

# FMRI Data Analysis at Individual Level

SSCC/NIMH/NIH/HHS

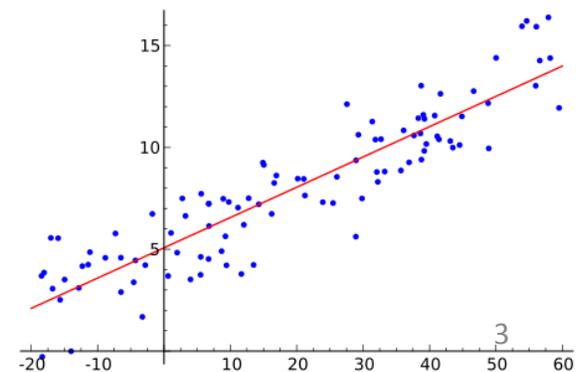


# Overview

- Basics of linear model
- FMRI experiment types
  - Block design; Event related experiment; Mixed
- FMRI data decomposition: three components
  - Baseline + slow drift + effects of no interest; Effects of interest; Unknown
  - Effects of interest - understanding BOLD vs. stimulus: IRF
- Three modeling strategies
  - Fixed-shape IRF
  - No assumption about IRF shape
  - One major IRF plus shape adjustment
- Other issues
  - Multicollinearity
  - Catenation
  - Percent signal change

# Basics of Linear Model

- **Regression**: relationship between a response / outcome (dependent) variable and one or more explanatory (independent) variables (regressors)
  - Simple regression: fit data with a straight line: Sir Francis Galton's original meaning: regression to mean
    - When 2 variables are not perfectly correlated, regression to mean exists
    - May show up in psychology (Daniel Kahneman): Rewards for good performance vs. punishment of mistakes (correlation vs. causation)
    - Lost in most cases including FMRI
  - Some statisticians just call it **linear model**
- Mathematical crystallization
  - $y_i = \alpha + \beta x_i + \varepsilon_i$ , or  $y_i = \alpha + \beta_1 x_{1i} + \dots + \beta_k x_{ki} + \varepsilon_i$
  - $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$ ,  $\mathbf{X} = [1, x_1, x_2, \dots, x_k]$
  - Assumption
    - linearity
    - white noise (independence) and Gaussianity  $\boldsymbol{\varepsilon} \sim N(0, \sigma^2 \mathbf{I})$



# Basics of Linear Model

- Solution for linear regression  $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$ 
  - Project data  $\mathbf{y}$  onto the space of explanatory variables ( $\mathbf{X}$ )
  - OLS  $\hat{\boldsymbol{\beta}} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$
- Meaning of coefficient:  $\beta$  value, slope, marginal effect or effect size associated with a regressor
- Various statistical tests
  - Student  $t$ -test for each  $\beta$  ( $H_0: \beta_3 = 0$ )
  - Student  $t$ -test for linear combination of  $\beta$  values - general linear test (GLT), *e.g.*,  $H_0: \beta_3 - \beta_5 = 0$ , or  $H_0: 0.5*(\beta_3 + \beta_4) - \beta_5 = 0$
  - $F$ -test for composite null hypothesis, *e.g.*,  $H_0: \beta_3 = \beta_4 = \beta_5$  or  $H_0: \beta_3 = \beta_4 = \beta_5 = 0$
  - Omnibus or overall  $F$ -test for the **whole** model, *e.g.*,  $H_0$ : all  $\beta$  values are 0, or  $H_0$ : all  $\beta$  values of interest are 0

# Linear Model with fMRI

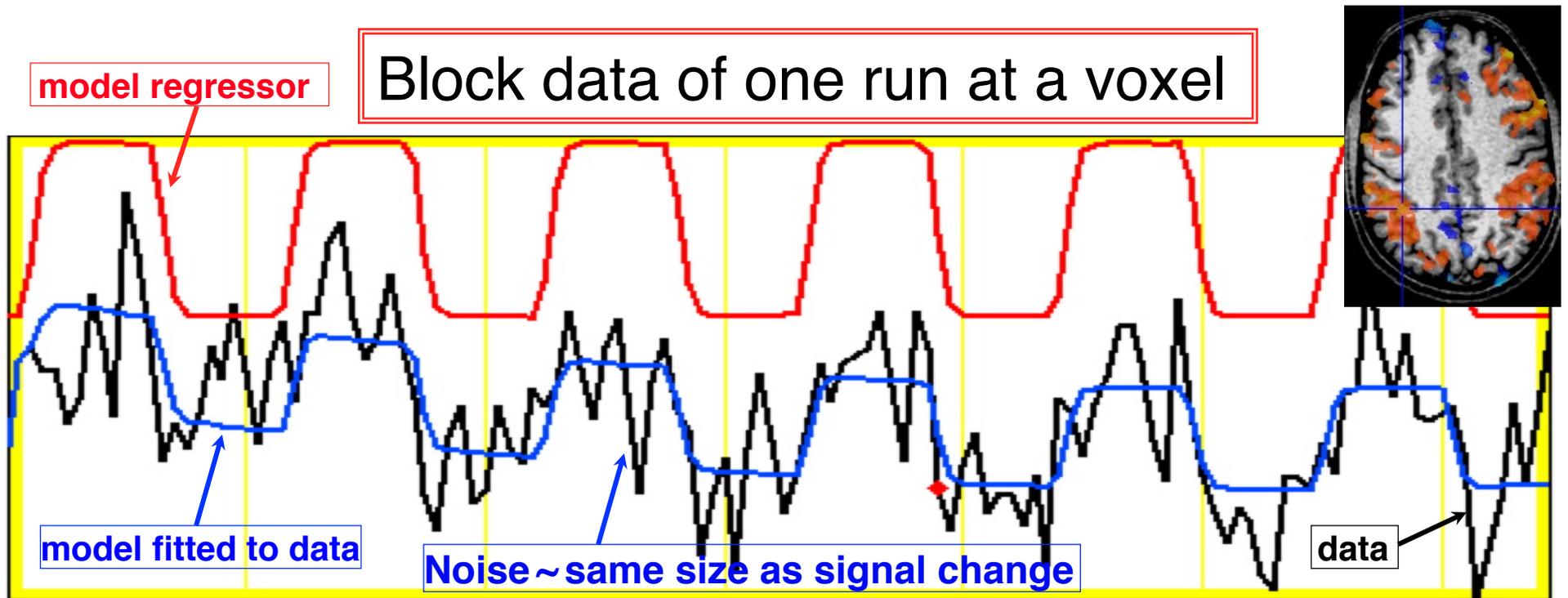
- Time series regression: data  $\mathbf{y}$  is time series
  - Regressors: idealized response or yardstick
    - We get what we're looking for
    - It may miss something when we fail to recognize it
    - Regressor construction is quite challenging
  - Special handling: noise not white  $\boldsymbol{\varepsilon} \sim N(0, \sigma^2 \boldsymbol{\Sigma})$  with temporal or serial correlation
    - Banded variance-covariance matrix  $\boldsymbol{\Sigma}$
  - AKA general linear model (GLM) in other fMRI packages
    - General vs. generalized
- Same model for all voxels in the brain
  - Simultaneously solve the models: voxel-wise analysis, massively univariate method

# FMRI Experiment Terminology

- Experiment setup
  - Number of subjects
  - Number of conditions (tasks, stimulus (trial, event) types): Factorial design?
  - Sample size (repetitions) per condition
  - Block, event-related, or mixed?
  - Inter-stimulus interval (ISI)
- Scanning parameters: TR, voxel size, data points (volumes), slice sequence (sequential or interleaved), slice thickness, removing first few TRs
- Scanning terms
  - Run: continuous scanning; a brief break before next run
  - Session: subjects come back after along period of time
  - Experiment or study

