

Martin Lindquist

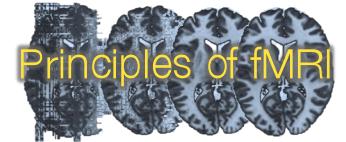
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Institute for Cognitive Science
University of Colorado, Boulder

Parametric modulation
designs for enhanced psychological
specificity

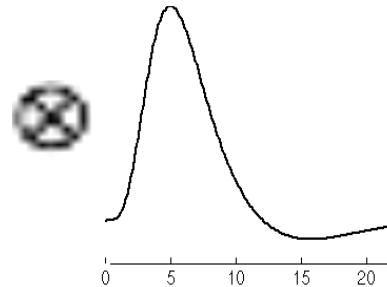
Convolution for multiple predictors: Discrete event types



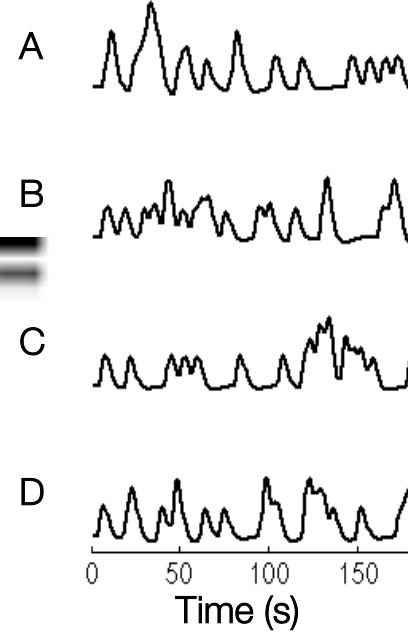
Indicator functions



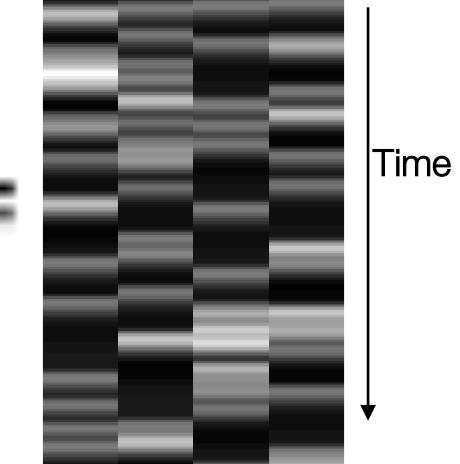
Assumed HRF
(Basis function)



Design Matrix (X^T)



Design Matrix (X)



Assumptions!

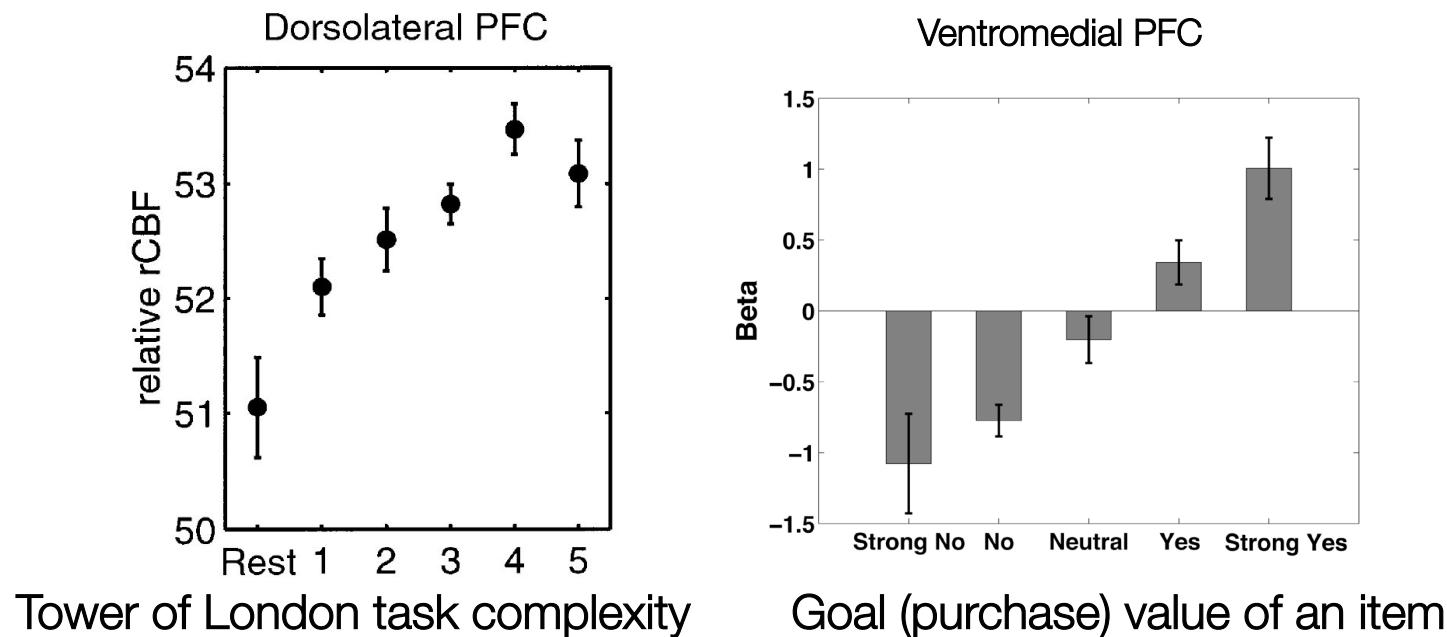
Assume neural activity function is correct

