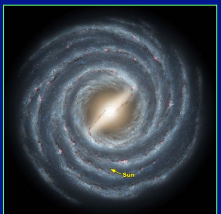
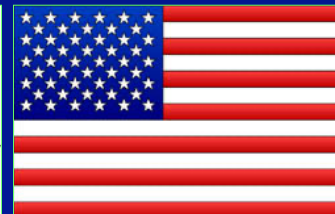


# Functional MRI Analysis with AFNI accents

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# To Dream the Impossible Dream

- *i.e.*, To explain fMRI Analysis in 60 min
- Basis for fMRI time series analysis:
  - 10-20 inter-twined concepts
  - pre-processing ; individual subject analysis ; group analysis ; connectivity ; ...
- Many seem simple, but *each* one has subtleties and points of contention
- ☀ Can only cover the most central ideas now
  - ◆ **Absolute Beginners**: Confusion Ahead!

# Some Goals of fMRI Analyses

- **Task-based experiments**
  - Per subject: estimate amplitude of BOLD response to each different type of stimulus
  - Model + Estimate inter-regional correlations between fluctuations in BOLD responses

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- **Resting-state experiments**
  - Measure spatial patterns in coherent fluctuations in spontaneous BOLD effects

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- **Group level**
  - Combine and contrast per subject results

# Conceptual Basis - 1

- **Time shifting** = pretend get 3D snapshot
- **Despiking** = remove large blips
- **Image Registration** (AKA alignment)
  - intra-EPI time series, and EPI-Structural
- **Blurring in space** = lower resolution :-(& less noise :-) & more group overlap :-)
- **Masking** = ignore non-brain voxels
- **Scaling** = normalizing data amplitude
  - Makes inter-subject comparisons more valid

# Conceptual Basis - 2

## ★ Time series regression

- model of the BOLD response in the data = Hemodynamic Response Function **convolved** with stimulus timing
- plus baseline (null hypothesis) model
- plus physiological noise
- plus allow for serial correlation
- plus allow for random response strength (!!)

## ★ Talairach-ing = Spatial Normalization

- Talairach, MNI-152, ...
- affine and nonlinear spatial transformations

# Conceptual Basis - 3

★ **Group Analyses** = Putting it all together

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★ – ANOVA, LME, MVM, Meta-Analyses, ...

- **Blobs** = Spatial models of activation

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★ – Assigning statistical significance to blobs

- **Connectivity** = Inter-regional analyses

- SEM, PPI, SVAR, DCM, ...

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- Resting state fMRI (Connectome!)

- **Dimensional factorization**

Components, such as PCA, ICA

# Conceptual Basis - 4

- **Data Formats** = NIfTI-1.x is your friend
- **Software** for fMRI analyses: \*open-source
  - AFNI<sup>\*</sup>, **BrainVoyager**, FSL<sup>\*</sup>, SPM<sup>\*</sup>, ...
  - Whichever you use, **don't blindly assume** the software works perfectly all the time
- Most important thing I will say today  
**Understand and check the steps applied to your data!**
- 2<sup>nd</sup> most important: **Is no "best" way to analyze data, just "reasonable" ways**