# Types of fMRI Experiments

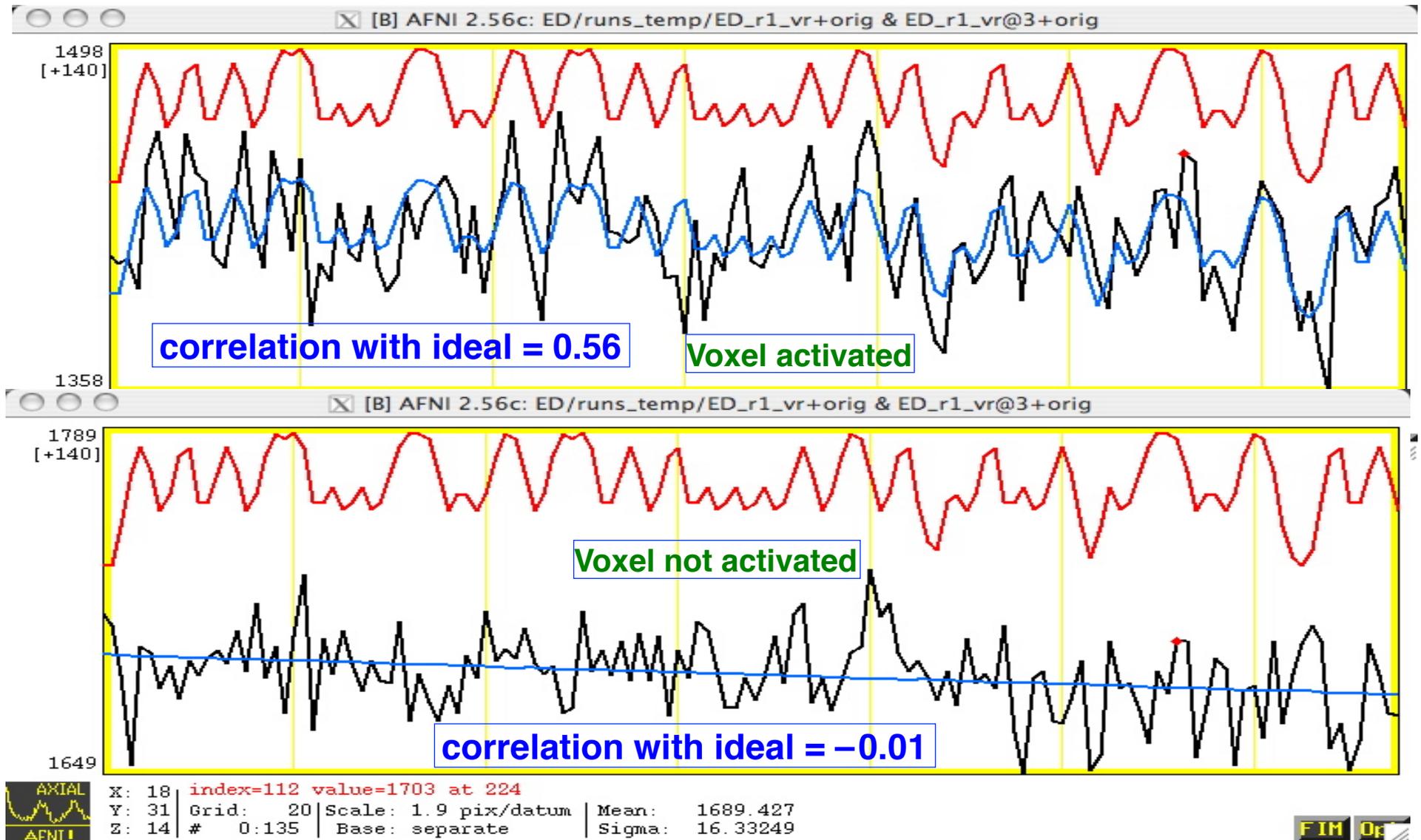
- Two classical types of experiment design
  - Block (boxcar) design
    - Each block lasts for more than one TR (e.g., 4 to 20s)
    - Each block is under one condition (e.g., watch a video clip), or a series of multiple trials (e.g., 10 consecutive blur images)
    - BOLD response is often visible in time series
    - SNR: noise magnitude about same as BOLD response
  - Event-related design
    - Each event or trial lasts for one TR or shorter
    - Events are randomly spaced and/or sequenced in time
    - BOLD response to stimulus tends to be weaker, since fewer nearby-in-time “activations” have overlapping signal changes
    - SNR: data looks more like noise (to the pitiful human eye)
- Mixed designs
  - Containing both events and blocks, e.g., cue + video watching



Block: 27 s “on” / 27 s “off”; TR=2.5 s; 130 time points

- This is best voxel; most voxels are not fitted as good as this
- Activation amplitude and shape vary across blocks
  - Subject attention variability
  - Habituation (or attenuation): psychological/physiological level
  - Linearity assumption
  - Pure random effects

# Event-Related Data at 2 Voxels



Lesson: ER-FMRI activation is not obvious via casual inspection

# FMRI Data

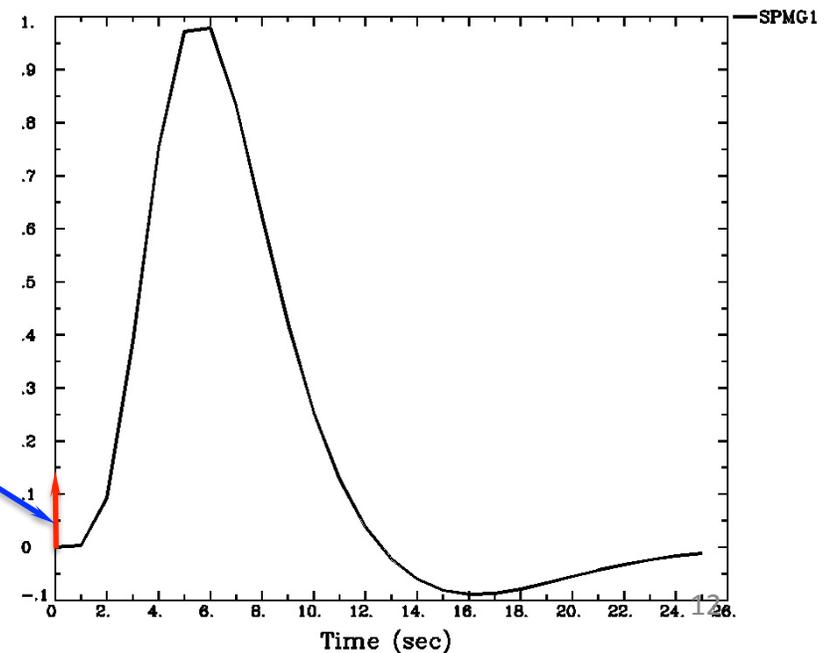
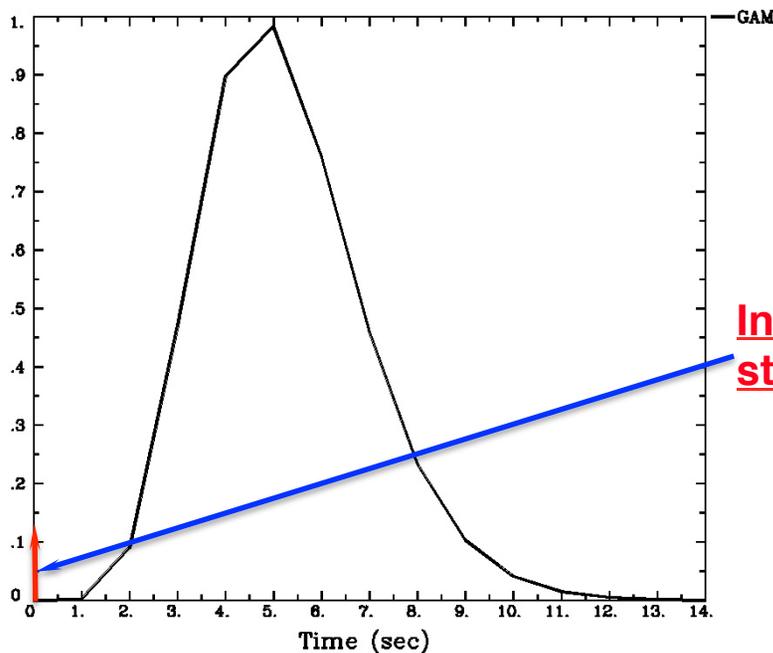
- Data partition: **Data = Signal + Noise**
  - **Data** = acquisition from scanner (voxel-wise time series)
    - What we have
  - **Signal** = BOLD response to stimulus; effects of interest + no interest
    - **We don't really know the real signal!!!**
    - Look for idealized components, or search for signal via repeated trials
    - Of interest: effect size (response amplitude) for each condition: beta
    - Of no interest: baseline, slow drift, head motion effects, ...
  - **Noise** = components in data that interfere with signal
    - Practically the part we have don't know and/or we don't care about; that is, noise is the part we can't explain in the model
    - Will have to make some assumptions about its distribution
- **Data = baseline + slow drift + other effects of no interest + response<sub>1</sub> + ... + response<sub>k</sub> + noise**
  - How to construct the regressors of interest (responses)?

