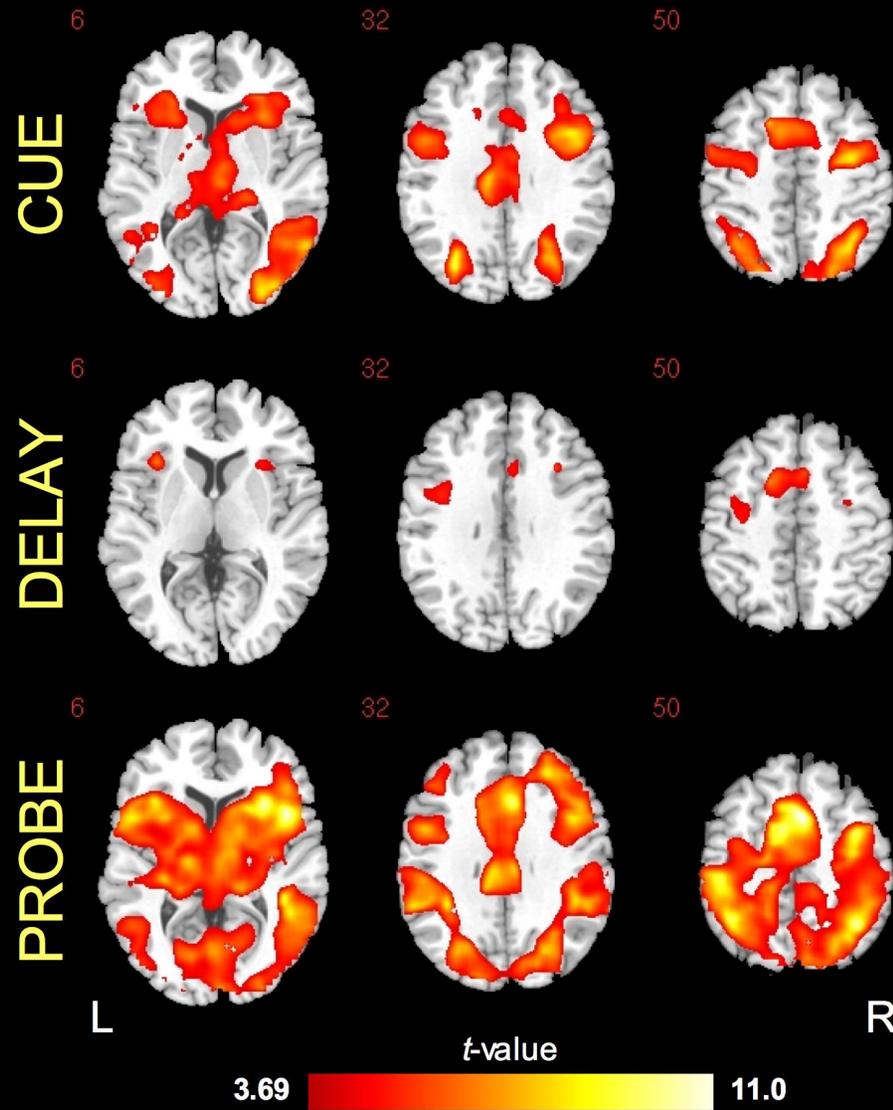
*Which brain regions are most strongly correlated with this seed region during face maintenance?*

# Univariate Activation



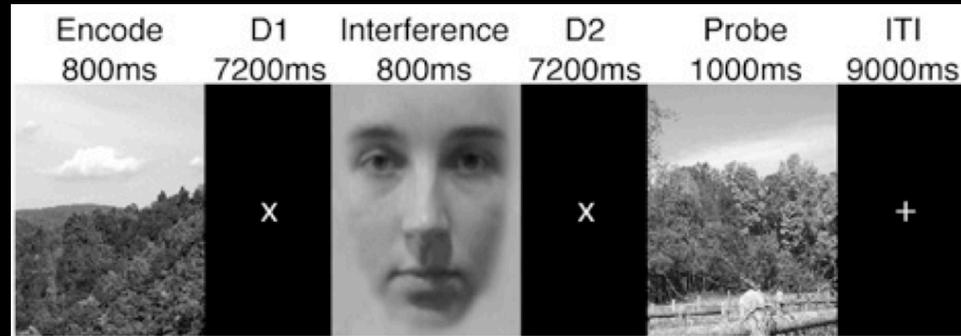
# Univariate Activation

# Right FFA Correlation



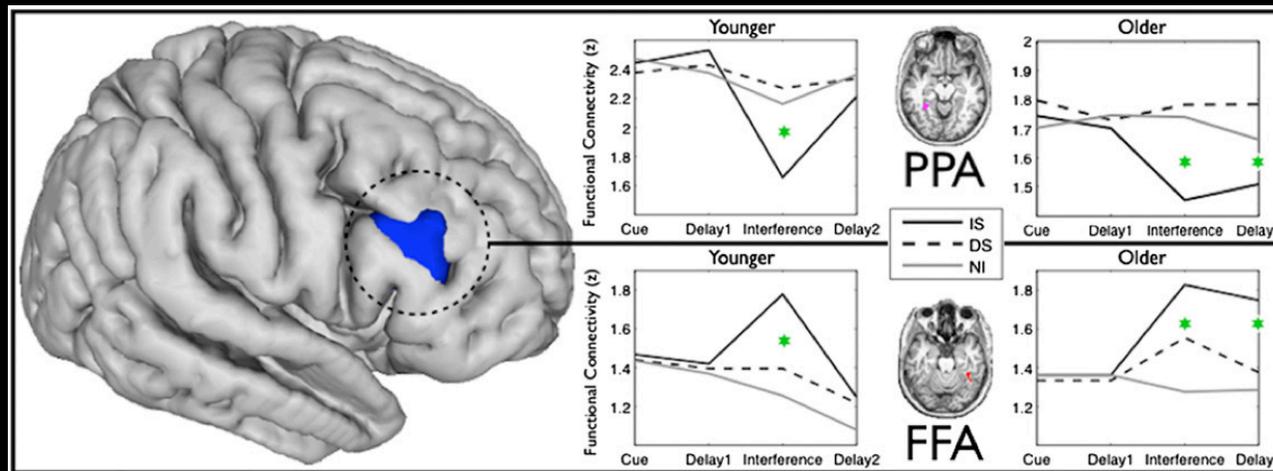
# Another example:

## Age-related changes in prefrontal coupling



### 3 task conditions:

- Interrupting stimulus (IS): *make judgment about face (male over 40?)*
- Distracting stimulus (DS): *ignore face; no decision required*
- No interference (NI): *no face stimulus presented*



Older adults failed to reestablish connectivity following interruptions!