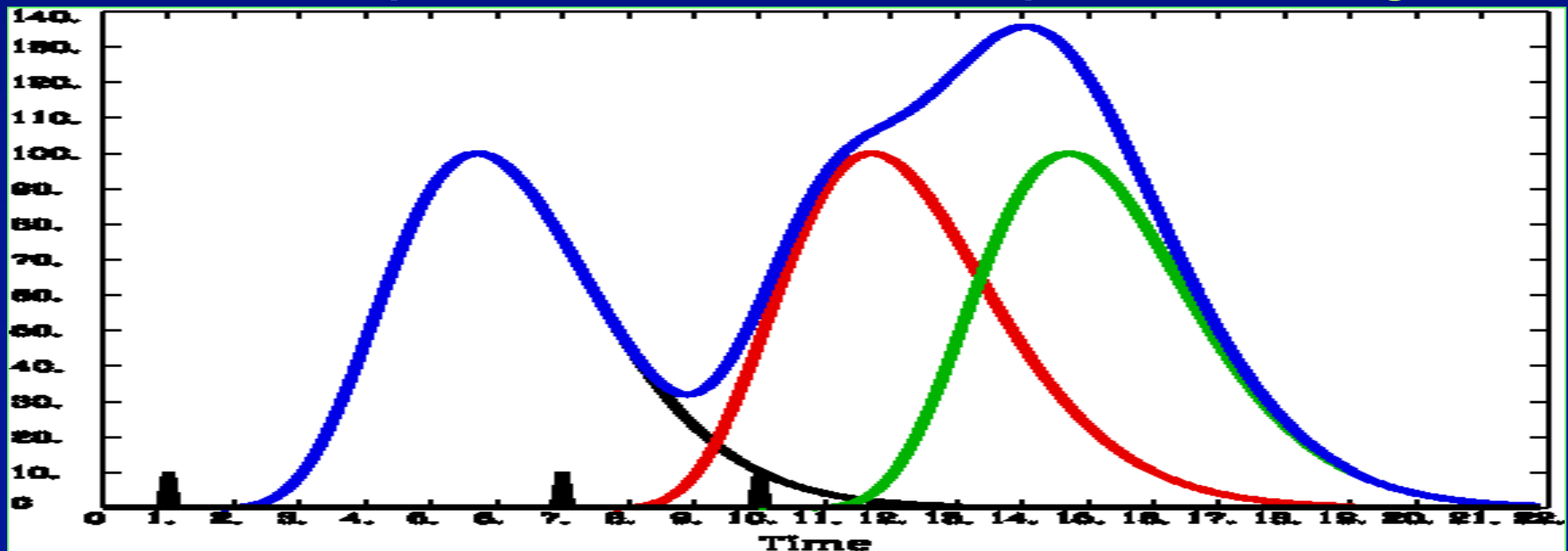
# ★ Time Series Regression

- After pre-processing the images:
  - Each voxel time series is (separately) fit to an idealized model of the BOLD effect
  - Plus other (nuisance) components
    - Baseline drift, physiological artifacts, ...
- Estimated parameters of the BOLD model are the "**activation map**"
  - Amplitude of response = most common
  - Shape of response can also be modeled
  - Statistics on parameters allow thresholding



# Basic BOLD Model

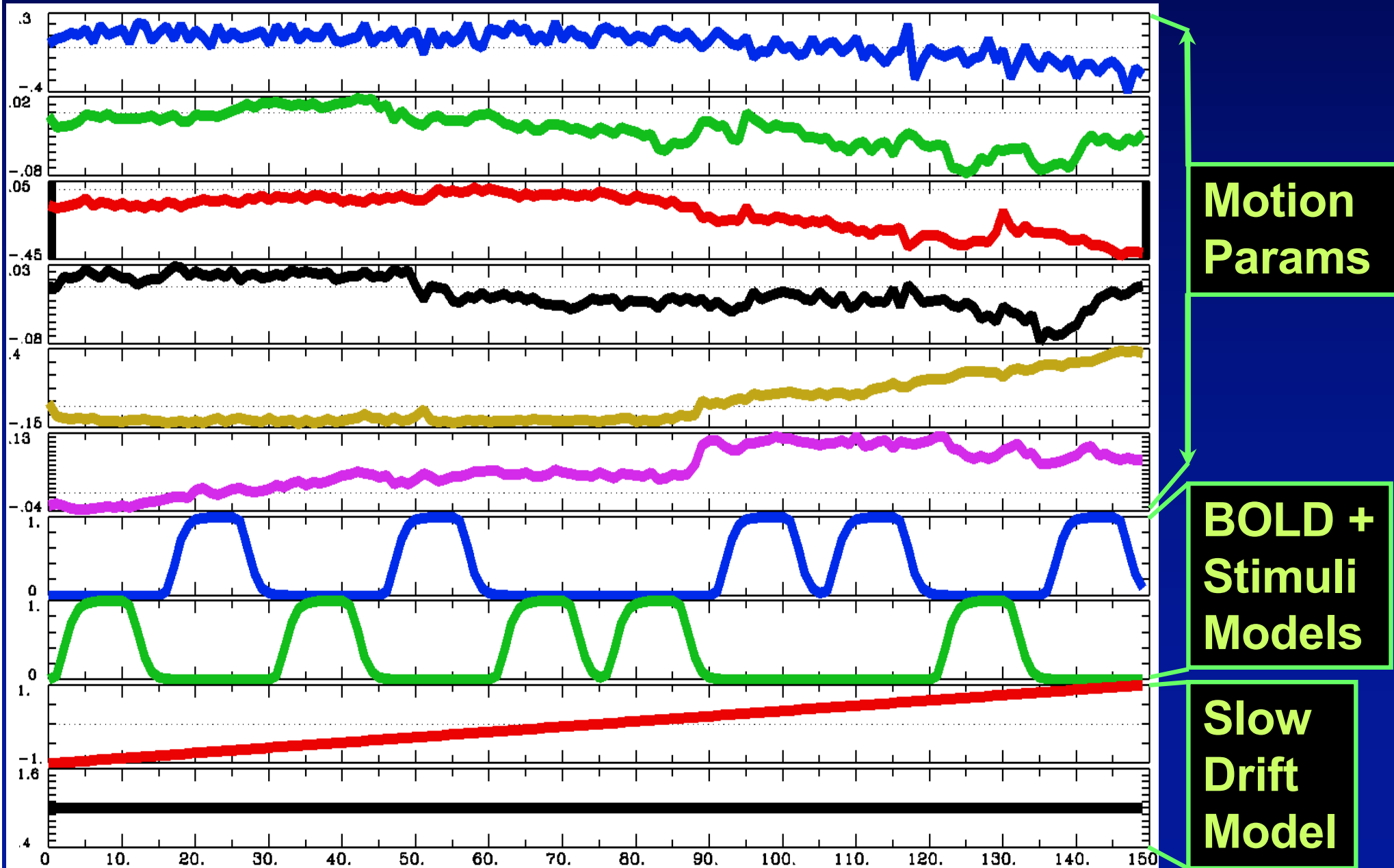
- Assumptions:
  - BOLD response starts  $\approx 2$ s after "activity"
  - Rises to peak about  $\approx 4$ s later
  - Drops to baseline about  $\approx 5$ s post-activity
  - Overlapping (in time) BOLD responses from separate events add up = **linearity**