# BOLD Response

- Hemodynamic response (HDR)
  - Brain response to stimulus / task / condition
    - Ideally we want to know the response (activation) at neuronal level, but that is beyond the fMRI capability
  - Indirect measure of neural response: dynamic regulation of blood flow
- Hemodynamic response function (HDF)
  - Mathematical formulation / idealization of HDR
  - BOLD signal is further an indirect measure of brain response
  - HDF bridges between neural response and BOLD signal
- How to build the bridge?
  - One extreme: Assume a fixed-shape (idealized) HDF
  - The other extreme: No assumption about HDR shape
  - Middle ground: major shape + wiggle room for shape adjustment

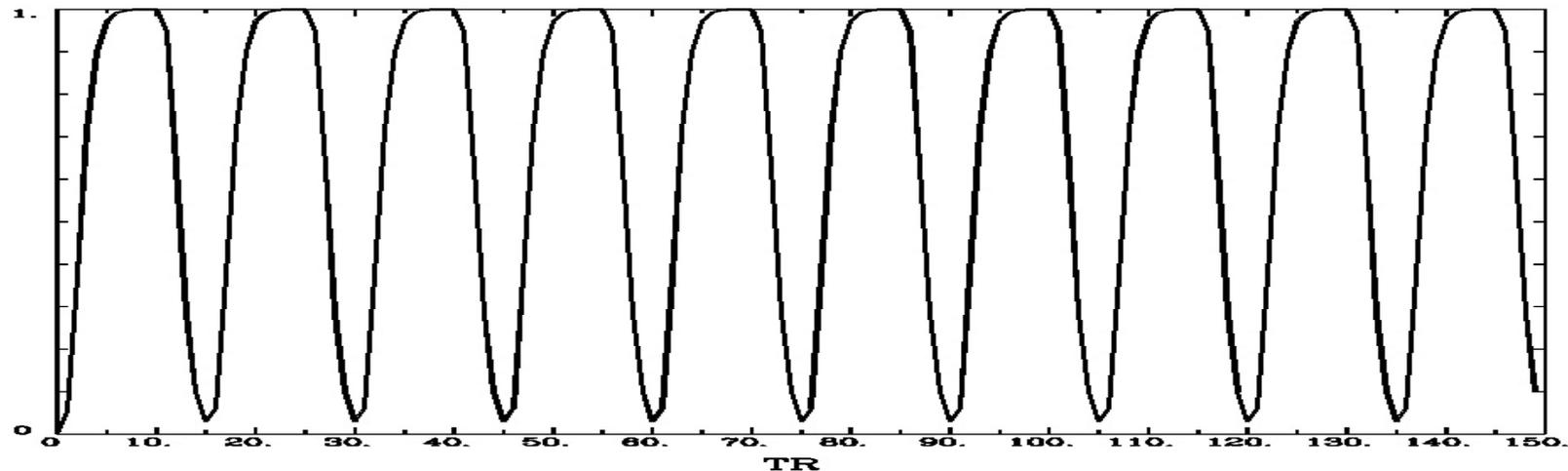
# Fixed-Shape IRF

- Assuming a fixed shape  $h(t)$  for HDF to an **instantaneous** stimulus: impulse response function (IRF)
  - GAM( $p,q$ ):  $h(t) = [t / (p \cdot q)]^p \cdot \exp(-t / q)$ 
    - Default IRF:  $h(t) = t^{8.6} \exp(-t / 0.547)$  [MS Cohen, 1997]
    - A variation: SPMG1 (undershoot)
  - Build HDF based on presumed IRF through **convolution**
    - Roll IRF  $h(t)$  with stimulus timing  $S(t)$ :  $x(t) = h(t) \otimes S(t)$



# Fixed-Shape HDF for Block Design

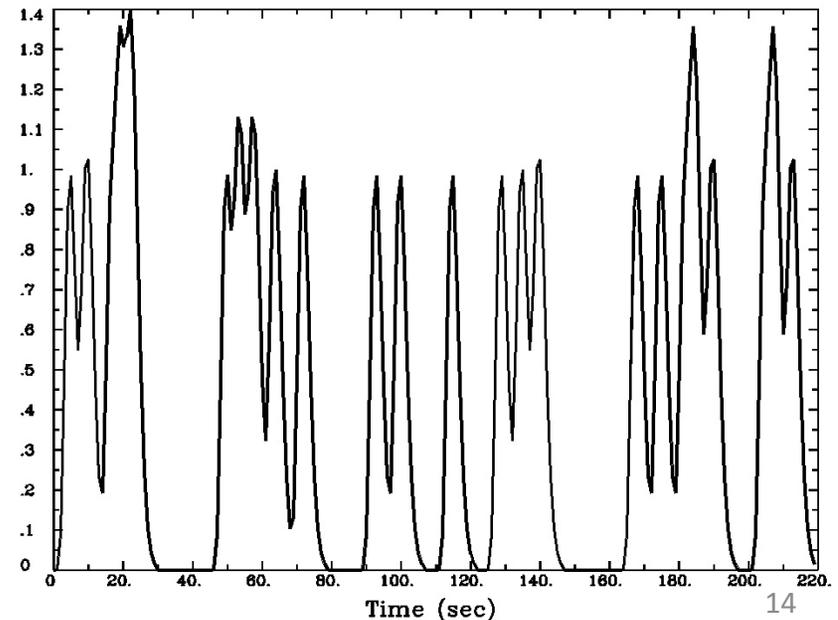
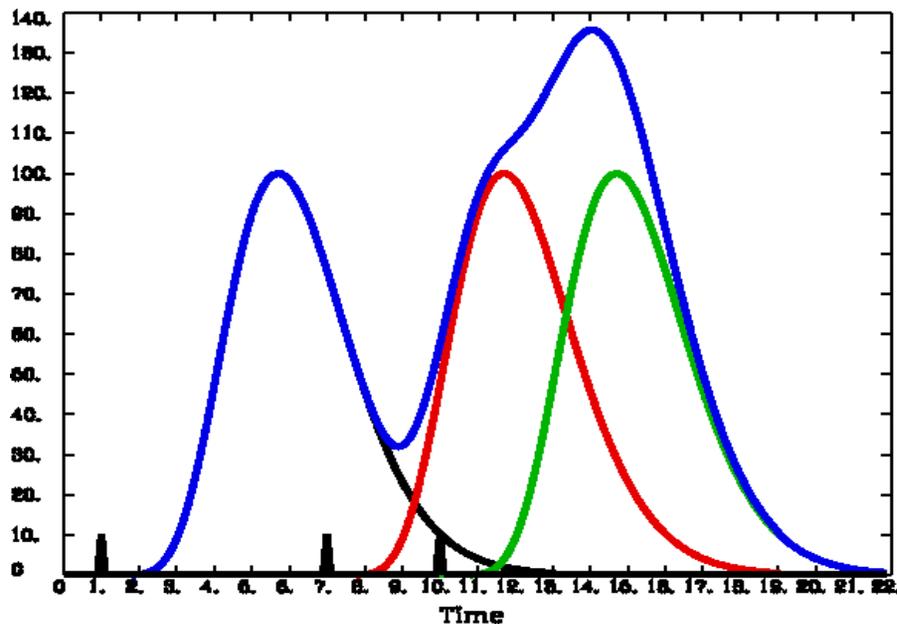
- Assuming a fixed shape  $h(t)$  for IRF to an **instantaneous** stimulus
  - For each block,  $h(t)$  is convolved with **stimulus timing** AND **duration** ( $d$ ) to get idealized response (temporal pattern) as an explanatory variable (regressor): **BLOCK**( $d,p$ )
    - Equivalent to convolving a series of consecutive events
    - Linearity assumed within each block: plateau-like response
    - $p$ : scale HDF to 1 for easy interpretation of  $\beta$



Block: 10 s on and 10 s off; TR=2 s; 150 time points

# Fixed-Shape HDF for Event-Related Design

- Fixed shape  $h(t)$  for IRF to an **instantaneous** stimulus
  - For multiple events of a condition/task,  $h(t)$  is convolved with **stimulus timing** to get idealized response (temporal pattern) as an explanatory variable (regressor):  $\text{GAM}(p,q)$ 
    - Linearity assumed when events are close with each other: overlapping impulse responses