# Pros & cons of beta series correlation method



- **Pros:**

- Can examine how functional interactions between regions evolve over the course of a multi-stage trial
- Relatively simple to implement (demo this afternoon)

- **Cons:**

- Single trial activity estimates can be quite noisy
- Serially-positioned HRF-convolved regressors may not provide ideal fit to data
- Not ideal for rapid, jittered event-related designs
- Cannot determine whether inter-regional correlations reflect direct or indirect communication

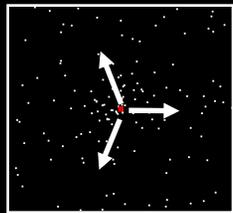
# Psychophysiological interaction (PPI)

- Bilinear model of how a psychological context **A** changes the influence of area **B** on area **C** :

$$B \times A \rightarrow C$$

- A PPI corresponds to a difference in regression slopes for different contexts
- Seed-based approach
  - PPI effects computed voxel-by-voxel across entire brain

# Psychophysiological interaction (PPI)



		Task factor	
		Task A <i>NO ATTENTION</i>	Task B <i>ATTENTION</i>
Stimulus factor	Stim 1 <i>STATIC DOTS</i>	$T_A/S_1$	$T_B/S_1$
	Stim 2 <i>MOVING DOTS</i>	$T_A/S_2$	$T_B/S_2$

GLM of a 2x2 factorial design:

$$y = (T_A - T_B) \beta_1$$

← main effect of task

$$+ (S_1 - S_2) \beta_2$$

← main effect of stim. type

$$+ (T_A - T_B) (S_1 - S_2) \beta_3$$

← interaction

$$+ e$$

- Replace one main effect in the GLM by the time series of an area that shows this main effect
- e.g., swap out the main effect of stimulus type with the time series of area V1

$$y = (T_A - T_B) \beta_1$$

← main effect of task

$$+ V1 \beta_2$$

← V1 time series ≈ main effect of stim. type

$$+ (T_A - T_B) V1 \beta_3$$

← psycho-physiological interaction

$$+ e$$

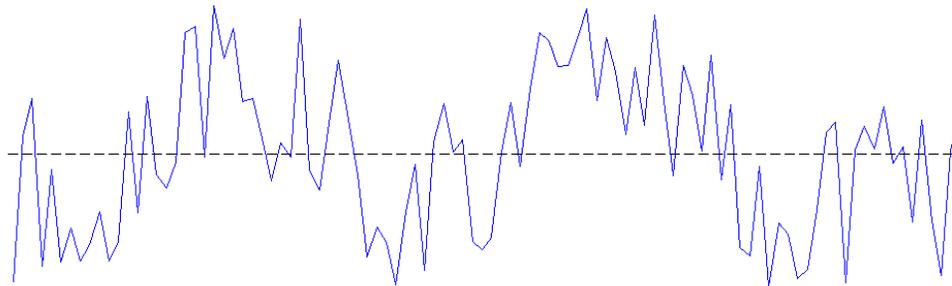
# Psychophysiological interaction (PPI)

**PSY main effect  
(task variable)**

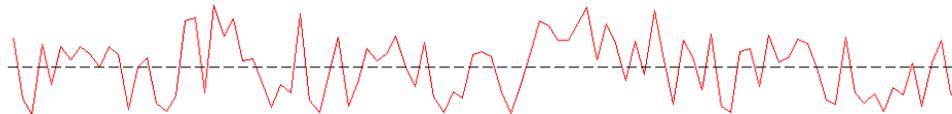


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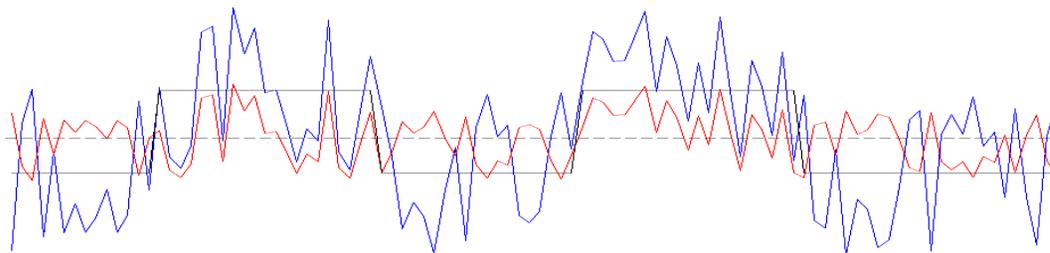
**PHYS main effect  
(time series from  
seed region)**



**PPI =  
PSY.\*PHYS**

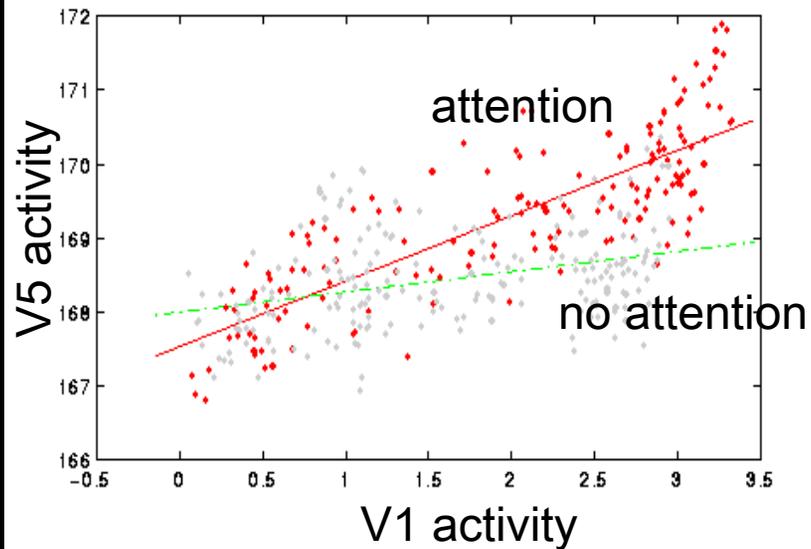
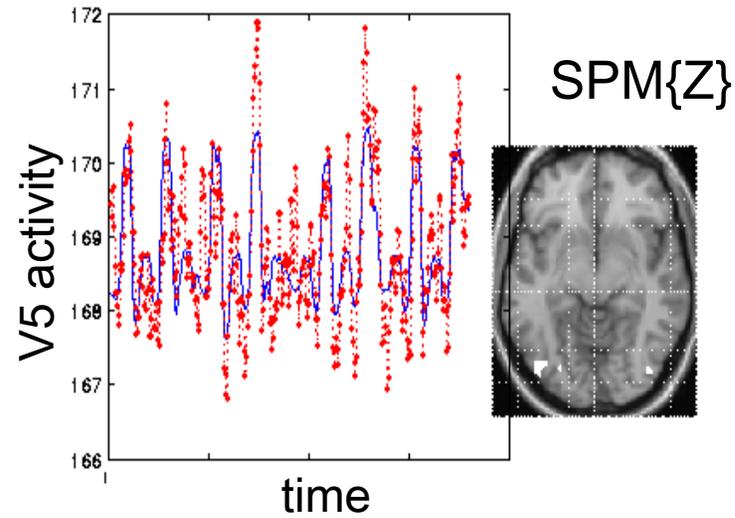