# Further Model Pieces-Parts

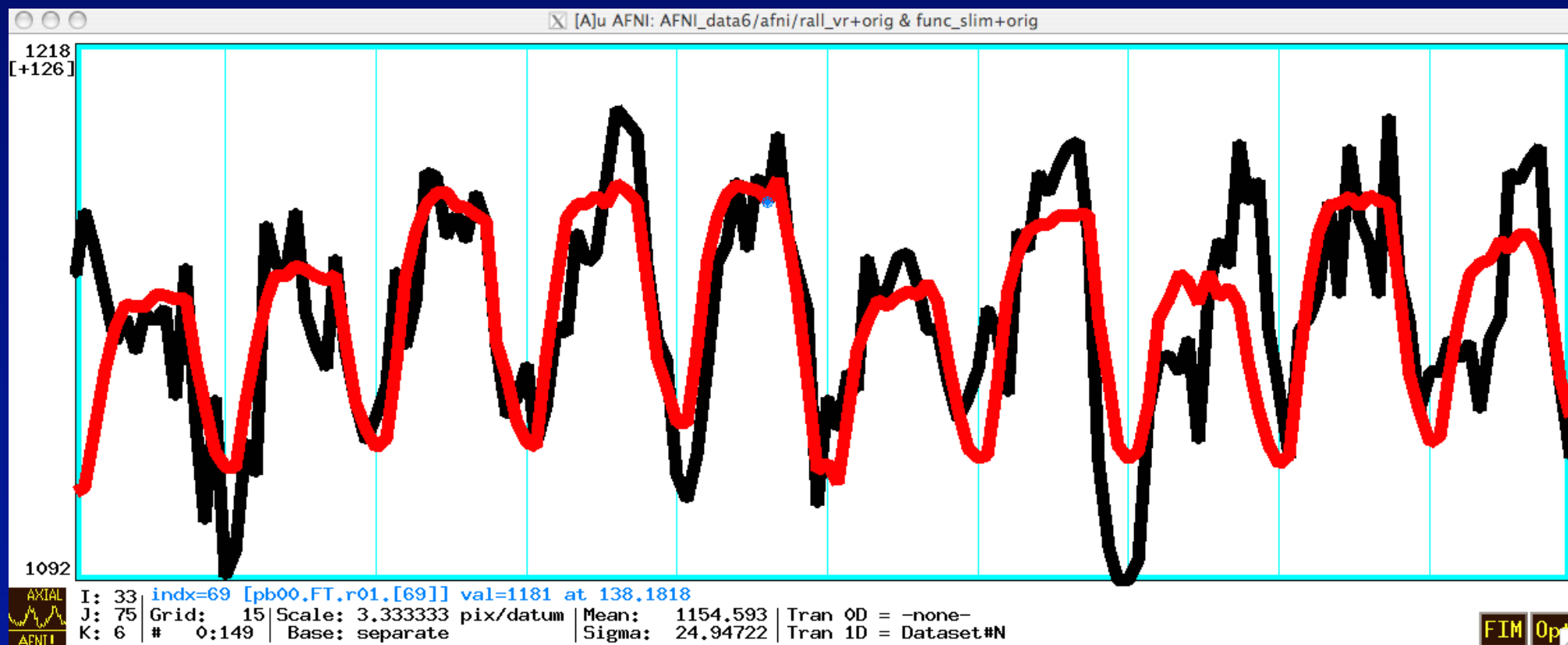
- FMRI data = pretty noisy →
  - Need multiple copies of each stimulus type
  - Assume BOLD responses same within type
- Slow drifting terms in baseline highpass filter
- Residual effects of subject movements
  - Use motion parameters as extra regressors
- Include physiological regressors (**RetrolCOR**)
  - From monitoring heartbeat & respiration
- Serial correlation in noise → use **GLSQ**

# Sample Linear Regression Model



# One Voxel's Time Series

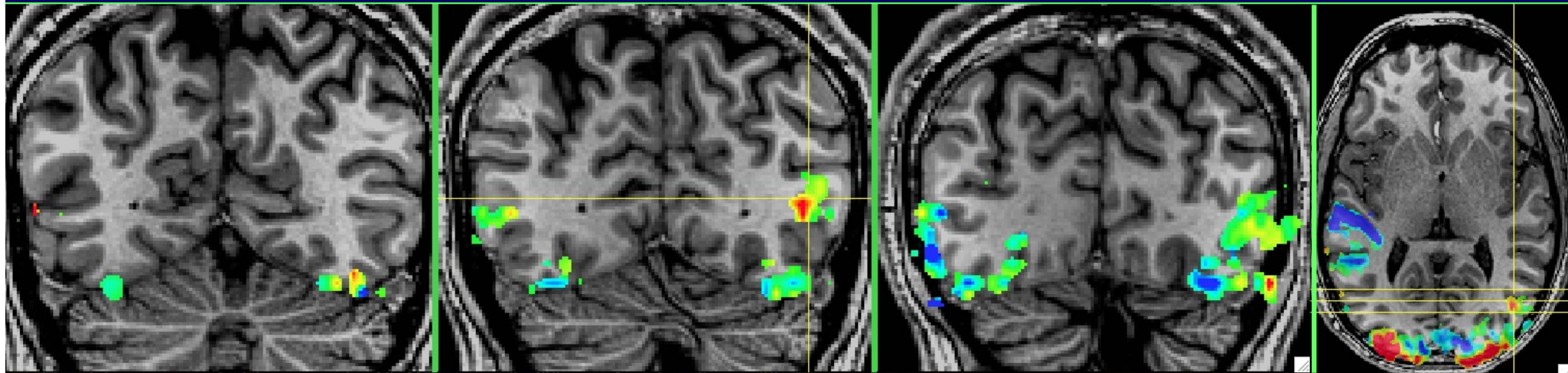
- This is good data at a high signal voxel
- 2 stimulus classes; 1 block every 30s; TR=2s