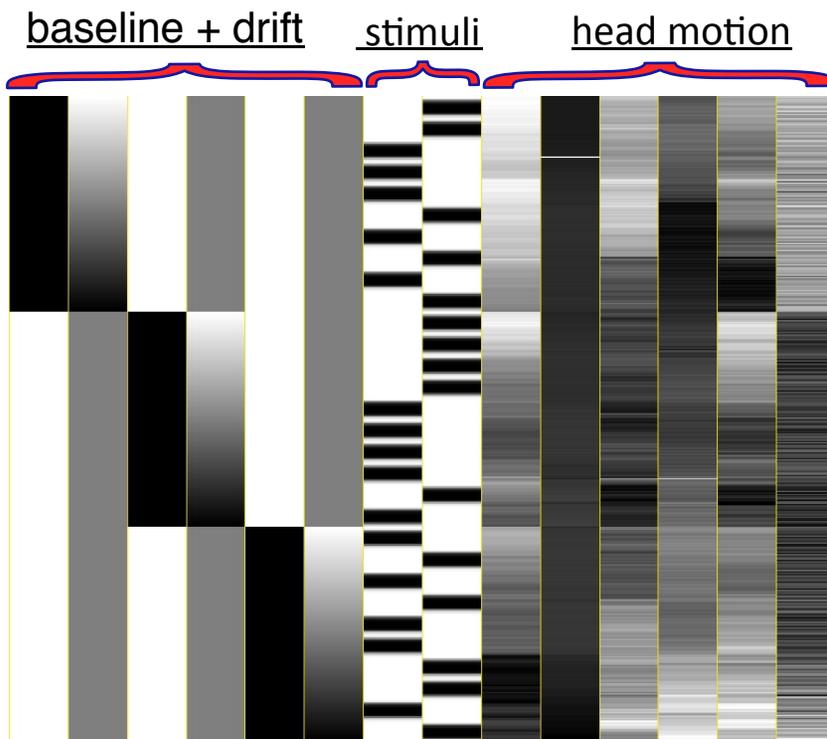


# Linear Model with Fixed-Shape IRF

- FMRI data = **baseline + drift + other effects of no interest + response<sub>1</sub> + ... + response<sub>k</sub> + noise**
- 'baseline' = **baseline** + drift + other effects of no interest
  - Drift: psychological / physiological effect, thermal fluctuation
  - Data = 'baseline' + **effects of interest** + noise
  - **Baseline condition (and drift)** is treated in AFNI as **baseline**, an additive effect, not an effect of interest (*cf.* SPM and FSL)!
- $y_i = \alpha_0 + \alpha_1 t_i + \alpha_2 t_i^2 + \beta_1 x_{1i} + \dots + \beta_k x_{ki} + \dots + \varepsilon_i$
- $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$ ,  $\mathbf{X} = [1, t, t^2, x_1, x_2, \dots, x_k, \dots]$
- In AFNI **baseline + slow drift** is modeled with polynomials
  - A longer run needs a higher order of polynomials
    - One order per 150 sec
  - With  $m$  runs,  $m$  separate sets of polynomials needed to account for temporal discontinuities across runs
    - $m(p+1)$  columns for **baseline + slow drift**: with  $p$ -order polynomials
- Other typical effects of no interest: head motion effects

# Design Matrix with Fixed-Shape IRF

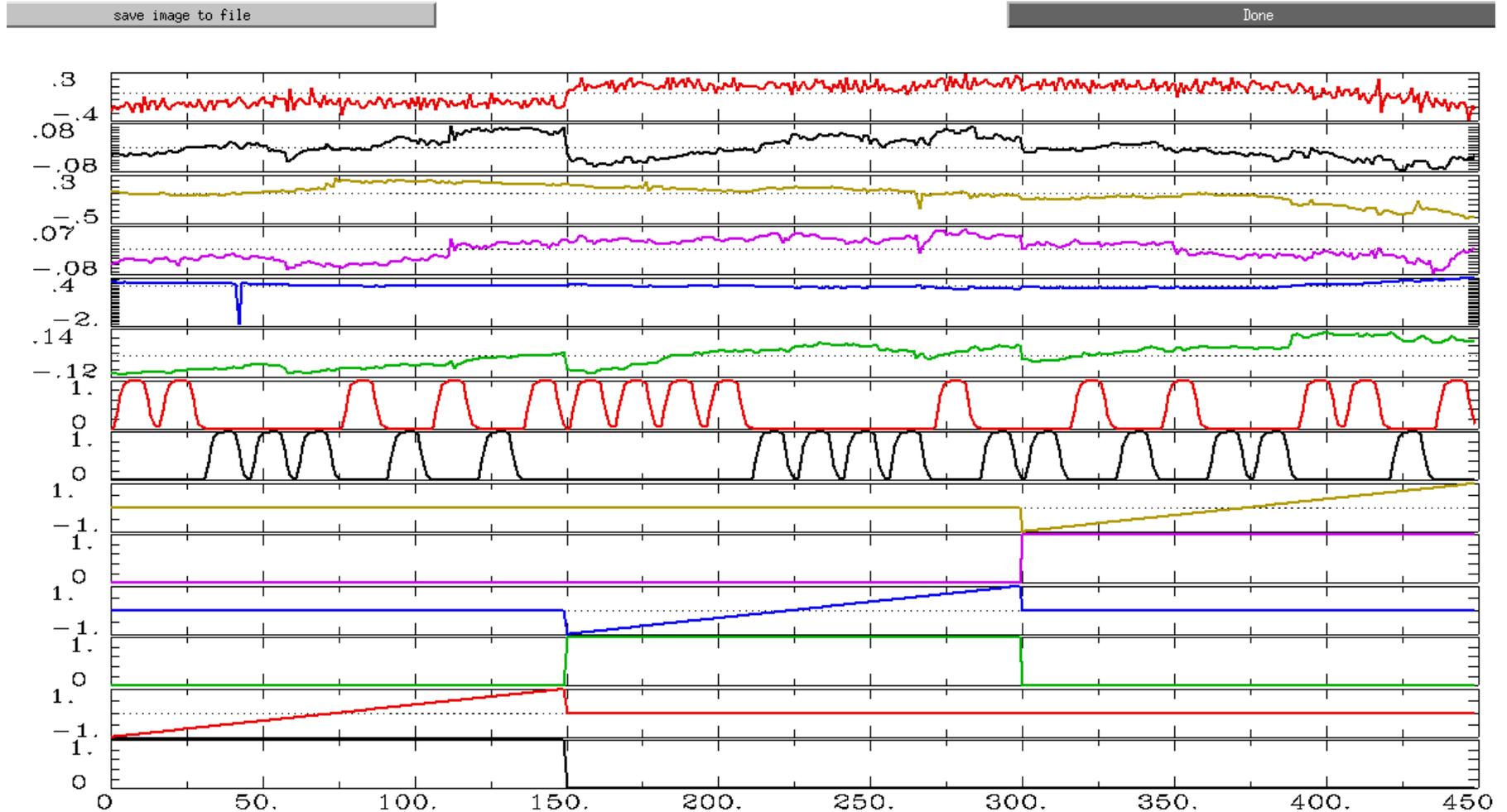
- Voxel-wise (massively univariate) linear model:  $y = X\beta + \varepsilon$ 
  - $X$ : explanatory variables (regressors) – **same** across voxels
  - $y$ : data (time series) at a voxel – **different** across voxels
  - $\beta$ : regression coefficients (effects) – **different** across voxels
  - $\varepsilon$ : anything we can't account for – **different** across voxels
- Visualizing design matrix  $X = [1, t, t^2, x_1, x_2, \dots, x_k, \dots]$  in grayscale



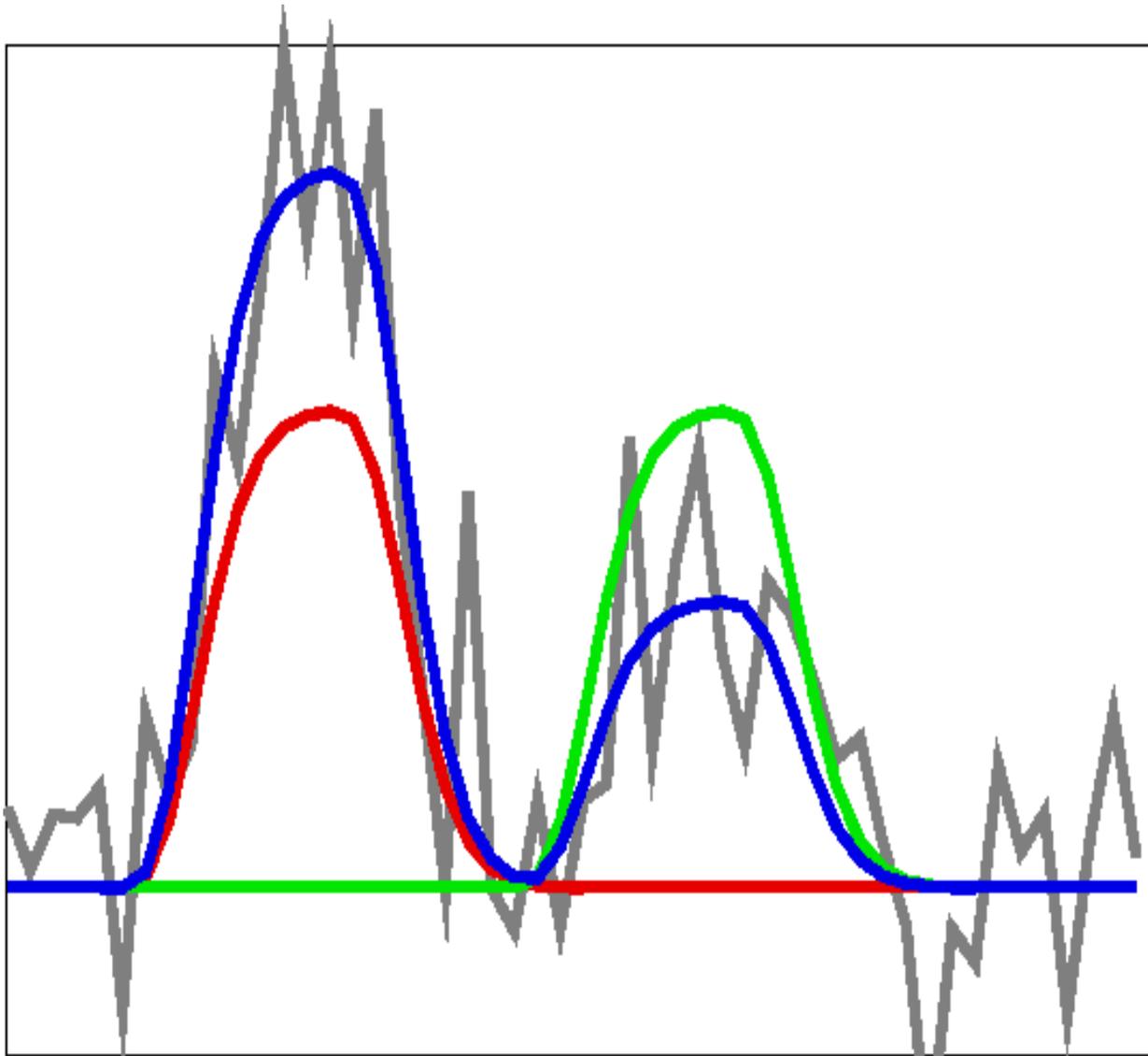
- 6 drift effect regressors
  - linear baseline
  - 3 runs x 2 parameters/run
- 2 regressors of interest
- 6 head motion regressors
  - 3 rotations + 3 shifts