Assume HRF is correct

Assume LTI system

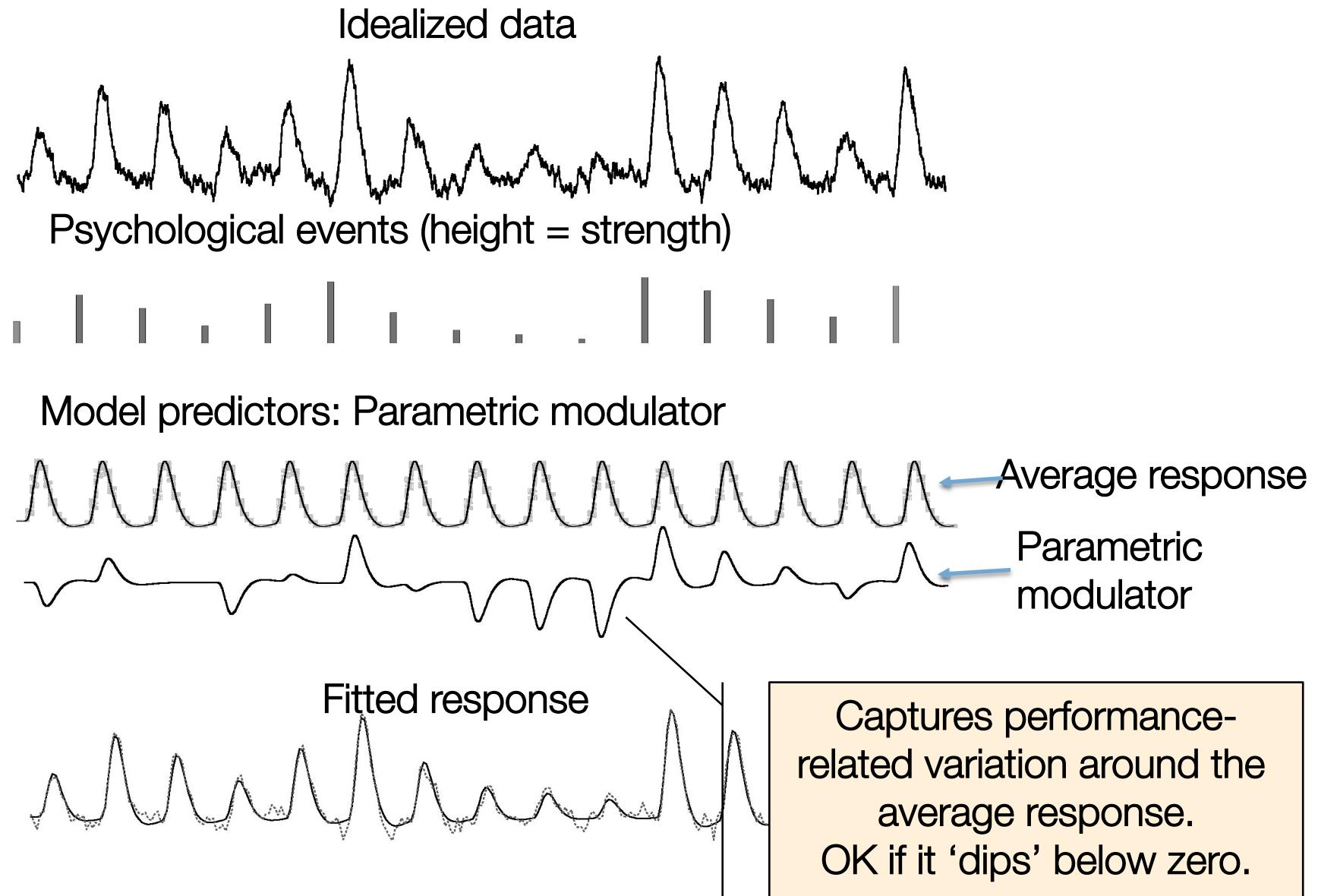
Parametric modulation

- Test for a linear relationship between brain activity and performance/ behavioral/psychological variable
- Modeling how brain activity varies across multiple levels can provide stronger evidence than contrasts for brain-performance relationships.



Dagher et al. 1999; Hare et al., 2009; Grinband et al., 2006; Wood et al., 2008

Parametric modulation



Key properties

- “Standard regressor” captures average event-related response
- Parametric modulator is orthogonal to the “standard regressor,” and so captures modulator variable-related activity above and beyond the average response
- Extendable:
 - Add linear modulators for time (e.g., linear change across trials), very useful for assessing habituation/fatigue and practice effects
 - Add other basis functions to capture some nonlinear effects across performance or time (e.g., quadratic or exponential terms)
- Implementation: All done by adding regressors to the GLM design matrix