**Overlay:**



# Psychophysiological interaction (PPI)

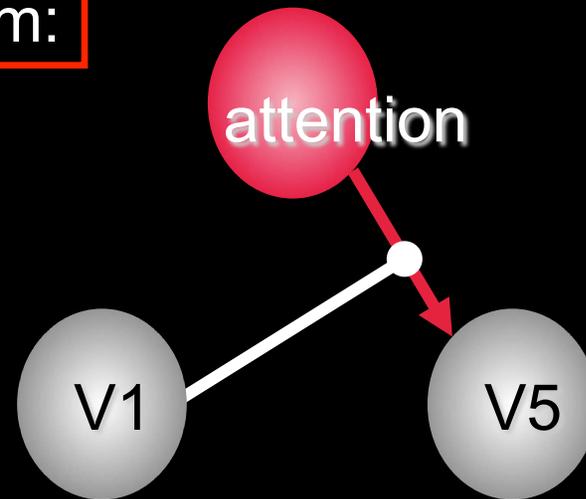
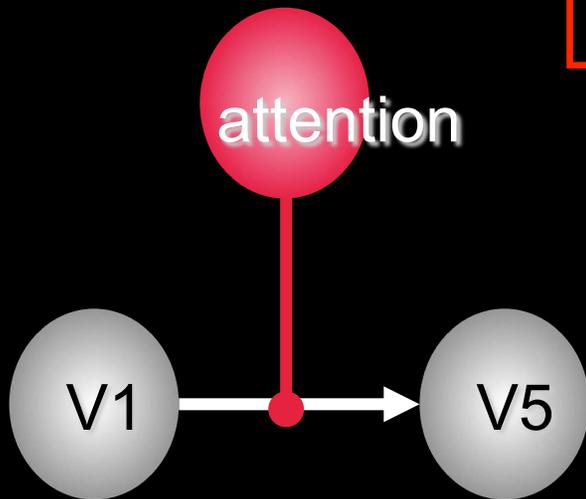
Friston et al. (1997), *NeuroImage*



# Psychophysiological interaction (PPI)

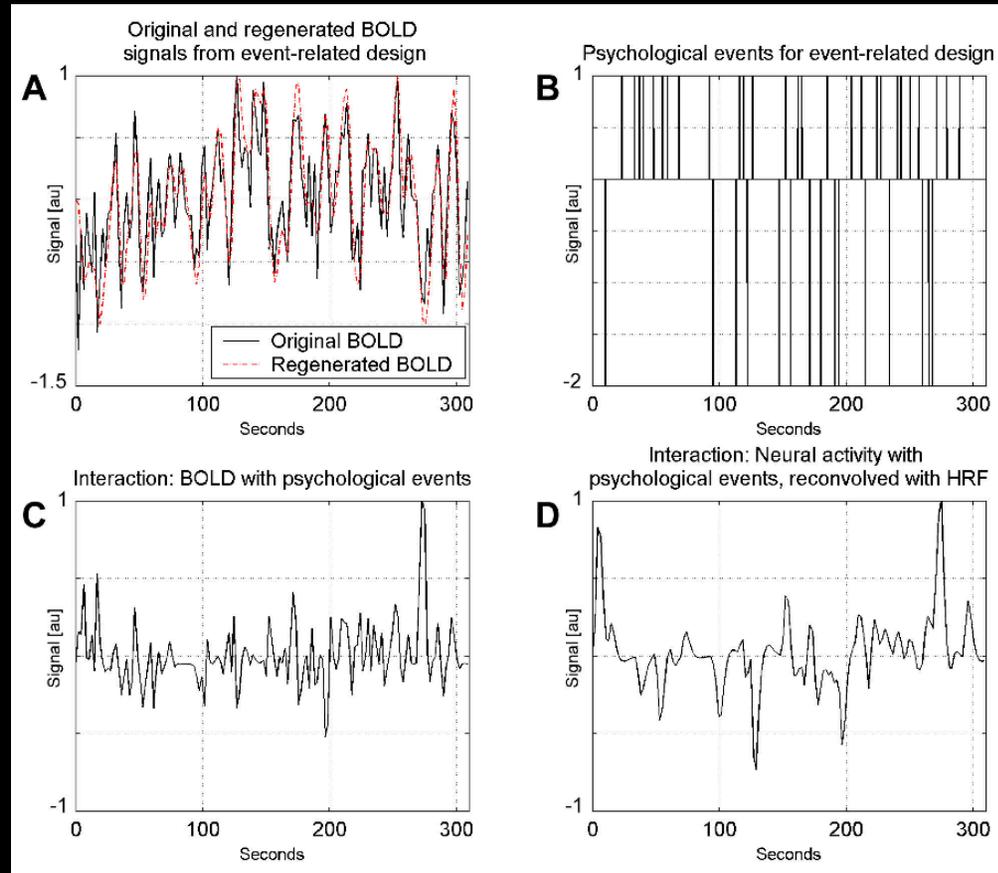
$$y = (T_A - T_B) \beta_1 \\ + V1\beta_2 \\ + (T_A - T_B)V1\beta_3 \\ + e$$

Two possible interpretations of the PPI term:



# PPI on event-related fMRI data

*The importance of hemodynamic deconvolution*



$$(A \otimes \text{HRF}) \times (B \otimes \text{HRF}) \neq (A \times B) \otimes \text{HRF}$$