# Design Matrix with Fixed-Shape IRF

- Visualizing design matrix  $X = [1, t, t^2, x_1, x_2, \dots, x_k, \dots]$  in curves



# Solving Linear Model



- Gray = data
  - Red = Expected response shape for #1
  - Green = Expected response shape for #2
  - Blue = modeled response =  $\beta_1 x_1 + \beta_2 x_2$  where  $\beta_1$  and  $\beta_2$  vary from 0.1 to 1.7
- 
- OLS: wiggle around and find  $\beta_1$  and  $\beta_2$  that make best fit with blue curve to data
- Gray = data =  $1.5x_1 + 0.6x_2 + \text{noise}$

# Model Quality Check

- First thing to do!
  - Unfortunately most users in fMRI simply jump to specific effects of interest, their contrasts and their significance. They simply don't pay any attention (or lip service) to overall model performance at all!
- Approaches to judge your model
  - Design matrix report from 3dDeconvolve
    - \*+ WARNING: !! in Signal-only matrix:
    - \* Largest singular value=2.37503
    - \* 7 singular values are less than cutoff=2.37503e-07
    - \* Implies strong collinearity in the matrix columns!
  - Full  $F$ -statistic (automatically provided in AFNI)
    - Data = 'baseline' + effects of interest + noise versus  
Data = 'baseline' + noise
  - Determination coefficient  $R^2$  at activated regions (-rout in 3dDeconvolve): poor modeling in fMRI!
    - Block design: ~50%
    - Event-related experiments: 10-20%
  - Modeled vs. not modeled: -fits and -errts in 3dDeconvolve
    - Fitted curve = 'baseline' + effects of interest
    - Residuals = noise = components we have no idea about

# Statistical Testing

- Everything is about contrast!
  - Even true for happiness in life
- Effects (regression coefficients) of interest
  - $\beta$ : effect relative to baseline condition by default in AFNI
    - $\beta_A = \text{Effect}_A - \beta_{\text{base}}$
  - $t$ -statistic: significance
- Pairwise comparisons (contrasts)
  - Conditions  $\beta_A$  vs.  $\beta_B$  (e.g., house vs. face)
    - $\beta_A - \beta_B = (\text{Effect}_A - \beta_{\text{base}}) - (\text{Effect}_B - \beta_{\text{base}}) = \text{Effect}_A - \text{Effect}_B$
  - $t$ -statistic: significance
- General linear test – linear combination of multiple effects
  - $t$ -statistic:  $0.5 \cdot \text{happy} + 0.5 \cdot \text{sad} - \text{neutral}$
- Composite tests
  - $F$ -statistic for composite null hypotheses:  $\text{happy} = \text{sad} = \text{neutral} = 0$ ;  
or,  $\text{happy} = \text{sad} = \text{neutral}$

# Assessing Fixed-Shape IRF Approach

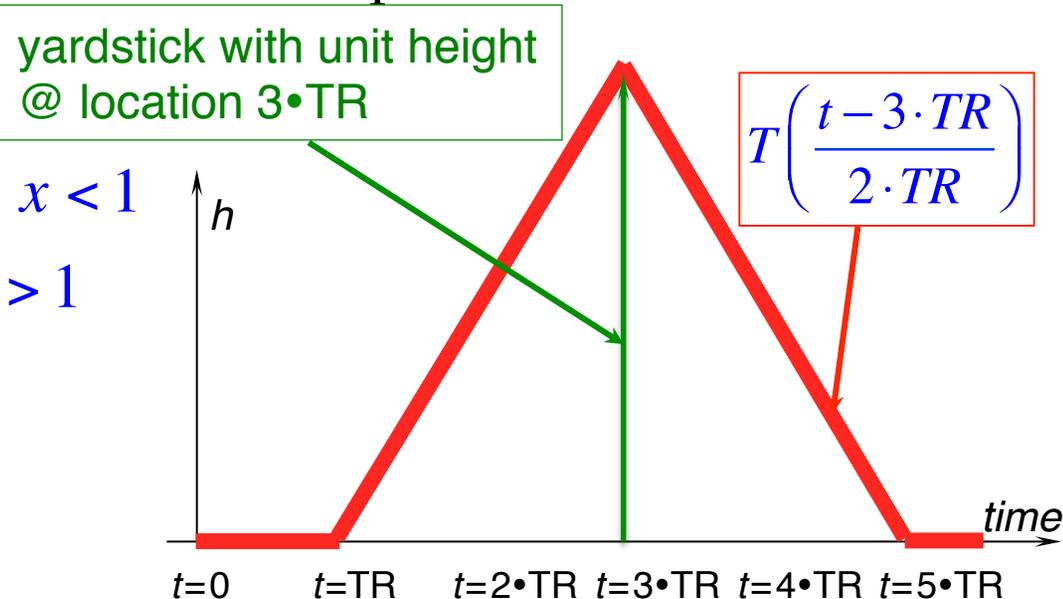
- Used 99% of time: Why popular?
  - Assume brain responds with **same shape** across 4 levels: **subjects**, activated **regions**, stimulus **conditions/tasks**, **trials**
    - Difference in **magnitude**  $\beta$  and its significance: what we focus on
    - Strong assumption about **four** levels of shape information?
  - Easy to handle: one value per effect
  - Works relatively well
    - Block design: shape usually not important due to accumulating effects (modeled via convolution) of consecutive events
      - Really plateau? Same magnitude across blocks?
    - Event-related experiment: OK most of time
      - Linearity when two responses overlap? Same effect across events?
- Not what you want if you
  - care about subtle shape difference across subjects, across regions, across conditions, and across trials
  - improve modeling

# No Constraint on IRF Shape

- Yardstick (or TENT) perspective
  - Set multiple yardsticks (or tents) at various equally-spaced locations to cover the potential BOLD response period
    - Each yardstick or TENT is a **basis function**
  - BOLD response measured by yardstick heights at all locations
    - Condition effect is reflected by as many as number of yardsticks
- Yardsticks (percent signal change sticks): TENT functions
  - Also known as 'piecewise linear splines'

$$T(x) = \begin{cases} 1 - |x| & \text{for } -1 < x < 1 \\ 0 & \text{for } |x| > 1 \end{cases}$$

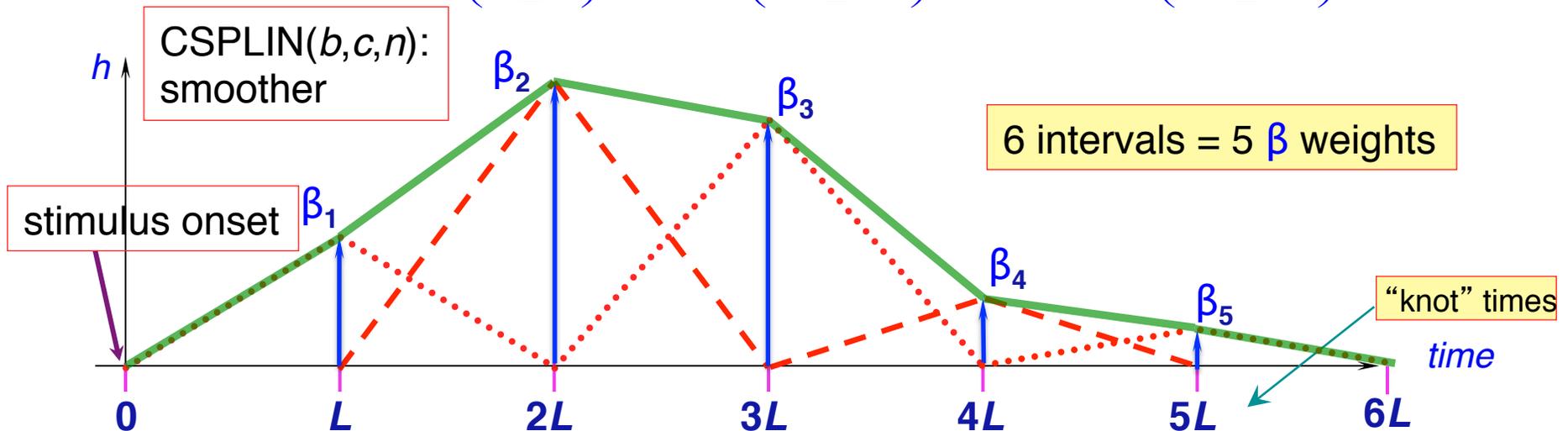
Cubic splines  
are also available



# Tent Functions = Linear Interpolation

- 5 equally-spaced tent functions (yardsticks): linear interpolation between “knots” with  $\text{TENTzero}(b,c,n) = \text{TENTzero}(0,12,7)$