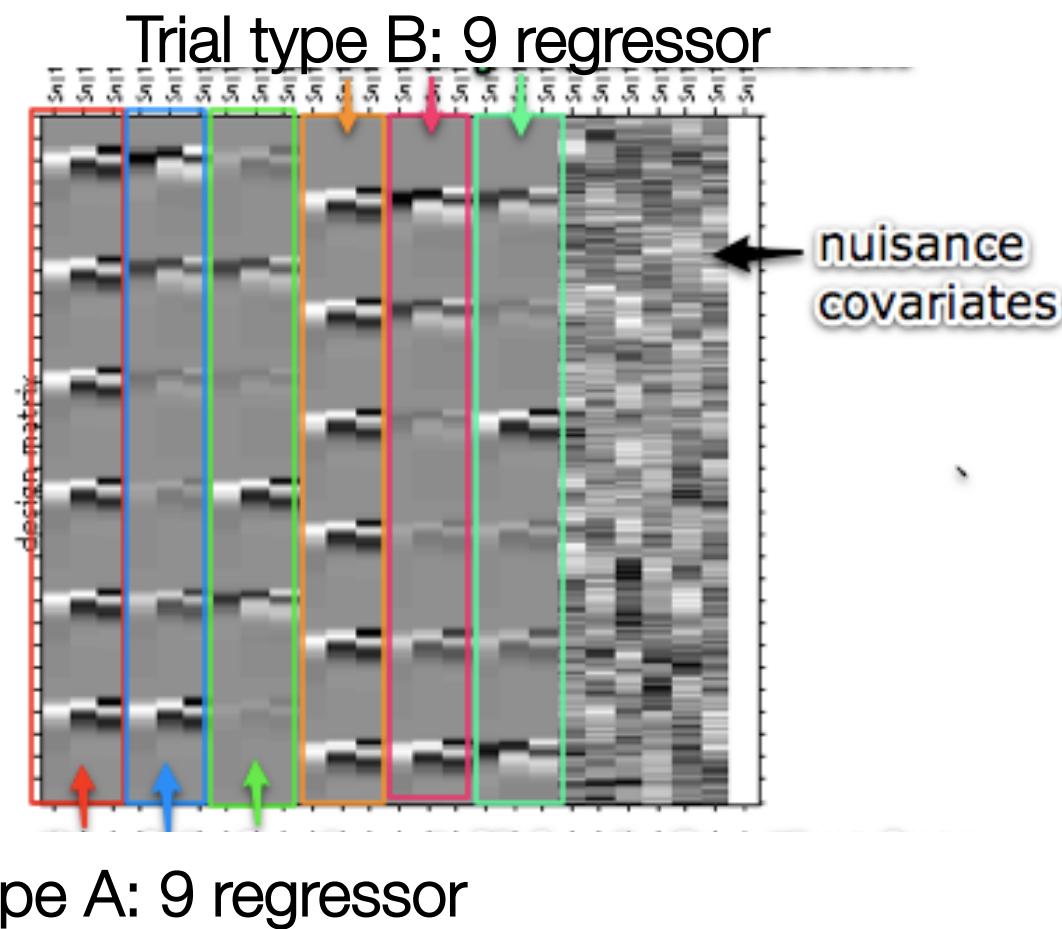
Cautions

- Standard approach is to modulate only the amplitude of a brief event ('stick function') or epoch
 - Ok for many purposes, but may not be accurate model
 - It is also possible to modulate the duration of an epoch with some tricks
 - As with other regressors, may not be straightforward to do a t-test that captures the parametric modulation effect if you have multiple basis functions.
-
- If you enter multiple modulators: Caution
 - In some software (e.g., SPM), subsequent modulators orthogonalized with respect to earlier ones, so that they are only allowed to explain variance not explained by other modulators.
 - This is not a standard multiple regression, but can be changed by modifying the code.



Parametric modulation

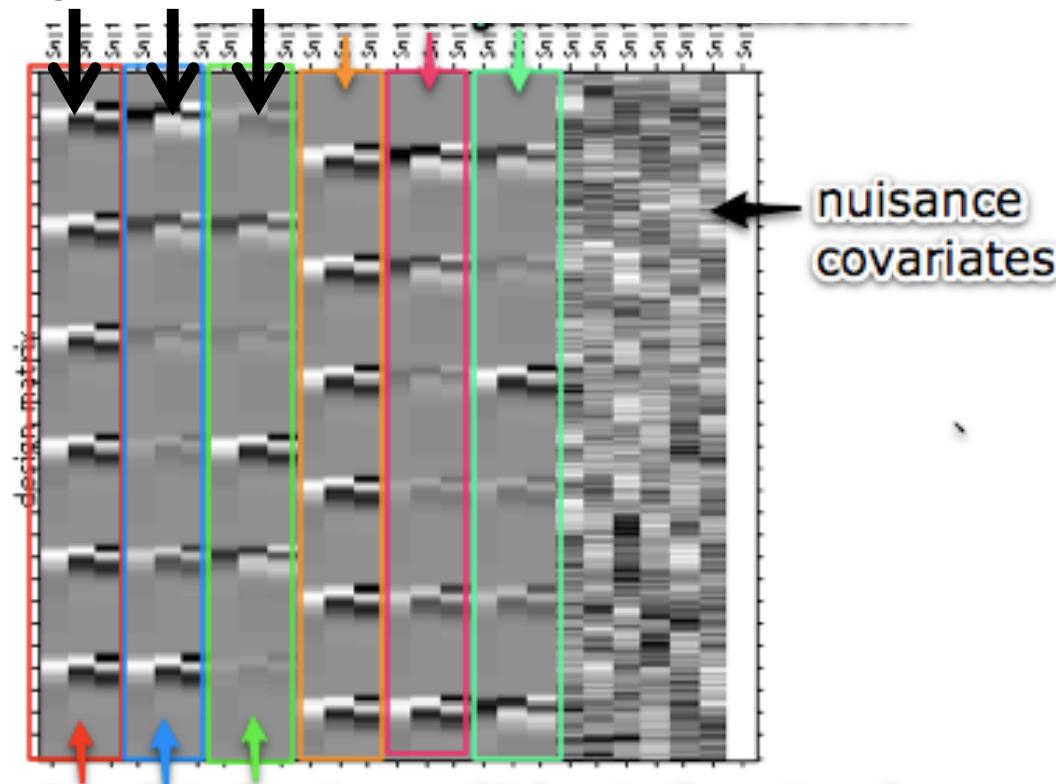
- Image of design matrix:



Parametric modulation

- Image of design matrix:

Average trial of the BOLD signal (shape)



Trial type A: 9 regressor

End of this section.



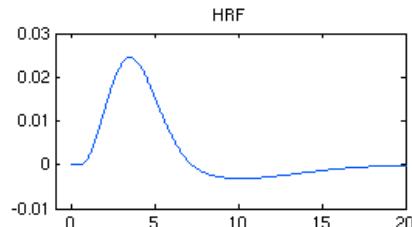
Key properties

- “Standard regressor” captures average event-related response
- Parametric modulator is orthogonal to the “standard regressor,” and so captures modulator variable-related activity above and beyond the average response
- Assumptions:
 - ▶ Linear brain responses based on input variables; sensitive to interval scaling properties of input variables
 - ▶ Only stick function amplitude is modulated; may not capture changes in shape/duration
- Subsequent modulators orthogonalized wrt earlier ones, so that they are only allowed to explain variance not explained by other modulators
 - ▶ This is done by a call to `spm_orth.m` and can be changed by modifying the code)

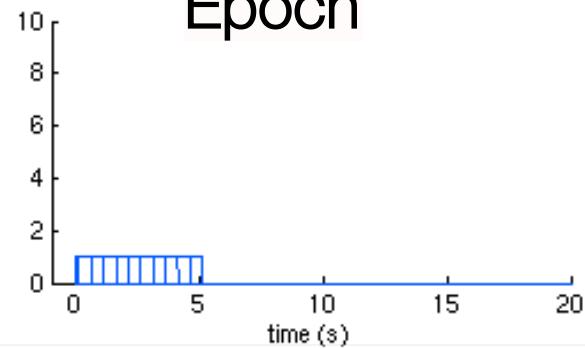
Parametric modulation of
duration vs. amplitude