Gitelman et al. (2004), *NeuroImage*

# Pros & cons of PPI analysis



- **Pros:**

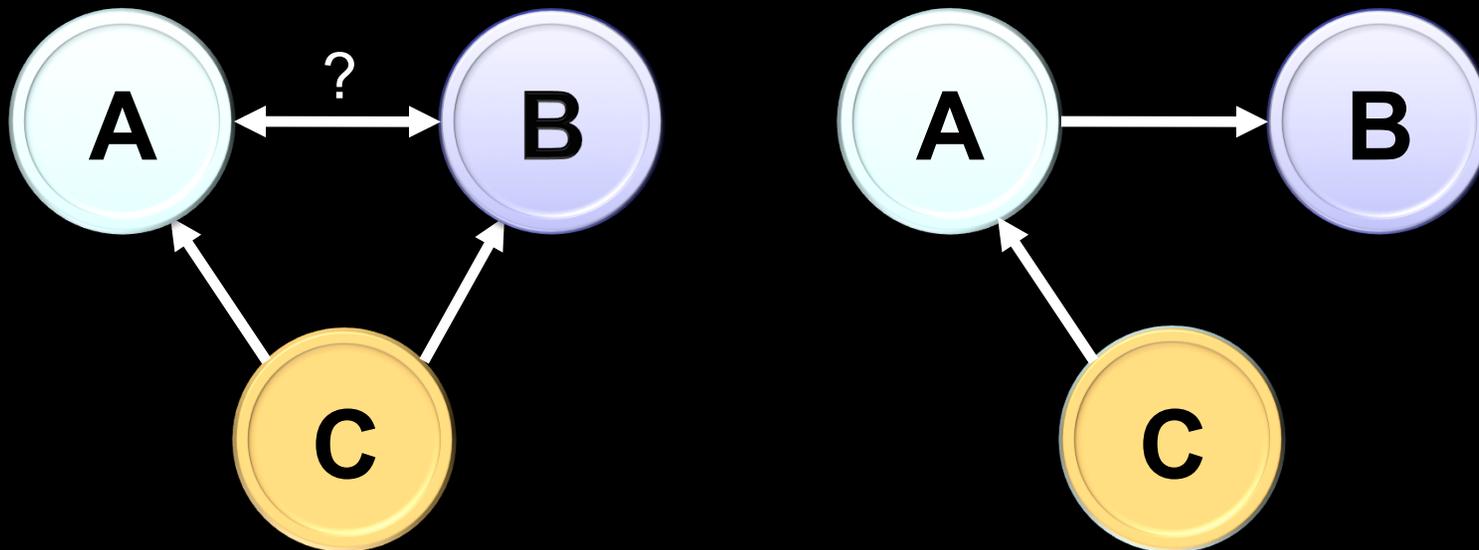
- Provides useful exploratory assay of how a given region's connectivity with the rest of the brain is modulated by task context
- Easy to implement (FSL demo this afternoon)

- **Cons:**

- Can only model contributions from a single area
- PPI regressor may be highly correlated with psychological task regressor, reducing power
  - Factorial designs help avoid this problem!
- Limited causal interpretability

## Inferring causality from fMRI data

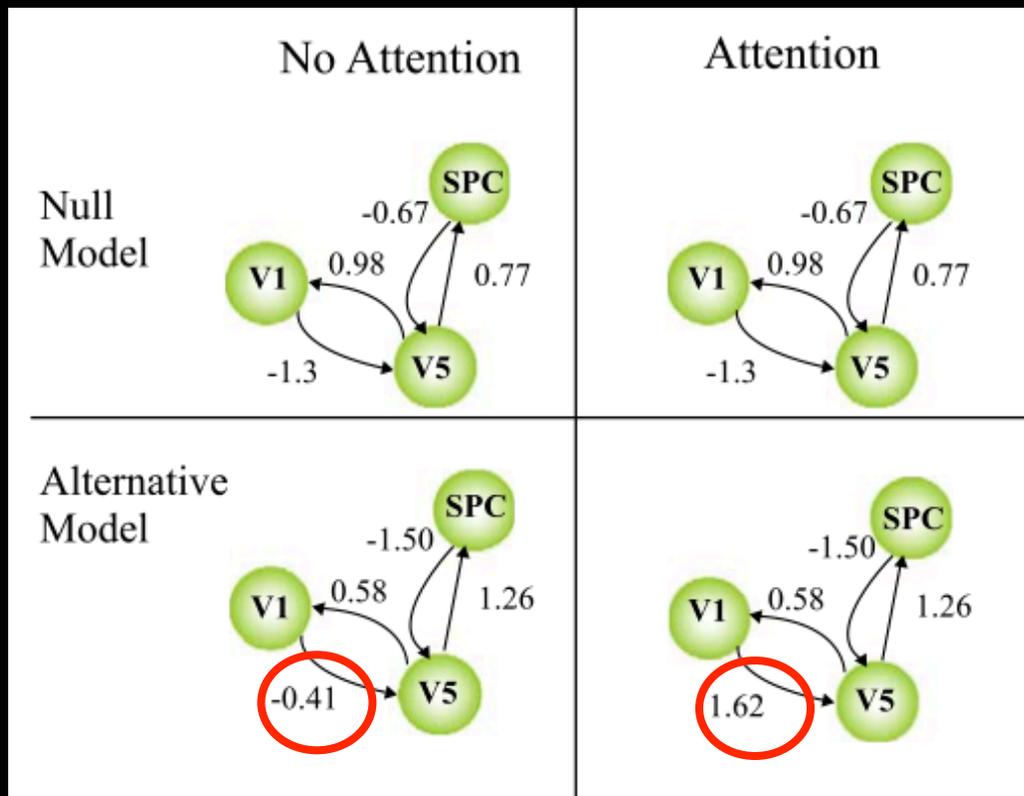
- How can we determine whether the apparent relationship between activity fluctuations in two regions is due to their *direct* communication or the *indirect* influence of another region?



# Structural Equation Modeling (SEM)

- A technique for characterizing the causal relationship between variables based on their covariance structure
  - Widely used in the social sciences; Also referred to as “path analysis”
  - First applied to PET data (McIntosh & Gonzalez-Lima, 1992,1994) and later extended to fMRI data (Büchel & Friston, 1997)
- Requires specification of anatomical graph model indicating hypothesized connections between brain regions
  - Connection paths are unidirectional (reciprocal connections allowed)
  - Model should be motivated by anatomical data (tracer studies / DTI)
- Parameters of the model are estimated by minimizing the difference between the *observed* covariances and the covariances *implied* by the structural model
  - Compare goodness of fits across models