$$h(t) = \beta_1 \cdot T\left(\frac{t-L}{L}\right) + \beta_2 \cdot T\left(\frac{t-2 \cdot L}{L}\right) + \dots + \beta_5 \cdot T\left(\frac{t-5 \cdot L}{L}\right)$$



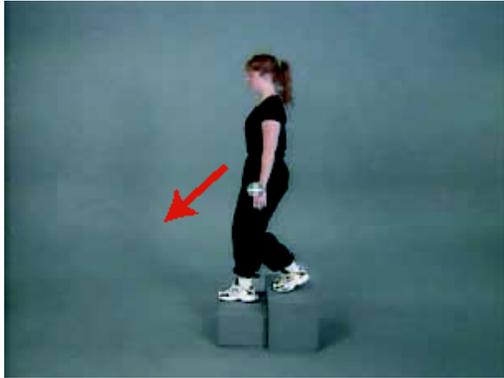
- Tent parameters are easily interpreted as function values (e.g.,  $L$ : tent radius;  $\beta_2$  = response (tent height) at time  $t = 2L$  after stimulus onset)
- Relationship of tent spacing  $L$  and TR ( $L \geq \text{TR}$ ), e.g., with TR=2s,  $L=2, 4$ s
- In `uber_subject.py` or `3dDeconvolve` with  $\text{TENTzero}(0, D, n)$ , specify duration ( $D$ ) of HRF and number ( $n$ ): radius  $L = D/(n-1)$  with  $(n-2)$  full tents, each tent overlaps half tent with two neighboring ones.
  - In above example,  $D=12$ s, then  $L=2$ s  $n=7$ ; covering 12s;  $\text{TENTzero}(0,12,7) \sim \text{TENT}(2,12,6)$

# Modeling with TENTs - Example

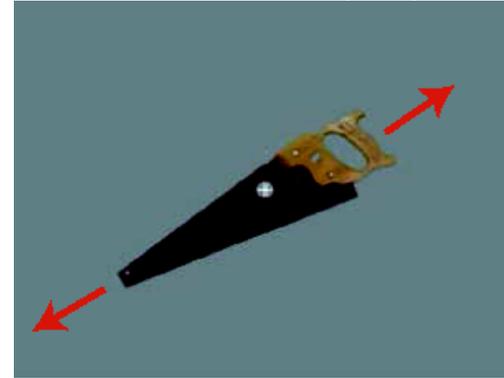
- Event-related study (Beauchamp et al., J Cogn Neurosci 15:991-1001)
  - 10 runs, 136 time points per run, TR=2 s
  - Two factors
    - Object type: human vs. tool
    - Object form: real image vs. points
  - 4 types (2x2 design) of stimuli (short videos)
    - Tools moving (e.g., a hammer pounding) - ToolMovie
    - People moving (e.g., jumping jacks) - HumanMovie
    - Points outlining tools moving (no objects, just points) - ToolPoint
    - Points outlining people moving - HumanPoint
  - Goal: find brain area that distinguishes natural motions (HumanMovie and HumanPoint) from simpler rigid motions (ToolMovie and ToolPoint)

- **Experiment: 2 x 2 design**

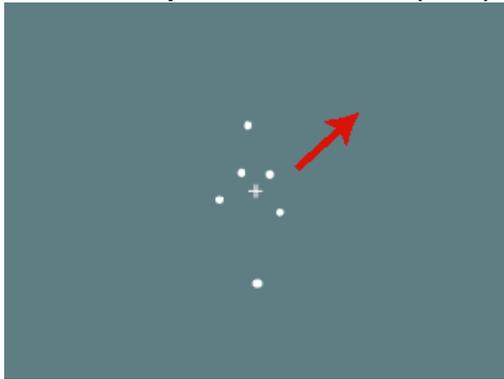
Human whole-body motion (HM)



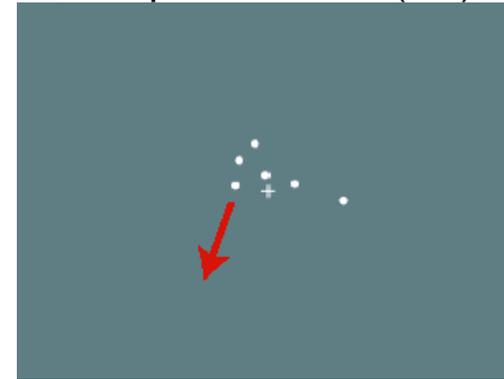
Tool motion (TM)



Human point motion (HP)



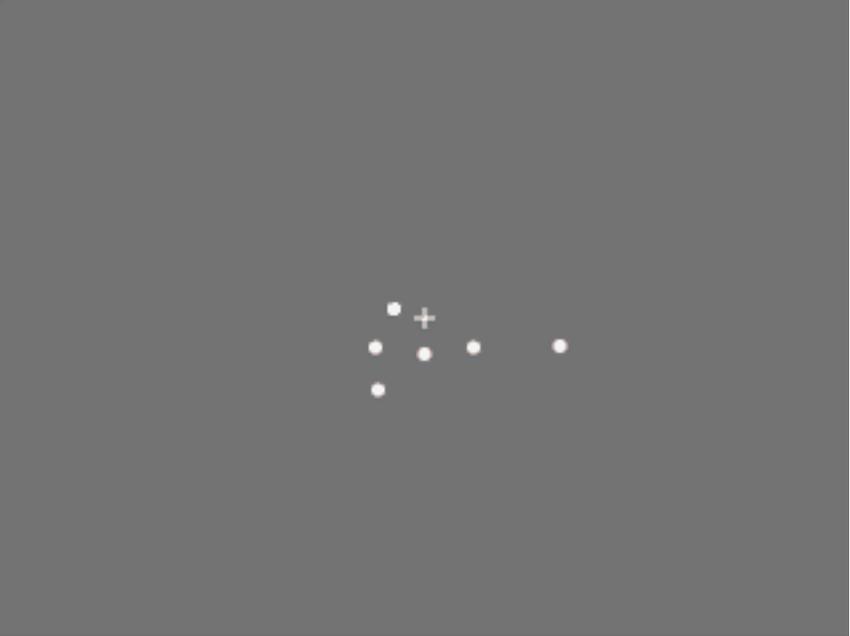
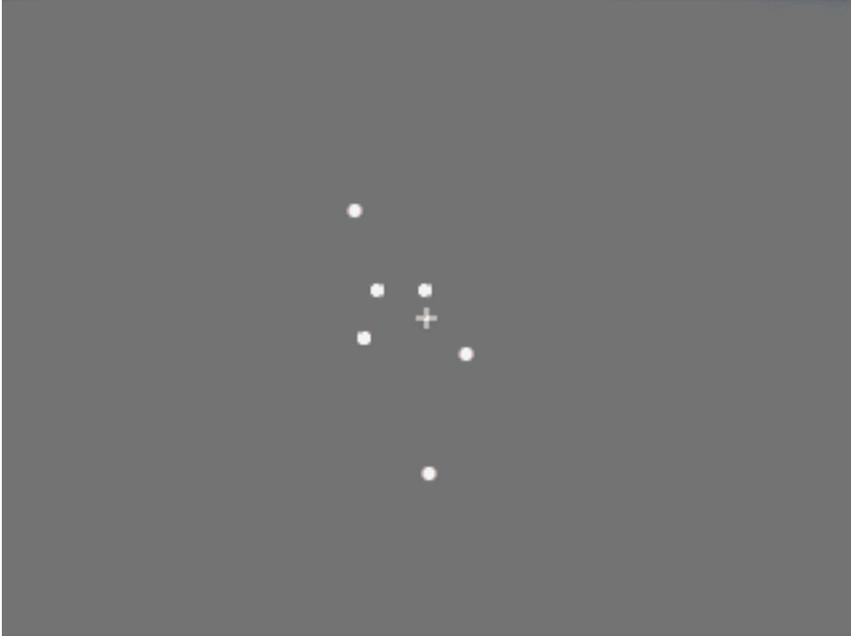
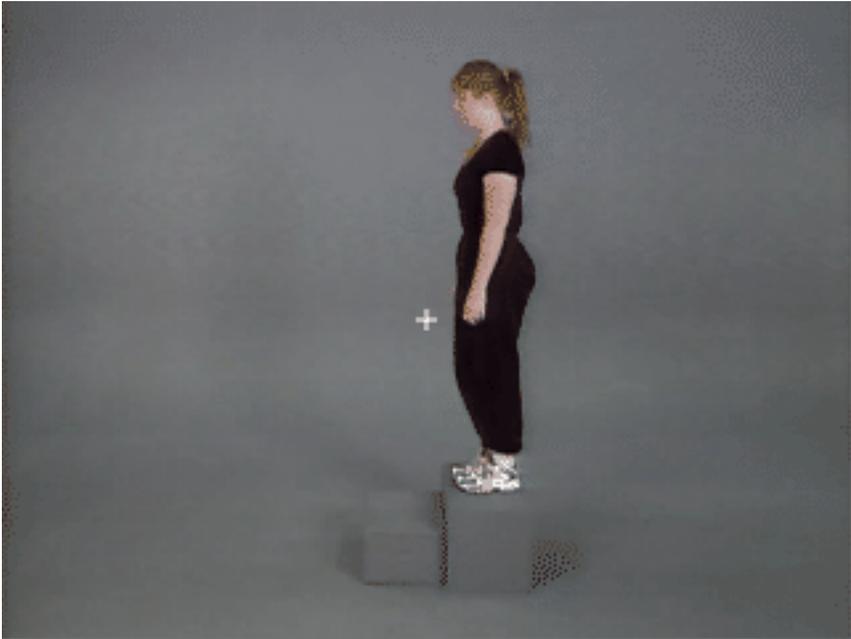
Tool point motion (TP)