Choosing a stimulus model: Events vs. epochs

HRF

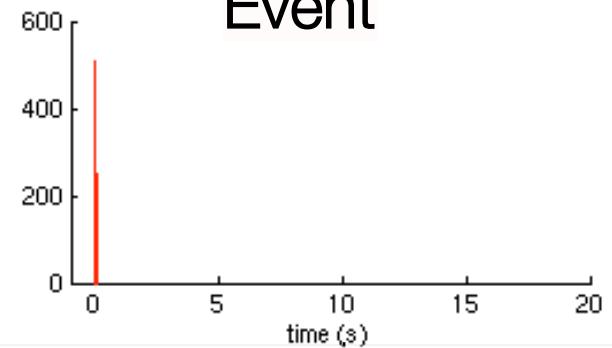


Epoch



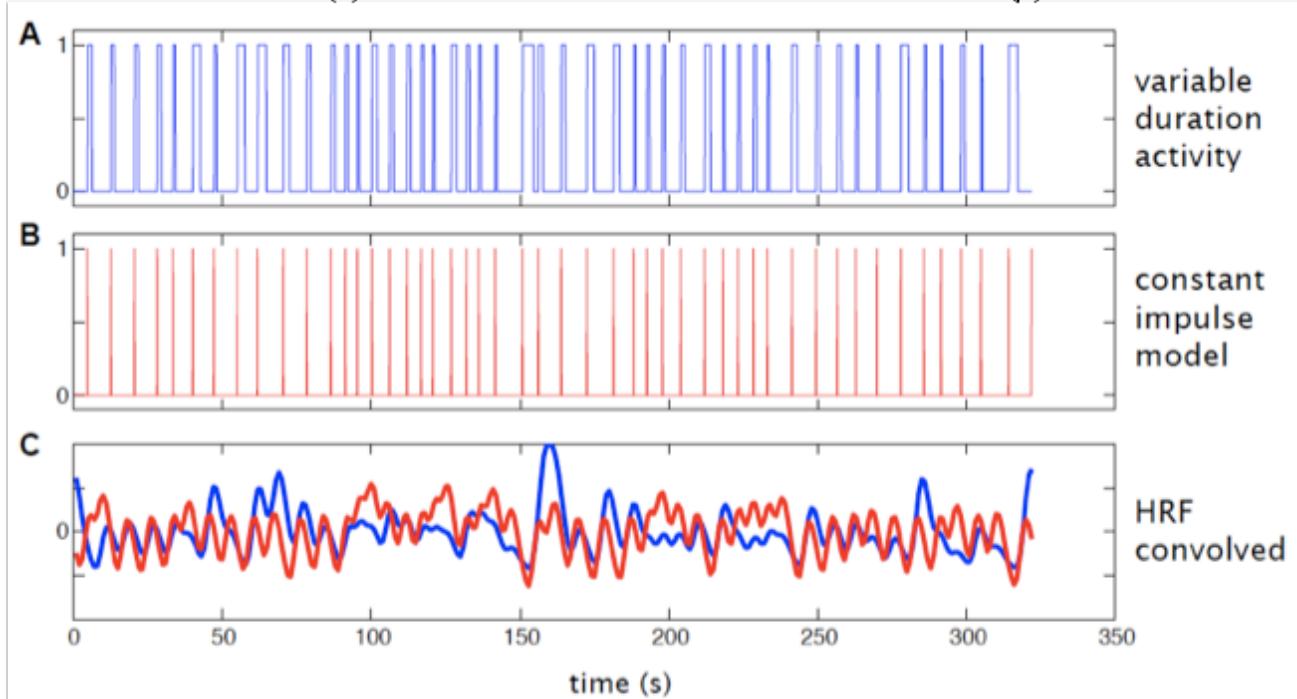
Stimulus Function

Event

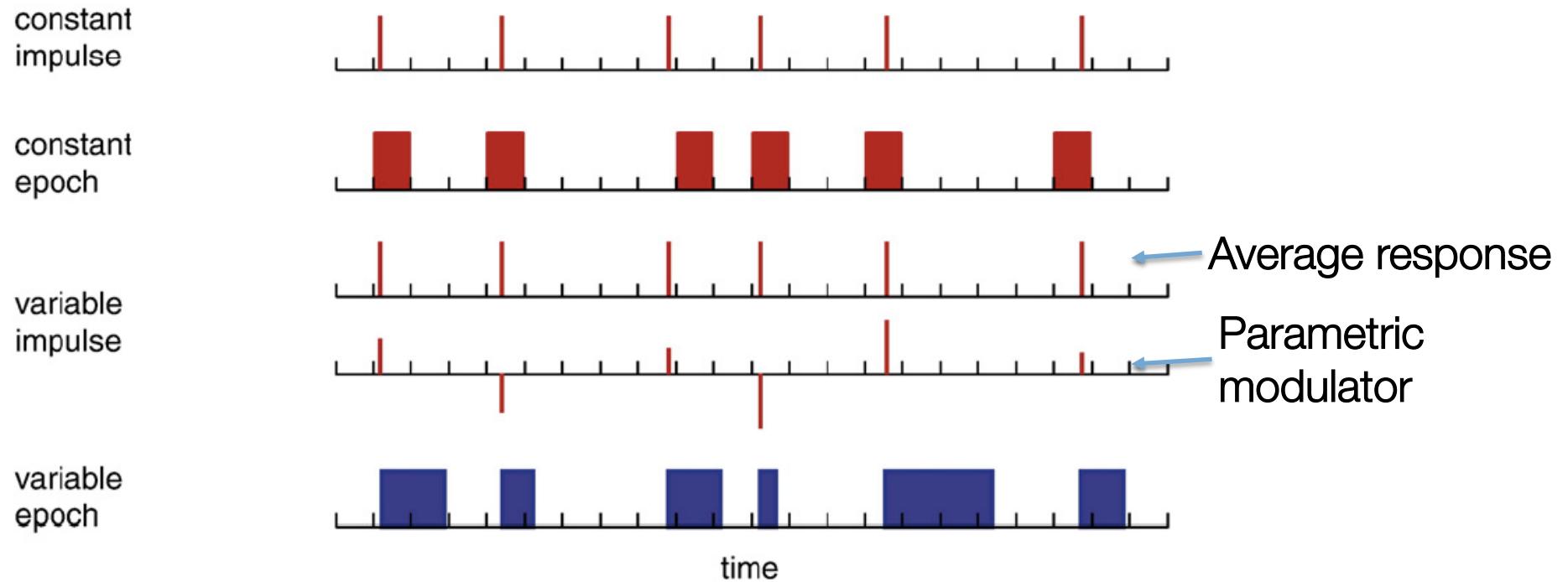


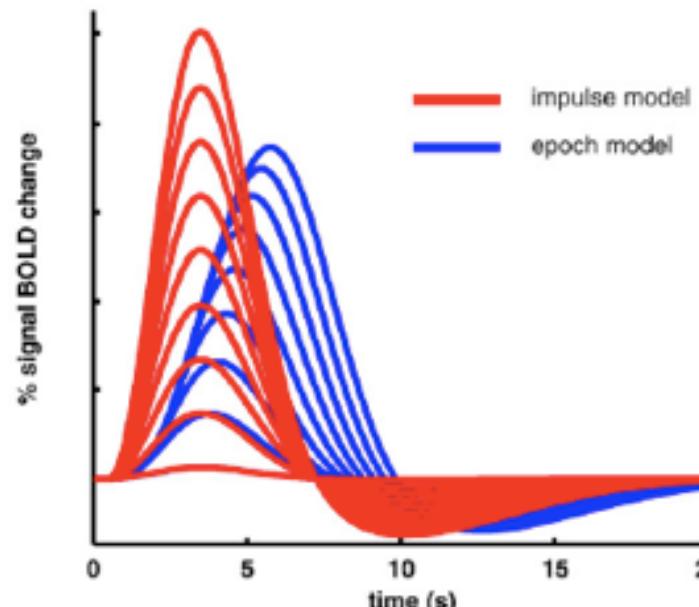
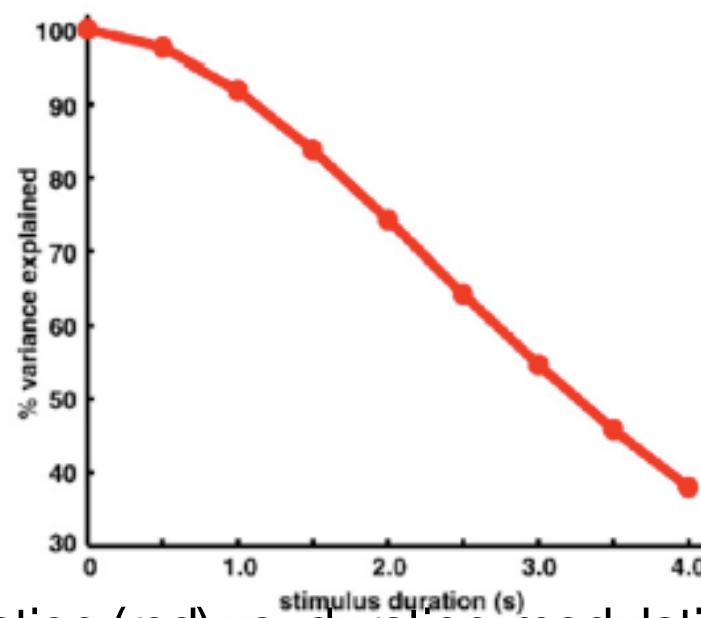
Does it matter for event-related fMRI?
* answer: More than you might think!

Grinband et al.,
2008



What to modulate: amplitude or duration?



A**B****C**

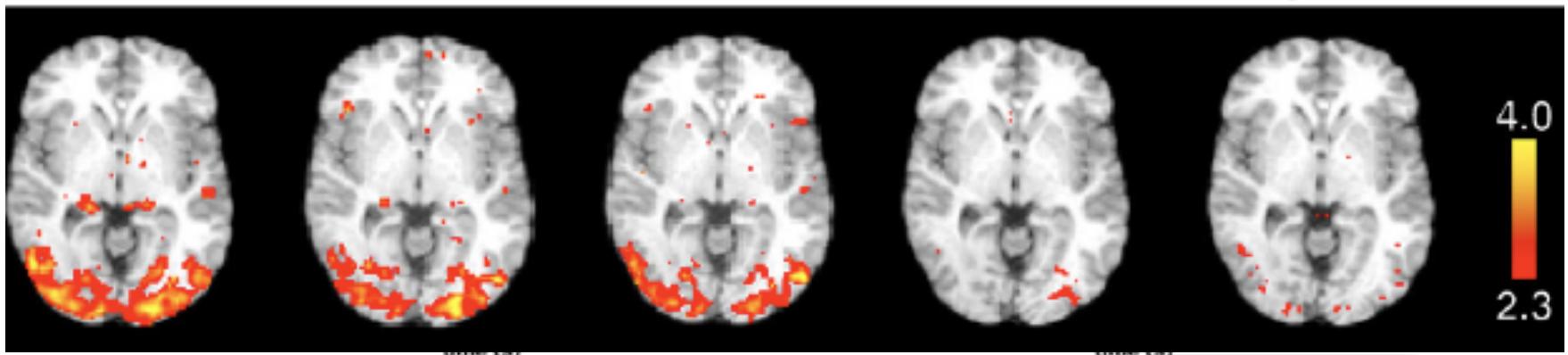
Visual cortex: Contrast-modulation (red) vs. duration-modulation (blue)

Variable
Epoch
Model

Constant
Epoch
Model

Constant
Impulse
Model

Variable Impulse Model
Constant
Regressor
Modulator
Regressor



Grinband et al., 2008



Key issue

- “Variable epoch” model is not orthogonalized with respect to the average trial response (i.e., there is no “standard regressor”)
- How do you test whether a behavioral modulator has an effect “above and beyond” the effect of the average trial response?
- Need to create both “standard regressors” and “variable epoch” regressors, and orthogonalize the 2nd wrt the 1st.
- You can do this using getPredictors (in the CANlab toolset / OptimizeDesign toolbox) and spm_orth (in SPM).