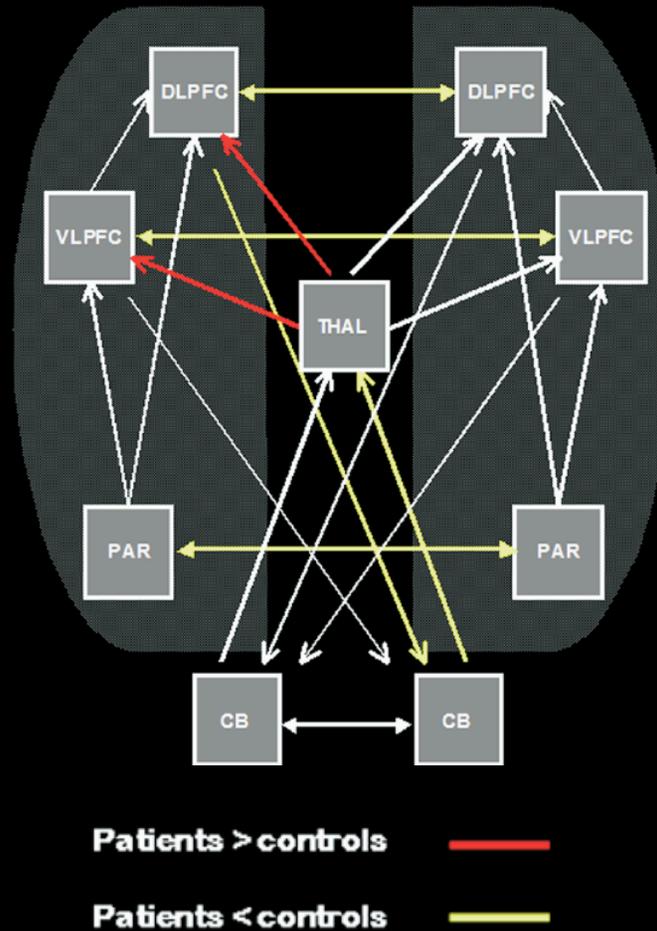
# Assessing effects of task context



Penny et al. (2004), *NeuroImage*

- Let's revisit the Attention-to-Motion data set
- How does attention influence connectivity of these nodes?
- Null model: constrain path coefficients to be identical
- Alternative model: allow path coefficients to vary across conditions
- Allowing  $V1 \rightarrow V5$  path to vary significantly improves goodness of fit

# Comparing path coefficients across populations



Schlösser et al. (2003), *NeuroImage*

# Pros and cons of SEM



## Pros:

- Can examine the functional interplay of many brain areas at once
- Allows inferences about the causal structure within a network

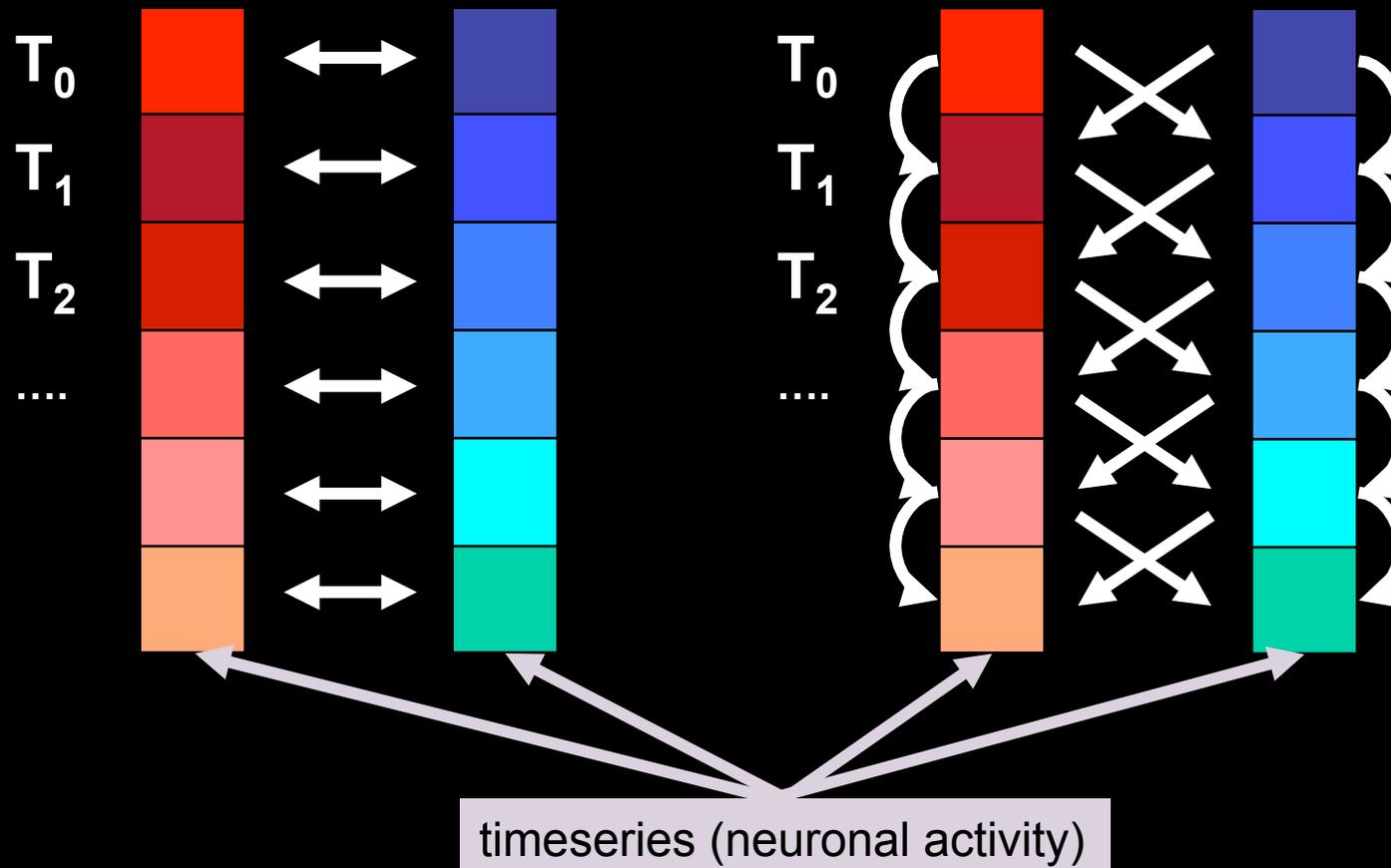
## Cons:

- Need to partition time series data to calculate condition-specific covariances
- No explicit inputs to system
  - use primary sensory regions and/or estimate residual influences
- Some models are not identifiable
  - especially when lots of reciprocal connections are allowed
- Searching for the best model can be unreliable
  - adding connections vs. pruning connections?
  - constrain model to be the same across subjects?

# From static to dynamic models...

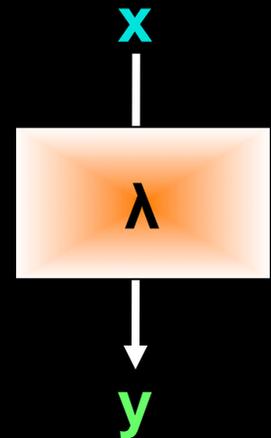
Seed voxel approach, PPI, SEM, etc.