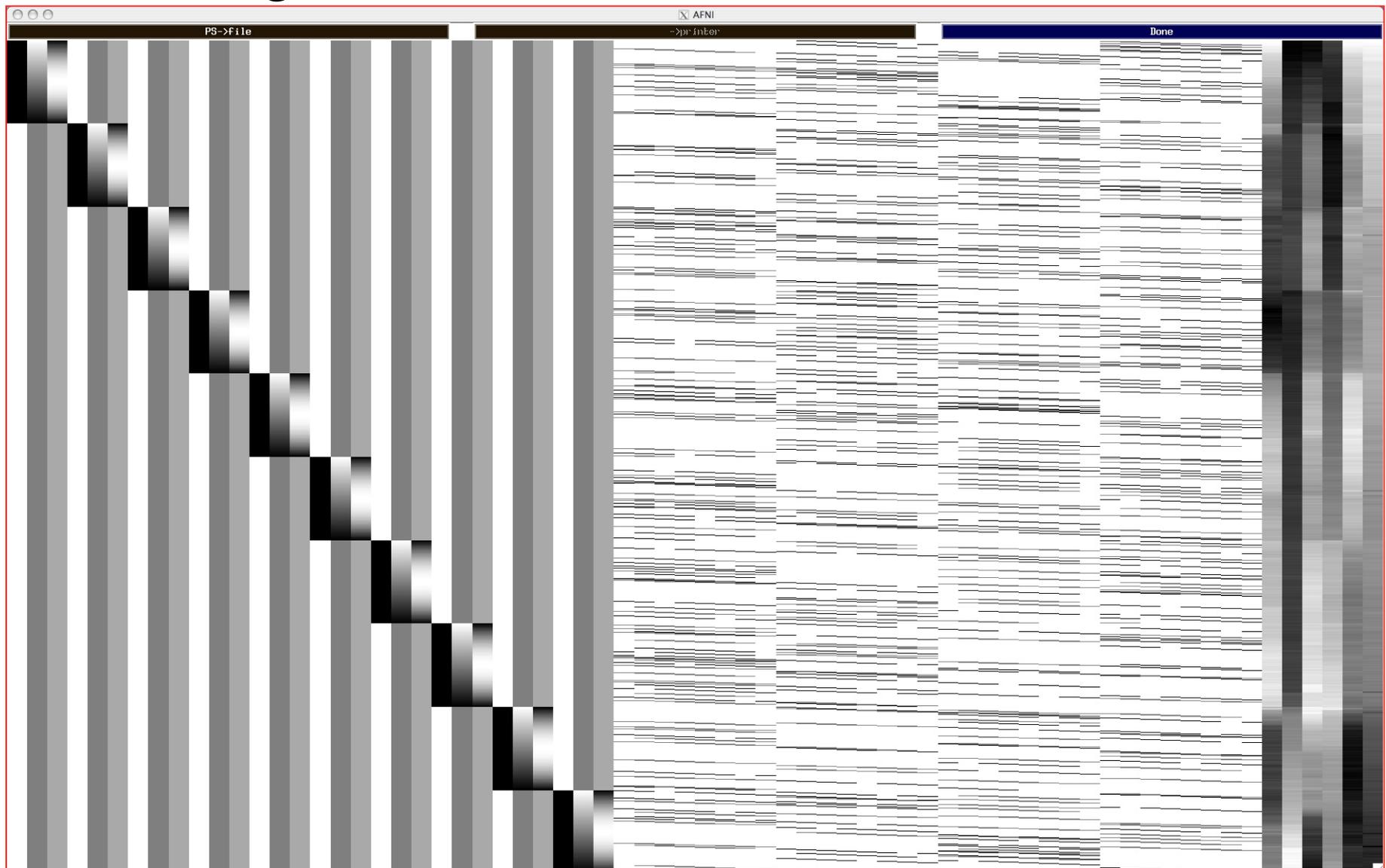
From Figure 1  
*Beauchamp et al. 03*

**Hypotheses to test:**

- Which areas are differentially activated by any of these stimuli (main effect)?
  - point motion versus natural motion? (type of image)
  - human-like versus tool-like motion? (type of motion)
- Interaction effects?
  - Point: human-like versus tool-like? Natural: human-like versus tool-like?
  - Human: point versus natural? Tool: point versus natural?



# Design Matrix with **TENTzero(0,16,9)**



Baseline + quadratic trend for 10 runs

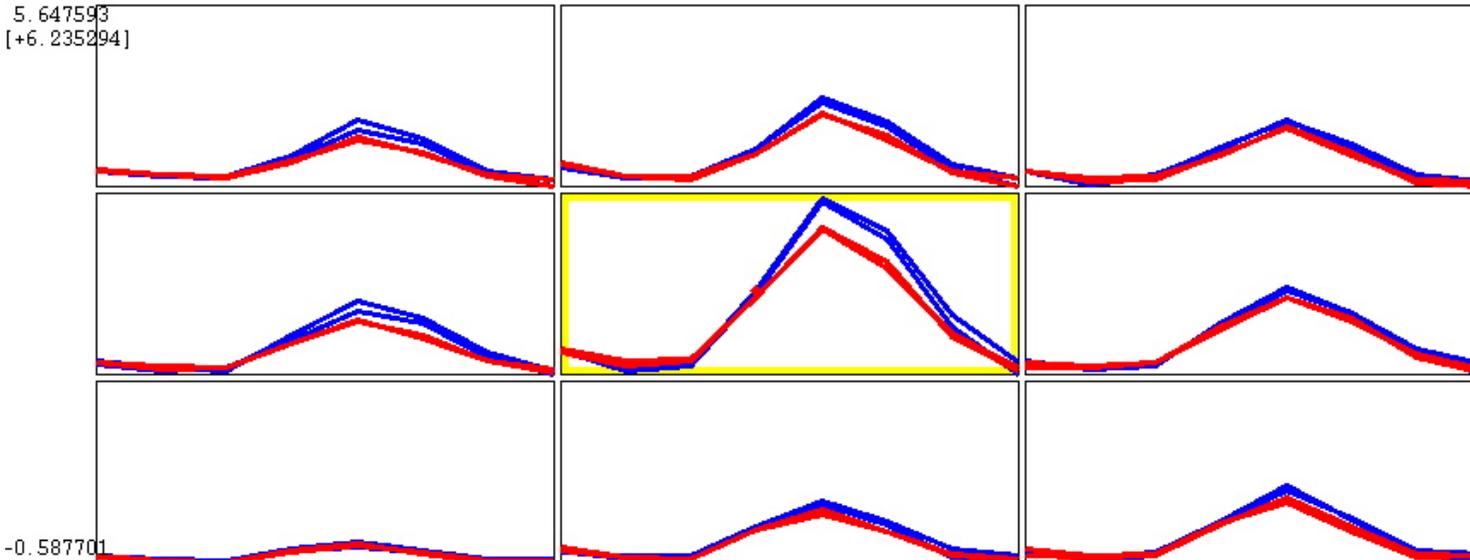
7 tents per condition  $\times$  4 conditions

head motion

# Results: Humans vs. Tools



5. 647593  
[+6. 235294]