- For example code see:
http://www.scan.psych.columbia.edu:16080/pubwiki/doku.php/help/fMRI_help/fMRI_statistical_models/



```
% This code will turn a series of onset times and durations (e.g., RT)
% into a duration-modulated parametric modulator
% (See Grinband et al., 2008)
% You would add the two variables in X to the SPM design matrix, and repeat
% for each event type.
%
% Needed: SCN_Core_Support (onsets2fmridesign)
```

```
% The onsets and modulator are stored in SPM already, assuming:
% You have set up both time and parametric modulation, so that parametric
% modulation is the 2nd modulator (SPM.Sess.U(*).P(2).P)
% AND assuming you have entered things in units of seconds.
% First load SPM.mat. Then:
onsets_in_sec = SPM.Sess.U(1).ons; % for the first trial type
durations = SPM.Sess.U(1).P(2).P; % e.g., reaction time; the modulator
```

```
Xavg = onsets2fmridesign(onsets_in_sec, SPM.xY.RT, SPM.nscan, 'hrf');
Xdur = onsets2fmridesign([onsets_in_sec durations], SPM.xY.RT, SPM.nscan, 'hrf');
X = [Xavg(:, 1) Xdur(:, 1)];
figure; subplot(1, 2, 1); plot_matrix_cols(X);
title('Original (non-orthogonalized)')
```