Dynamic causal models



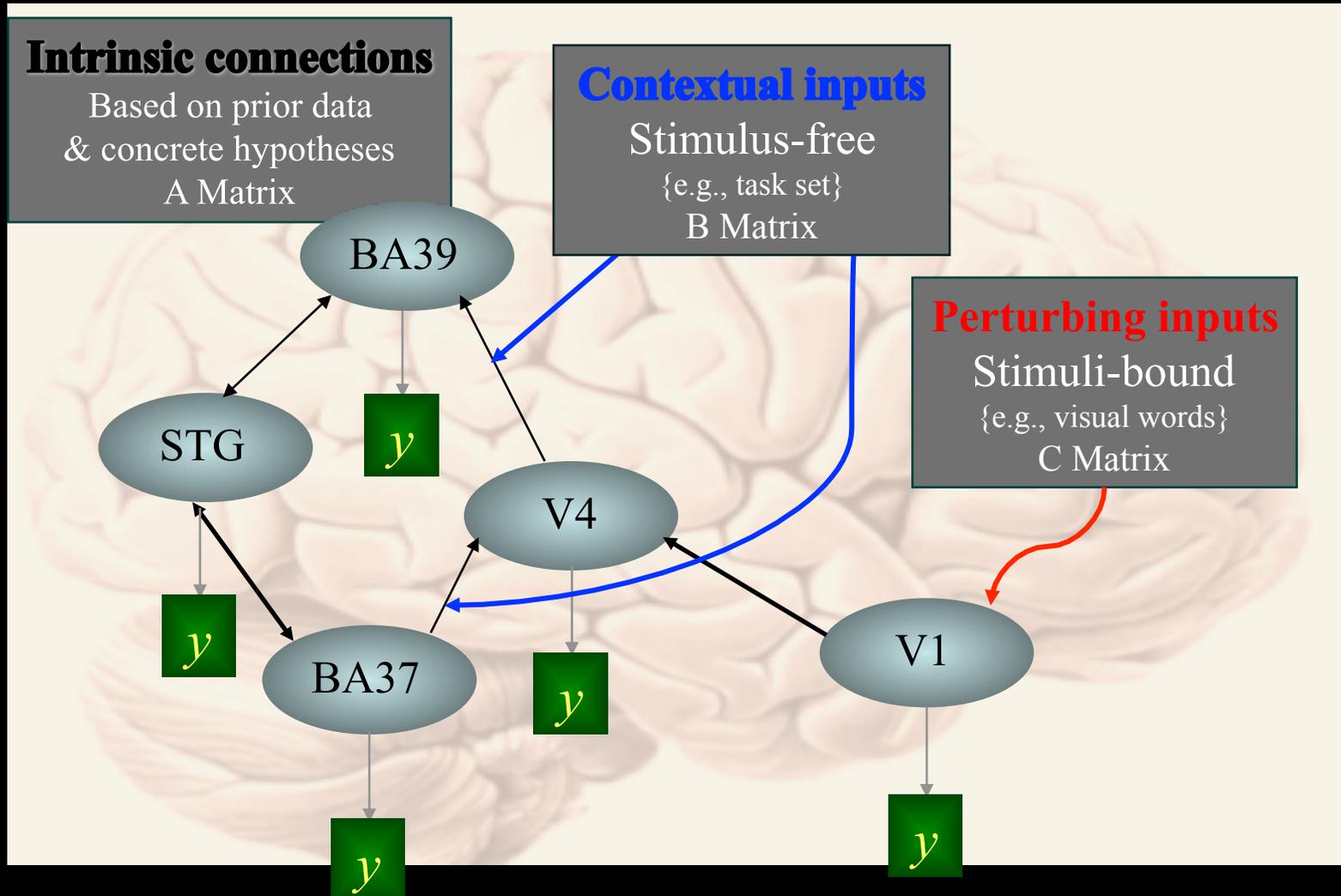
# Dynamic causal modeling (DCM)

- The dynamic state of a given neural circuit is modelled at its underlying neuronal level (not directly accessible with fMRI).
- The modeled neuronal dynamics ( $x$ ) are transformed into area-specific 'simulated' BOLD signals ( $y$ ) by a hemodynamic model ( $\lambda$ ).
- The aim of DCM is to estimate parameters at the neuronal level such that the modelled BOLD signals are maximally similar to the experimentally measured BOLD signals.
  - Model parameters estimated using a Bayesian framework