- Color overlay: **Human vs Tool**

- **Blue** (upper) : Human

- **Red** (lower) : Tool

# No Constraint on IRF Shape: Deconvolution

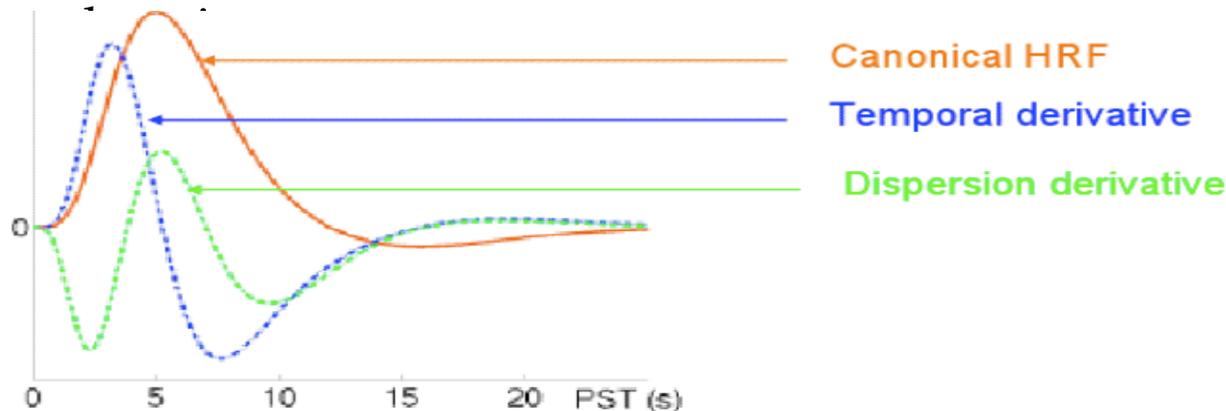
- Deconvolution perspectives: inverse process of convolution
  - **IRF**  $\otimes$  **stimulus** = **unit BOLD response**
    - Like multiplication, we have to know two and estimate the 3<sup>rd</sup>
  - Fixed-shape approach: Convolution + regression
    - Known: **impulse response**, **stimulus**
    - Use convolution to create **regressors** (hidden: waver or 3dDeconvolve)
    - Response strength ( $\beta$ ) estimated via linear model with 3dDeconvolve/3dREMLfit
  - Shape estimation: Deconvolution + regression
    - Known: **stimulus** + **BOLD response**; unknown: **impulse response**
    - **HRF**  $\otimes$  **stimulus** = **BOLD response** (note: HRF, not IRF)
    - **HDR** estimated as a linear combination of multiple yardsticks (basis TENT functions)
      - Each yardstick (TENT)  $\otimes$  stimulus = regressor
      - Deconvolution: **HDF** = a set of  $\beta$ 's estimated via regression

# No Constraint on IRF Shape: Pros + Cons

- What is the approach good at?
  - Usually for event-related experiments, but can be used for BLOCK
    - Multiple basis functions for blocks: within-block attenuation
  - Likely to have more accurate estimate on HDR shape across
    - subject
    - conditions/tasks
    - brain regions
  - Likely to have better model fit
  - Likely to be statistically more powerful on test significance
  - For block design, may detect within-block attenuation
    - Cross-block attenuation?
- Why is the approach not popular?
  - Difficult to summarize at group level
  - A few times more regressors than alternatives: DF's
  - Risk of highly correlated regressors: Multicollinearity
    - May change the number of basis functions
  - Overfitting: picking up something (head motion) unrelated to HDR

# Moderate Approach: SPMG1 / 2 / 3

- Balance in shape flexibility and basis functions
  - Constrain the HDR shape with a principal basis function
    - SPMG1 (similar to GAM in AFNI):  $e^{-t}(a_1 t^{p1} - a_2 * t^{p2})$  where  
 $a_1 = 0.0083333333$   $p1 = 5$  (main positive lobe)  
 $a_2 = 1.274527e-13$   $p2 = 15$  (undershoot part)
  - 2 or 3 basis functions: parsimonious, economical
    - SPMG1+SPMG2+SPMG3
    - SPMG2: temporal derivative capturing differences in peak latency
    - SPMG3: dispersion derivative capturing differences in peak



# Multicollinearity

- Voxel-wise regression model:  $y = X\beta + \varepsilon$ 
  - Regressors in design matrix  $X = [1, t, t^2, x_1, x_2, \dots, x_k, \dots]$
- Multicollinearity problem
  - Two or more regressors highly correlated
  - Difficult or impossible to tear apart the effects
- Multicollinearity scenarios
  - Collinearity -  $x_i = \lambda x_j$ : **model specification error**, e.g., 2 identical regressors (mistake in stimulus timing specifications)
  - Exact multicollinearity: linear among regressors: **faulty design** (rare)
  - High degree of correlation (+ or -) among regressors: **design problem**
    - E.g., cue + movie watching
  - Too many basis functions
- Diagnosis tools: ExamineXmat.R, timing\_tool.py, xmat\_tool.py

# Serial Correlation in Residuals

- Why temporal correlation?
  - In the residuals (not the time series data)
  - Short-term psychological and physiological effect
  - Other unknown reasons
- What is the impact of temporal correlation?
  - With white noise assumption,  $\beta$ 's are unbiased, but the statistics tend to be inflated
  - Little impact on group analysis if only taking  $\beta$ 's
  - May affect group analysis if considering effect reliability
- Approach in AFNI
  - ARMA(1,1) for residual time series
  - Slightly different from other packages

# Dealing with Multiple Runs

- Concatenation?
  - Analyze each run separately: AFNI, FSL
    - Have to have enough repetitions per run
    - Can test **cross-run** difference (trend, habituation) at **group** level
    - Summarize multiple  $\beta$ 's before group analysis
  - Concatenate but analyze with separate regressors across runs for each condition type: AFNI, SPM
    - Can test **cross-run** difference (trend, habituation, etc.) at **both individual and group** level
    - Summarize multiple  $\beta$ 's before group analysis
  - Concatenate but analyze with same regressor across runs for each condition type: default in AFNI
    - Assume no attenuation across runs
- Cross-block (or cross-event) attenuation
  - Crude method: `-stim_times_IM`

# Percent Signal Change

- Why conversion? Comparable across subjects
  - BOLD data don't have any physical / physiological meaning
  - Baseline is different across subjects
  - It's the relative changes that can be compared across subjects
- AFNI approach
  - Pre-processing: data scaled by voxel-wise mean
    - % signal change relative to mean, not exactly to baseline
    - Difference is tiny: less than 5%
  - Tied with modeling baseline as additive effects in AFNI
    - Sometimes baseline explicitly modeled in SPM and FSL
    - Global mean scaling (multiplicative) for whole brain drift
    - Grand mean scaling for cross-subject comparison: not %
    - Global and grand mean scaling, although not usually practiced, can be performed in AFNI if desirable

# Percent Signal Change

- Why not scaled  $\beta$ 's by real baseline???
  - No catenation: scale  $\beta$  per run by the run's baseline
    - Sample size in each run could be low
    - Have to summarize multiple  $\beta$ 's before group analysis
    - Better convert to percent signal change at run level before summing over runs
    - Be careful when motion parameters included in model
      - Uber\_subject.py automatically demeanes the head motion regressors
  - Catenation: **problematic**
    - Baseline may be different across runs
    - Effects are not comparable across runs

# Lackluster Performance in Modeling

- Essentially, all models are wrong, but some are useful (G.E.P. Box)
- Noisy data: easy excuse!
- Regressors: idealized response or yardstick
  - We get what we're looking for
  - It may miss something when we fail to recognize it
- Lots of variability across trials
  - Amplitude modulation if behavioral data are available
  - Model each trial separately
- Linearity assumptions
  - $\text{Data} = \text{baseline} + \text{drift} + \text{response1} + \text{response2} + \dots + \text{noise}$
  - When a trial is repeated, response is assumed same
  - Response for a block = linearity (no attenuation)
- Poor understanding of BOLD mechanism

# Summary

- Basics of linear model
- FMRI experiment types
  - Block design; Event related experiment; Mixed
- FMRI data decomposition: three components
  - Baseline + slow drift; Effects of interest; Unknown
  - Effects of interest - understanding BOLD vs. stimulus: IRF
- Modeling with fixed-shape IRF:  $GAM(p,q)$ ,  $BLOCK(d,p)$
- Modeling with no assumption about IRF shape
  - $TENT(b,c,n)$ ,  $CSPLIN(b,c,n)$
- Modeling with one major IRF plus shape adjustment
  - SPMG1/2/3
- Other issues
  - Multicollinearity
  - Catenation
  - Percent signal change