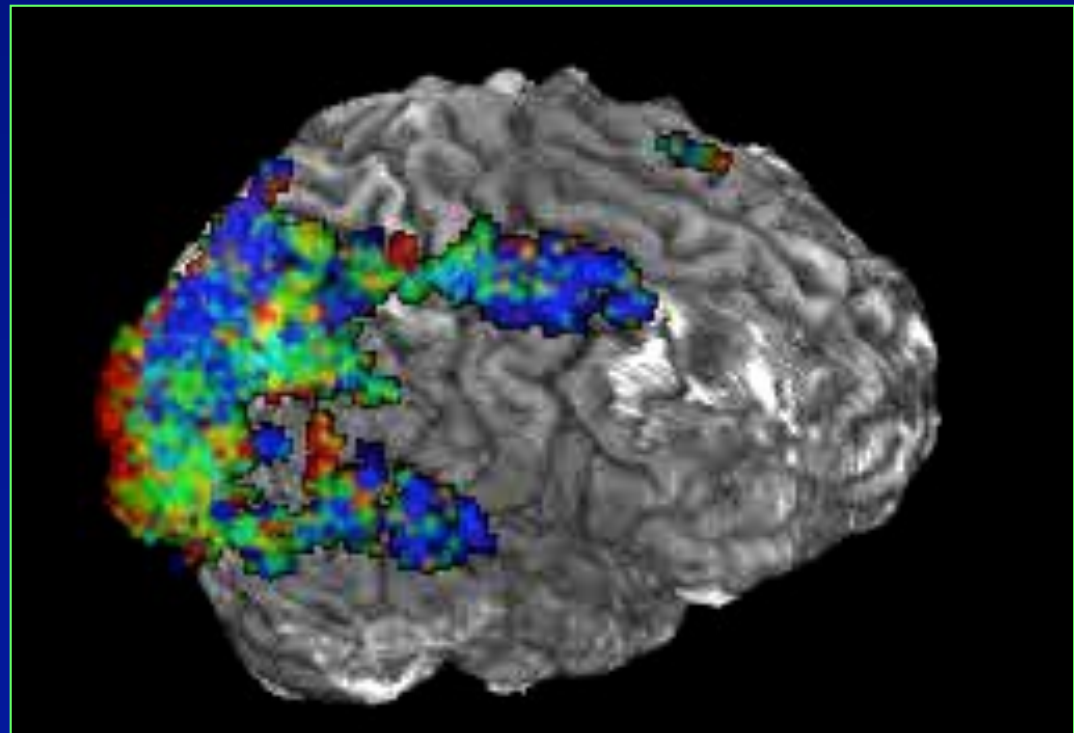
**Black** = 150 data points    **Red** = fitted model (BOLD++)

Data: Mike Beauchamp/UTexas

# Results from One Subject



ShowThru  
rendering of  
difference  
between visual  
and auditory  
BOLD  
amplitudes



# ★ Talairach-ing (AKA Normalizing)

- For combining and contrasting results across groups of subjects

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- Align 3D brain images (usually structural volumes) to a common template
  - Assumption: function follows structure

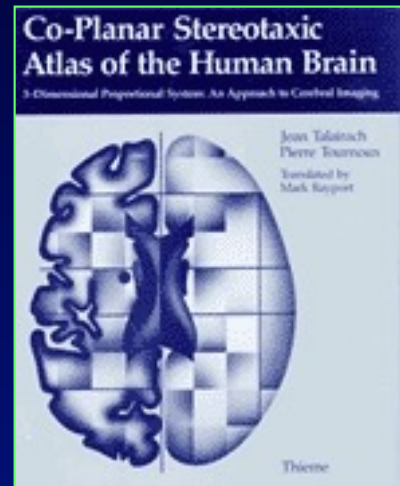
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- Then merge results on a voxel-wise basis on this new grid
  - Accuracy of alignment = 5-10 mm
  - Spatial blurring = more inter-subject overlap