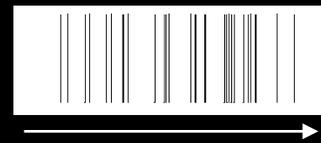


# Dynamic causal modeling (DCM)

## Schematic overview



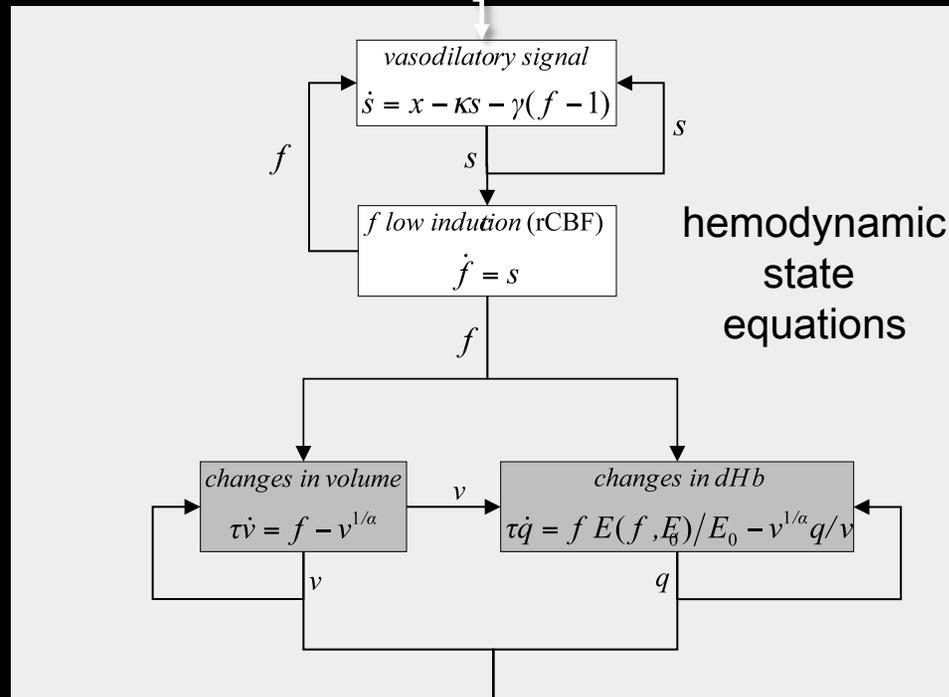
# The hemodynamic model



$u$  stimulus functions

activity  
 $x(t)$

neural state equation



Hemodynamic parameters computed separately for each brain area

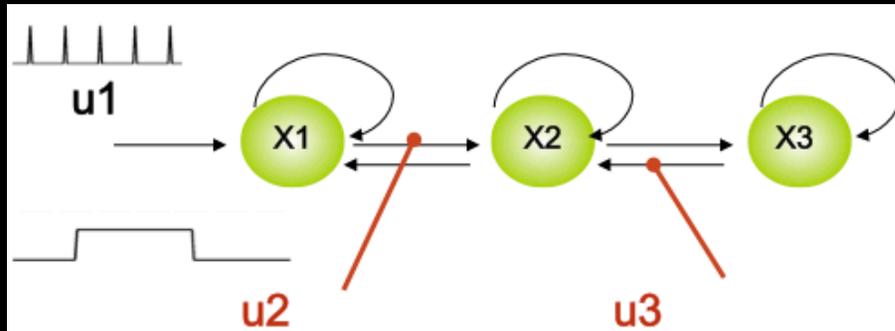
BOLD signal

$$y(t) = \lambda(v, q)$$

Estimated BOLD response

# Dynamic causal modeling (DCM)

## Mathematical overview



### Neural State Equation

$$\dot{x} = \left( A + \sum_{j=1}^m u_j B^{(j)} \right) x + Cu$$

$$\theta = \{A, B, C\}$$

$$\dot{x}_1 = a_{11}x_1 + a_{12}x_2 + c_1u_1$$

$$\dot{x}_2 = (a_{21} + u_2 b_{21}^{(2)})x_1 + a_{22}x_2 + (a_{23} + u_3 b_{23}^{(3)})x_3$$

$$\dot{x}_3 = a_{32}x_2 + a_{33}x_3$$

state changes

fixed effective connectivity

modulatory effective connectivity

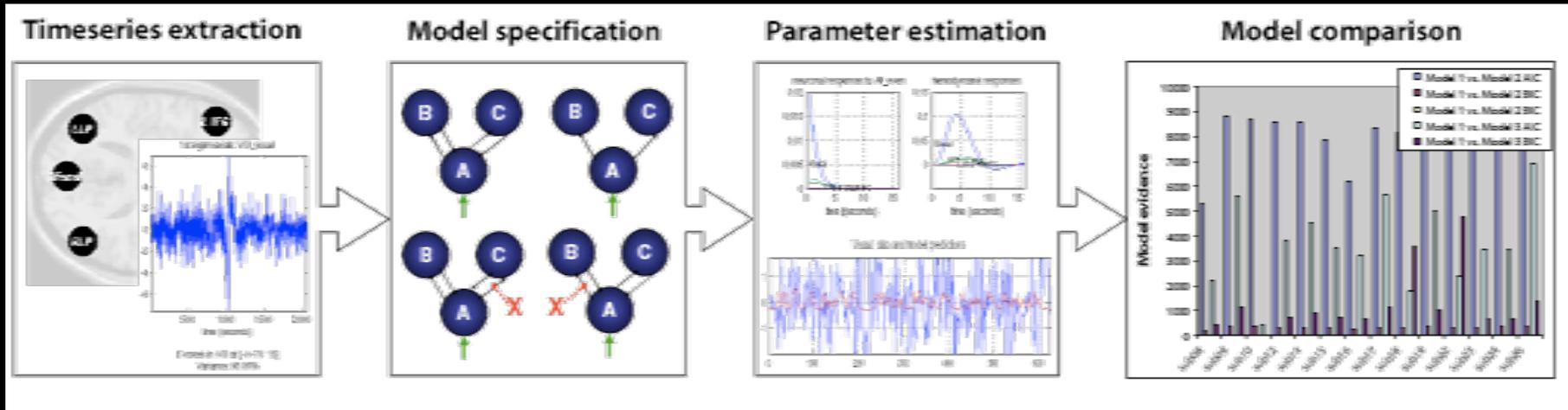
system state

input parameters

external inputs

$$\begin{bmatrix} \dot{x}_1 \\ \dot{x}_2 \\ \dot{x}_3 \end{bmatrix} = \begin{bmatrix} a_{11} & a_{12} & 0 \\ a_{21} & a_{22} & a_{23} \\ 0 & a_{31} & a_{33} \end{bmatrix} + u_2 \begin{bmatrix} 0 & 0 & 0 \\ b_{21}^{(2)} & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix} + u_3 \begin{bmatrix} 0 & 0 & 0 \\ 0 & 0 & b_{23}^{(3)} \\ 0 & 0 & 0 \end{bmatrix} \begin{bmatrix} x_1 \\ x_2 \\ x_3 \end{bmatrix} + \begin{bmatrix} c_{11} & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{bmatrix} \begin{bmatrix} u_1 \\ u_2 \\ u_3 \end{bmatrix}$$

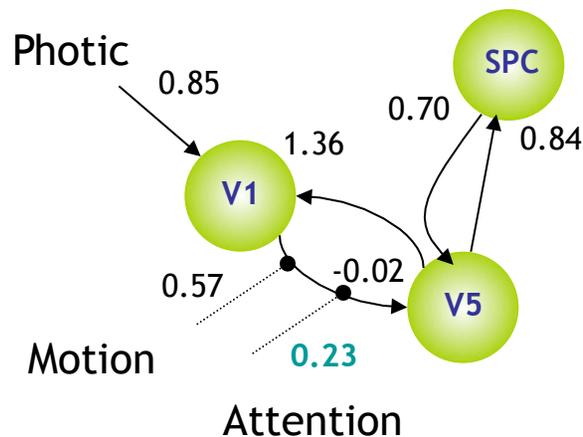
# The DCM processing stream



# Model comparison

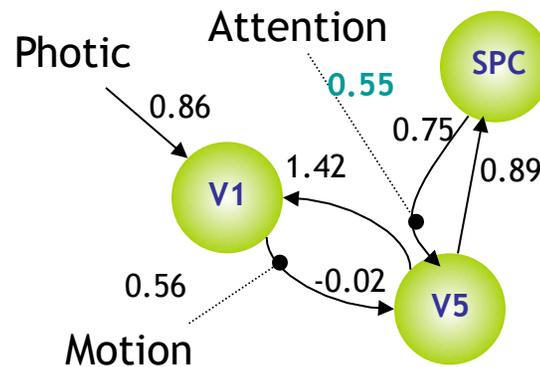
## Model 1:

attentional modulation  
of  $V1 \rightarrow V5$



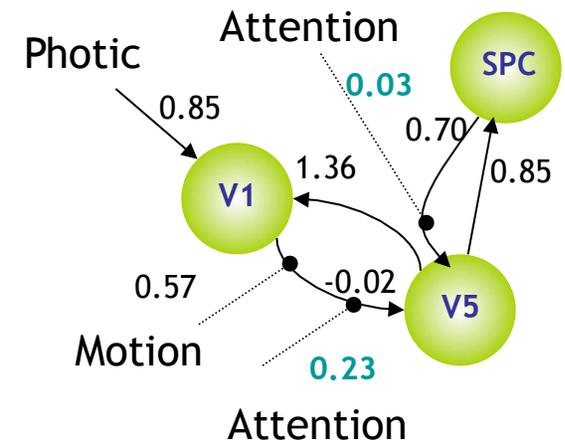
## Model 2:

attentional modulation  
of  $SPC \rightarrow V5$



## Model 3:

attentional modulation  
of  $V1 \rightarrow V5$  and  $SPC \rightarrow V5$



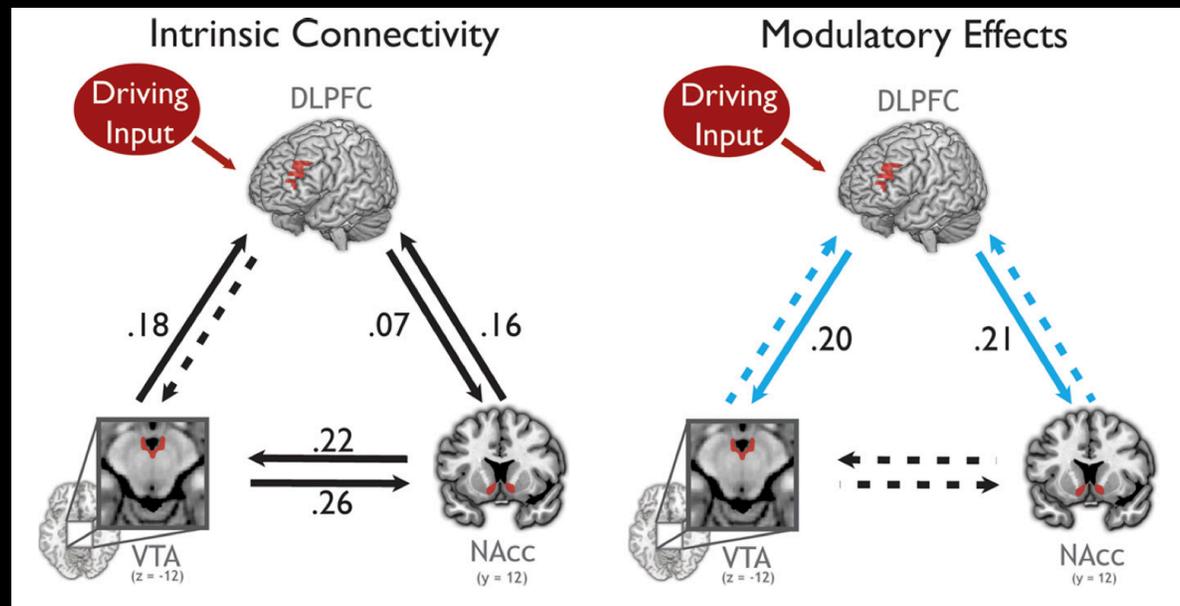
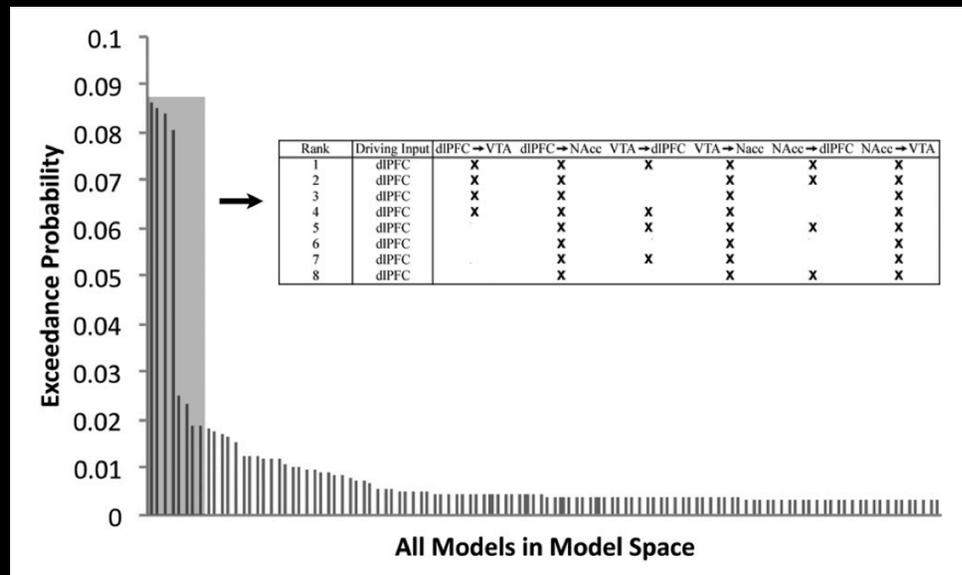
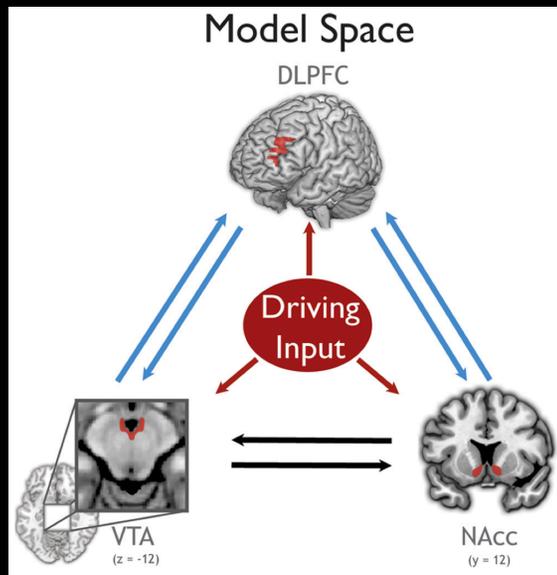
Bayesian model selection:

Model 1 better than model 2;  
model 1 and model 3 equal

Decide on Model 1 (since it is the simpler model):

*Attention primarily modulates  $V1 \rightarrow V5$  pathway*

# A recent DCM analysis example



Ballard (2011), *J Neuroscience*

# Pros and cons of DCM



## Pros:

- Like SEM, DCM examines the functional interplay of many brain areas at once
- Parameters estimated at the (inferred) neuronal level
- Stimulus-based and context-based perturbations of connectivity are explicitly coded into the model
  - Unlike SEM, where inputs are treated as unknowns and the path coefficients are separately computed for each condition
- Operates in the time series domain (dynamic)

## Cons:

- Complex methodology with many built-in assumptions
- Requires a relatively small number of nodes to reduce model complexity

**Thanks!**