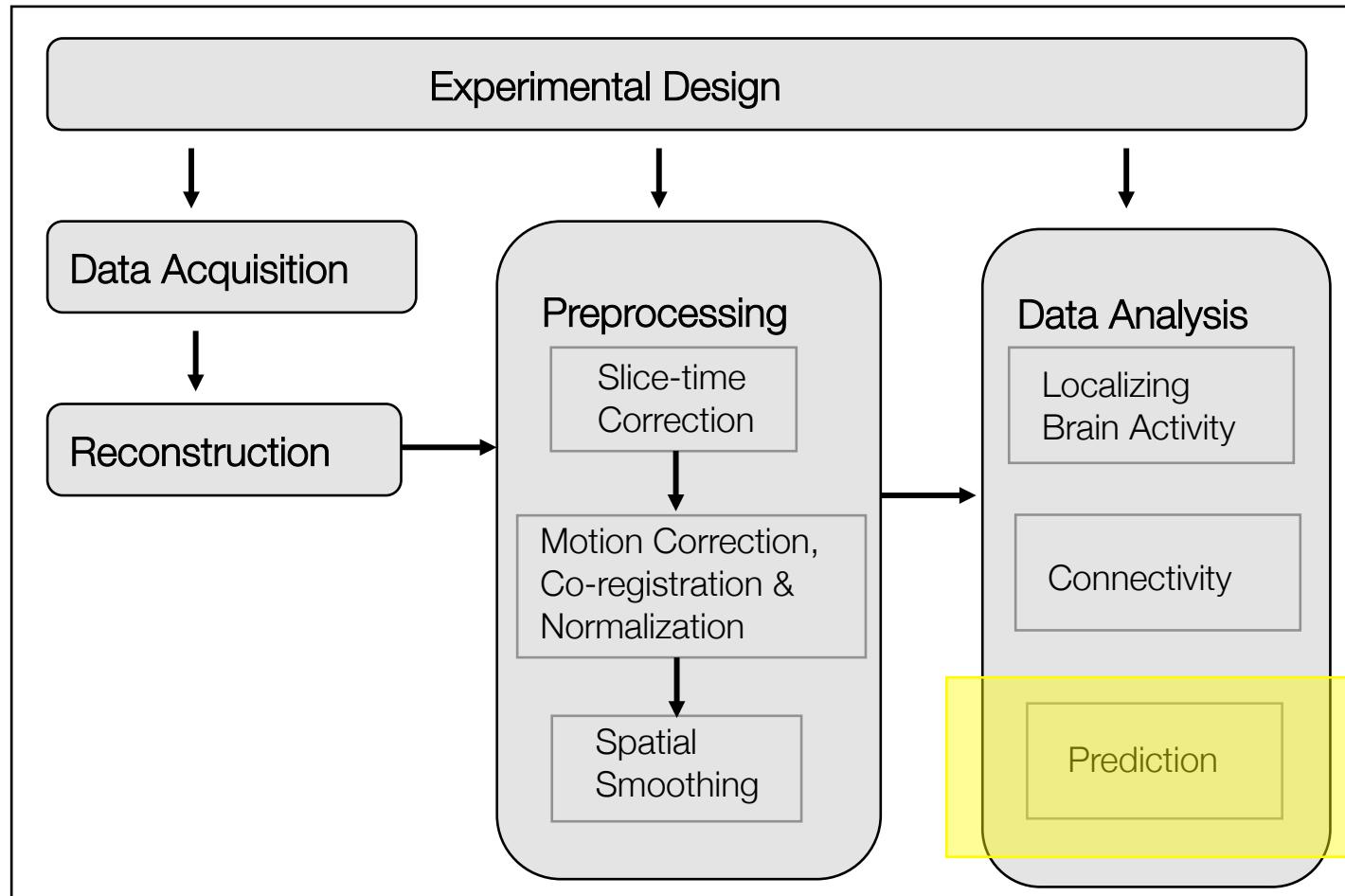
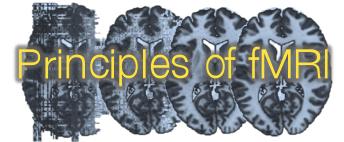
```
% We must orthogonalize the modulator with respect to the average response,
% so that significant effects for the modulator regressor are interpretable
% as duration-modulated effects above and beyond the average response to
% the events.
```

```
X = spm_orth(X); % orthogonalize
subplot(1, 2, 2); plot_matrix_cols(X)
title('Orthogonalized (add to SPM as regressors)')
```

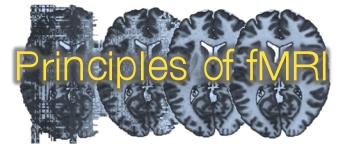
Example code

Multi-voxel Pattern Analysis

Data Processing Pipeline

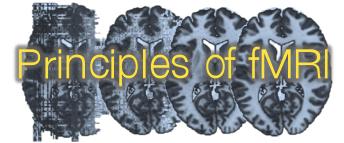


Classification and Prediction



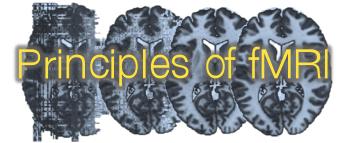
- There is a growing interest in using fMRI data for classification of mental disorders and predicting the early onset of disease.
- In addition, there is interest in developing methods for predicting stimuli directly from functional data.
- This opens the possibility of inferring information about subjective human experience directly from brain activation patterns.