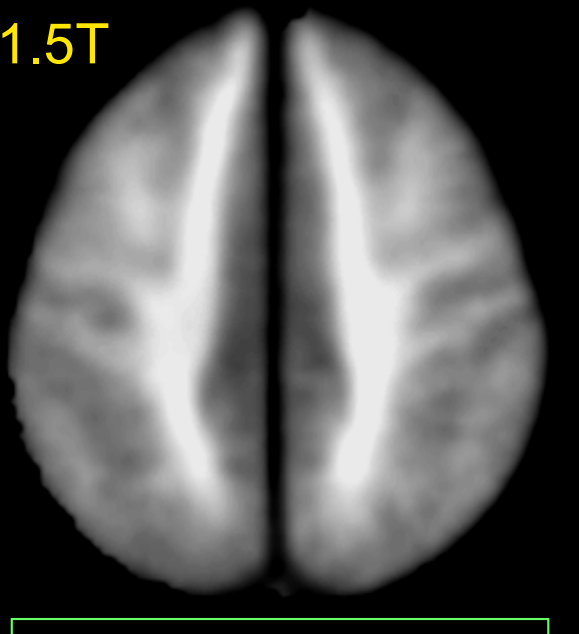


# Templates

- Ye Good Olde Days = Talairach-Tournoux atlas
- Decadent modern times = MNI-152 template = average of 152 brain images

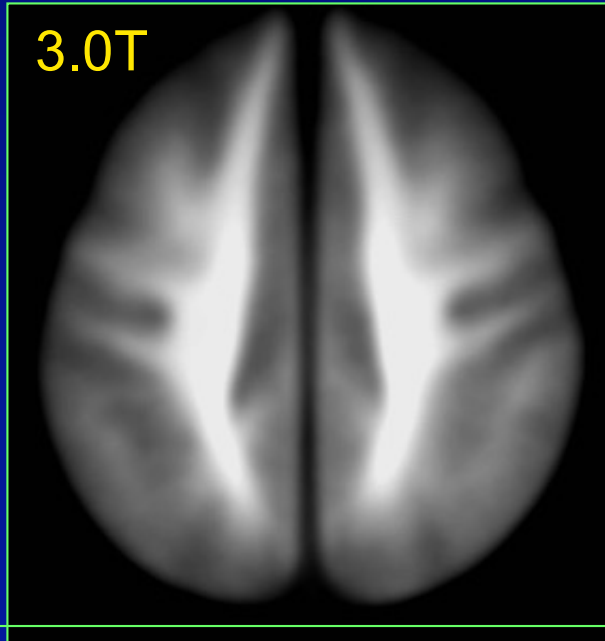


1.5T



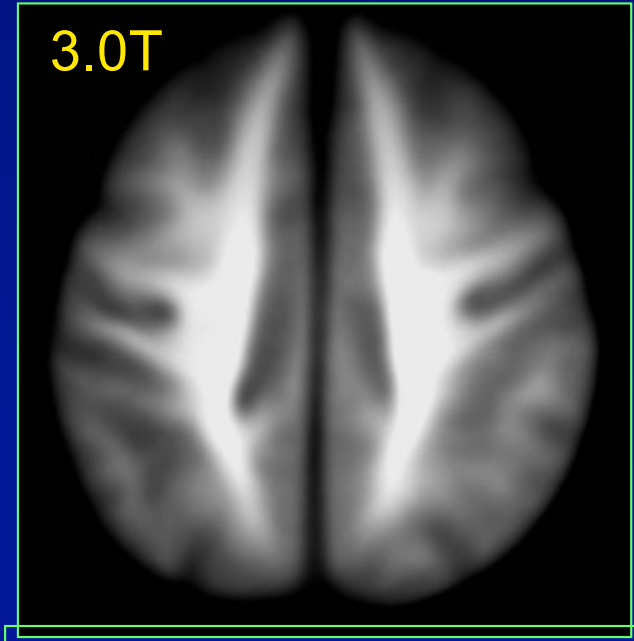
MNI-152 (affine)

3.0T



FCON-1000 (affine)

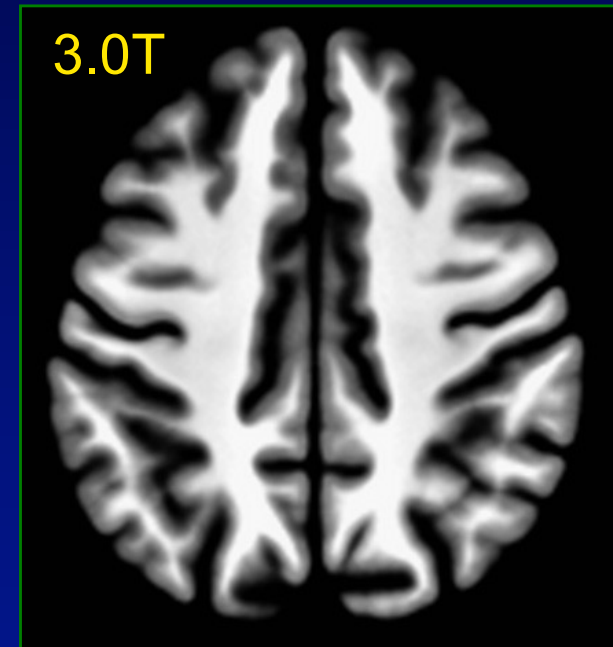
3.0T



FCON-1000 (poly9)

# Custom Templates

- Can register all subjects in study together iteratively
  - Then average to get a study-specific template
  - Can use high order nonlinear spatial warps for better structural feature alignment



101 subjects  
**AFNI 3dQwarp**

Data: MindBoggle

- Alternatives:
  - ROI or Atlas-based analyses
  - Cortical surface-based analyses [cf. **SUMA**]



# ★ Group Analysis

- Group analysis is done in 2 stages:
  - Individual subject analyses reduce EPI time series to BOLD response parameters
  - Inter-subject analysis is done on these parameters (e.g., measures of amplitude)
  - Usually have some estimate of reliability of these parameters (e.g., standard deviation)
- Normalization of data magnitude and spatial normalization are important steps

# Flavors of Group Analysis - 1

- $t$ -tests for simple 2-way contrasts
- **ANOVA** of various complexities for more intricate collective effects
  - Usually end up wanting to do  $t$ -tests on these "main" effects to see **What's Up** (*Doc*)
  - Plain **ANOVA** relies on many assumptions about distribution of randomness
    - Independent noise (or all samples are correlated the same way; "sphericity") ; Gaussianity ; **Homoscedasticity** (all samples have same variance) ; Balanced designs

# Flavors of Group Analysis - 2

- Linear Mixed Effects (**LME**) and Multi-Variate Models (**MVM**) generalize **ANOVA** to avoid such assumptions

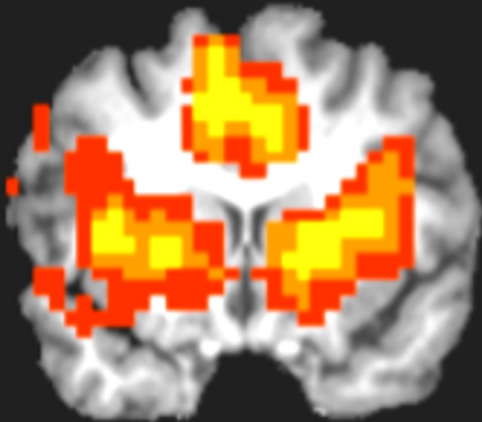
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- Meta-Analysis: Statistics run on statistics
  - Which is exactly what we are doing in FMRI
  - Many developments in the 1990s
  - Can be applied to FMRI data, using not just the individual subject amplitudes, but also their reliability estimates; cf. **AFNI** and **FSL**
  - Can also model non-Gaussianity (outliers)

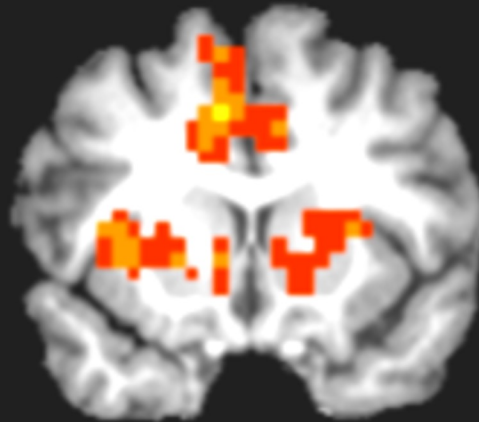
# Meta-Analysis Results

## Meta-Analysis

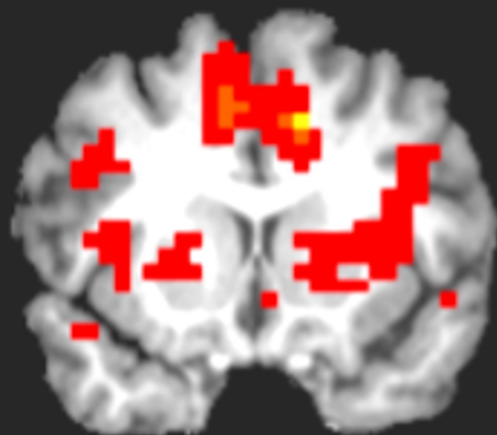
Task A: 20 sub



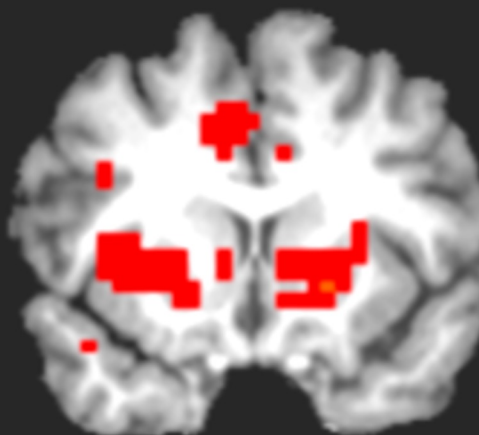
Task B: 12 sub



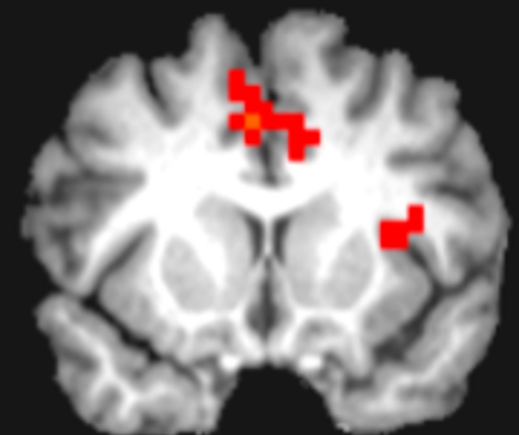
Task C: 11 sub



ANOVA



Yellow = Stronger




Data: James Bjork/NIH

## ★ Blobs! [AKA Clusters]

- For most purposes, if we saw that all the *statistically* significant activation was just in tiny and scattered clusters, we would be suspicious and upset
- We not only threshold on voxel-wise  $t$ - or  $F$ -statistics, but we also threshold on size of contiguous supra-threshold clusters
- Allows us to lower the per-voxel threshold by raising the cluster-size threshold to maintain error rate control