Machine Learning

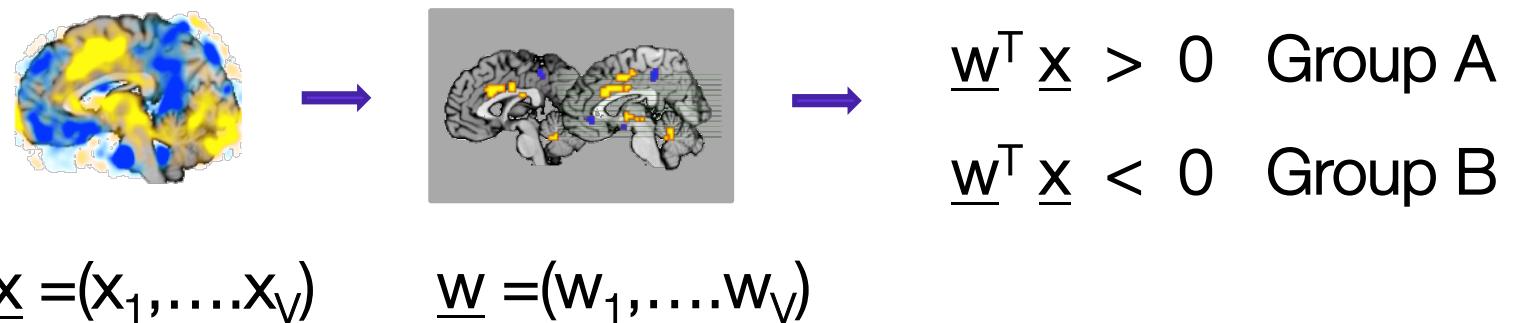


- Predicting brain states is challenging and requires the application of novel statistical and machine learning techniques.
- Various techniques have successfully been applied to fMRI data in which a **classifier** is trained to discriminate between different brain states and then used to predict the brain states in a new set of fMRI data.

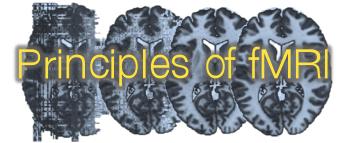
Machine Learning



- When applied to fMRI data the result is often a pattern of weights across brain regions that can be applied prospectively to new brain activation maps to quantify the degree to which the pattern responds to a particular type of event.



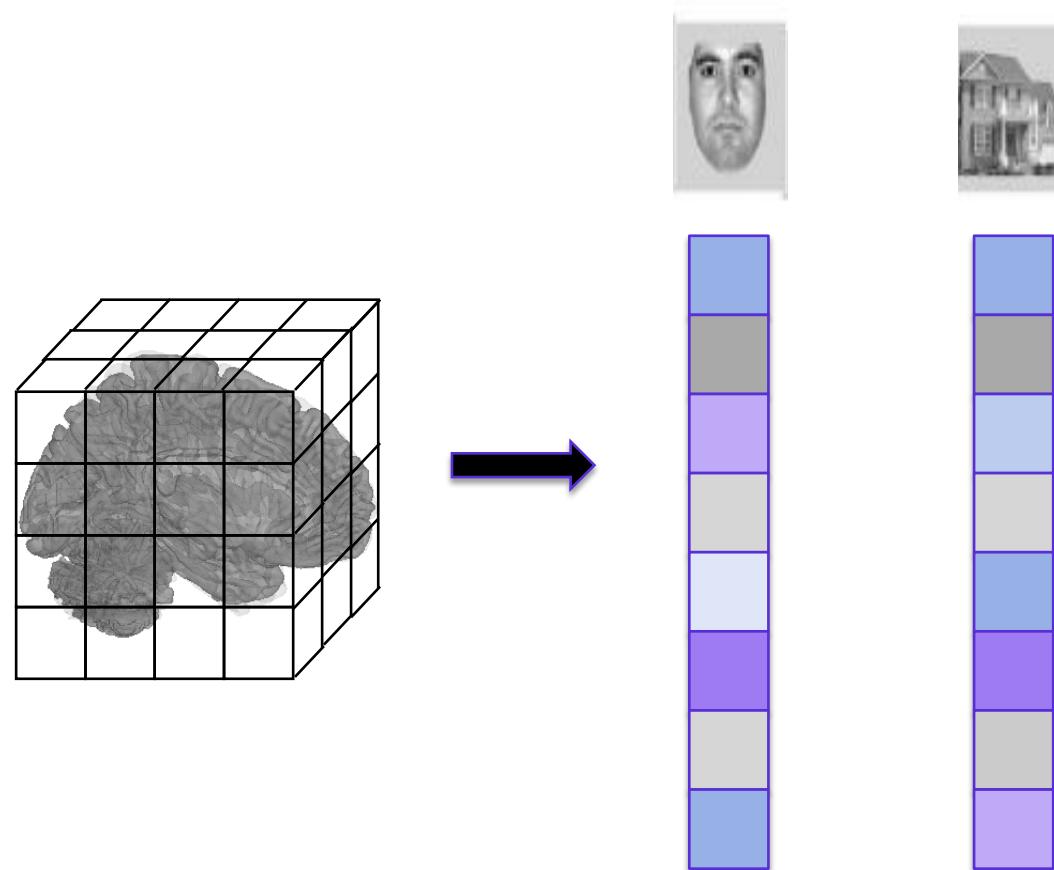
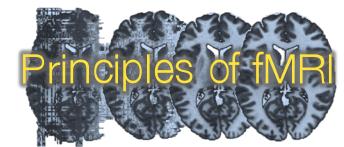
MVPA