# Blob Significance (!!)

- Probability that a smooth *noise-only* image would produce a blob, with each voxel above a given threshold and the size of the blob above a given cutoff
- Cannot be calculated in closed form :-(

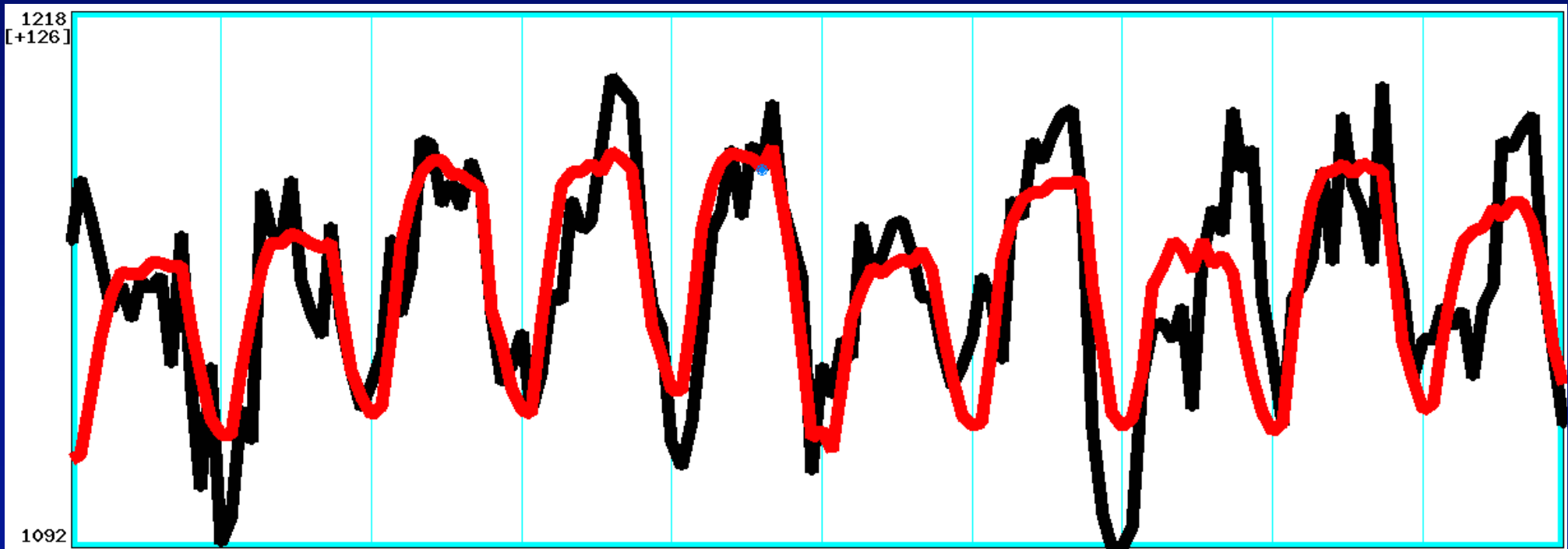
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- Two approaches available:
  - Asymptotic formula for high per-voxel thresholds and high degrees of smoothness K Worsley
  - Direct Monte-Carlo simulation (brute force) 
  - Which to use? Depends on your software! AFNI

# ★ Connectivity

- **FMRI basis:** Fluctuations in neural activity → fluctuations in BOLD signal
- → Task performance changes between repetitions can give coherent signal changes in remote brain regions
- Can look for *causal* changes also:
  - e.g., Predictability of one region's data from another's data at a *previous* time step
  - Limited TR of FMRI data makes this iffy
    - Probably need TR < 1s to do this well

# Task-Based Connectivity

- Fluctuations from model fit can be bigger than noise; recall this graph from 1 voxel

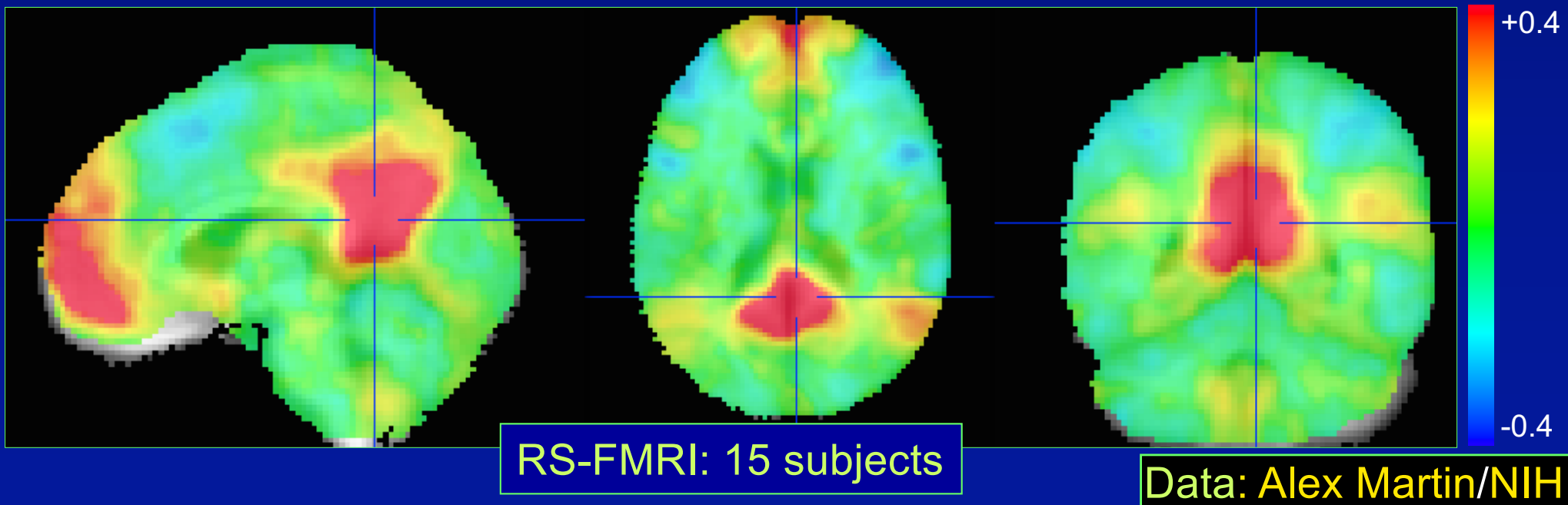


- So can subtract the fit and then look for inter-voxel correlations (if have enough data)



# Flavors of Connectivity - 1

- Simple correlation from a seed region (or voxel) with all other voxels in brain
  - Ambitiously: Correlate All-with-All :-)
- Then group analysis for statistical power
- Used in resting-state FMRI analyses



# Flavors of Connectivity - 2

- PPI = **P**sycho-**P**hysiological **I**nteraction
  - We call it "Context Dependent Correlation"
  - Augment the regression model with data time series averaged from a seed region
    - And multiplied by **+1** in context **A**, **-1** in context **B**, and **0** in other cases (e.g., baseline)
  - Fit parameter for new regressor measures fluctuations in EPI signal whose correlation with seed region **changes** between contexts (**A & B**)
  - Context = "Psycho" ; EPI/BOLD = "Physio"

# Flavors of Connectivity - 3

- Network **models** use multiple regions, try to fit multiple dependencies amongst them
  - FMRI noise means must average over pre-chosen regions before network hunting+fitting
  - Anatomical ROIs or Functional ROIs?

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- Instantaneous directional effects:
  - Structural Equation Modeling (**SEM**)

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- Delayed (causal) effects:
  - Dynamic Causal Modeling (**DCM**)
  - Structural Vector AutoRegression (**SVAR**)

# Connectivity Conclusions

- Often: make statement about *changes* in connectivity between two situations
  - Absolute connectivity statements are somewhat less common (though not rare)
- When contrasting subject groups, must be careful that physiological differences (e.g., respiration) are accounted for
  - Especially in resting-state studies, which don't have any task-timing to which we can tie down the analysis

# Final Gratuitous Remarks

- Multi-stimulus linear regression models for task-based fMRI appear in mid-1990s

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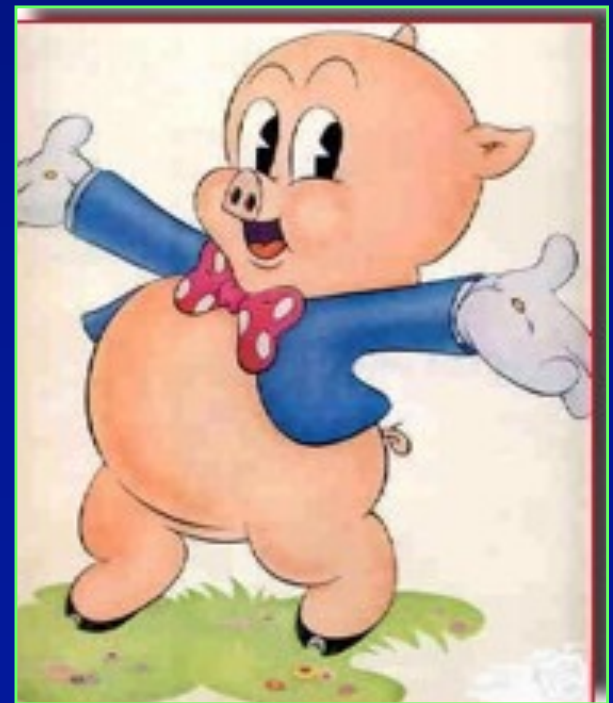
- Since then, most task-based fMRI time series analysis improvements have been incremental **and/or** controversial **and/or** of limited scope — **IMHO**

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- Are we near the limit of what kind of information can be gleaned from current-technology fMRI datasets?
  - **Or do we just need to be cleverer?**

# Th'Th'That's All Folks!

- Thanks must go to
  - Ziad Saad, Gang Chen, Rick Reynolds, Daniel Glen, Paul Taylor
  - James Hyde, Peter Bandettini, Andrzej Jesmanowicz
  - Elliot Stein, Jeff Binder, Steve Rao, Alex Martin, Leslie Ungerleider, Ted Deyoe, ... and too many more brain-ologists to name herein



Data: Warner Bros.