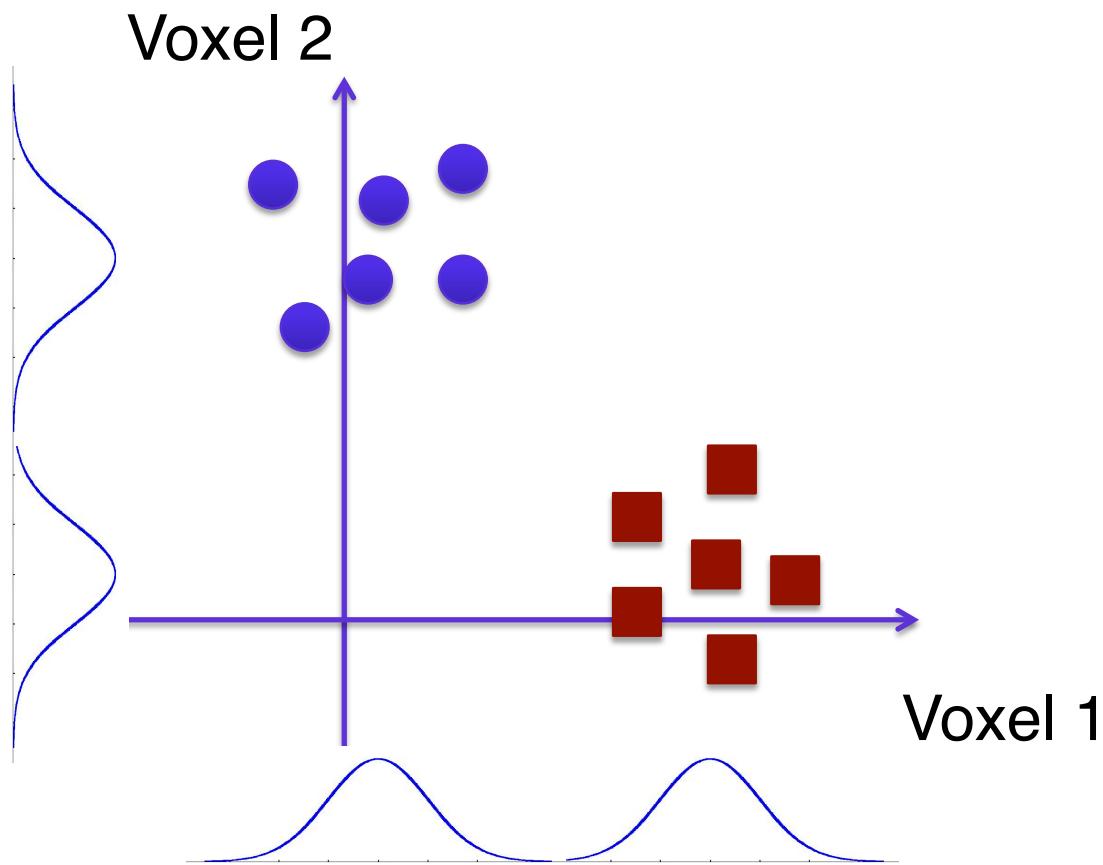
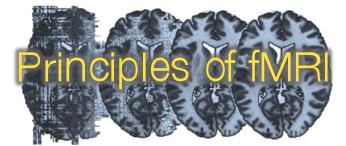
- The application of machine learning methods to fMRI data is often referred to as **multi-voxel pattern analysis** (MVPA)
- Instead of focusing on single voxels, MVPA uses pattern-classification algorithms applied to multiple voxels to decode the patterns of activity.



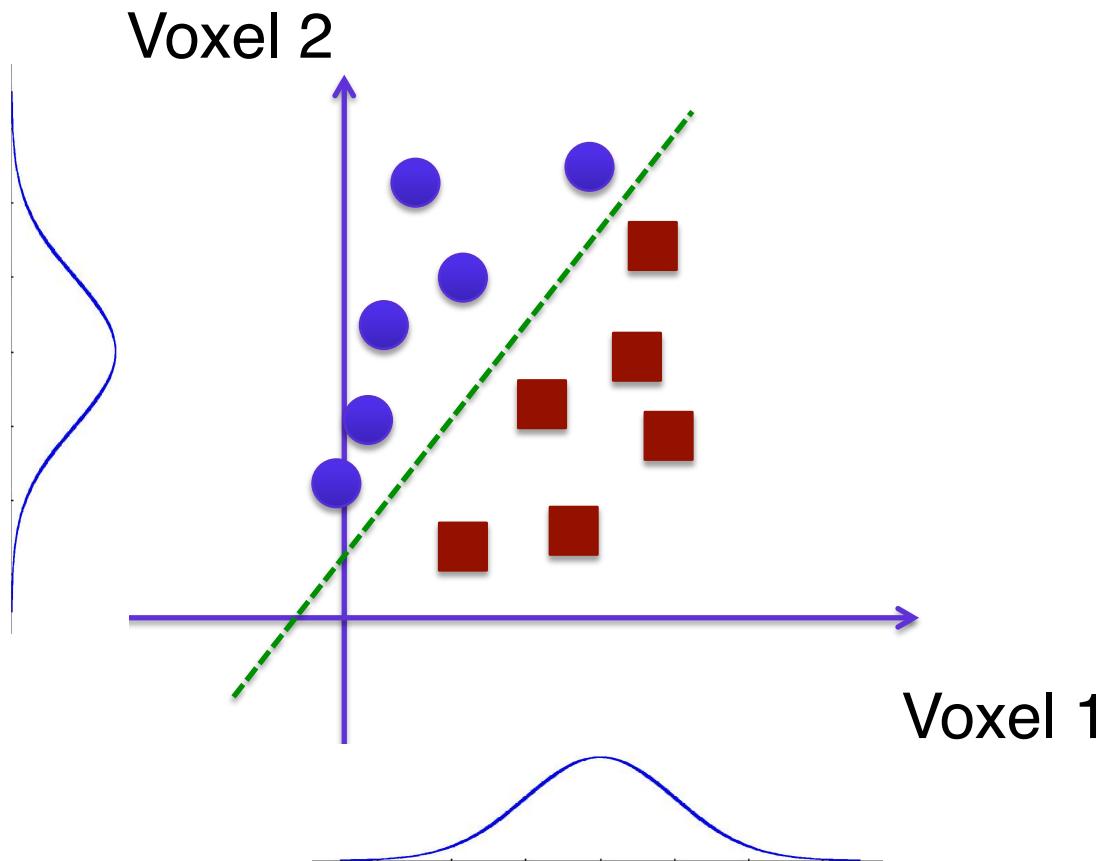
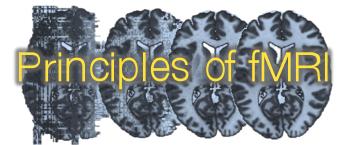
Multivariate Analysis



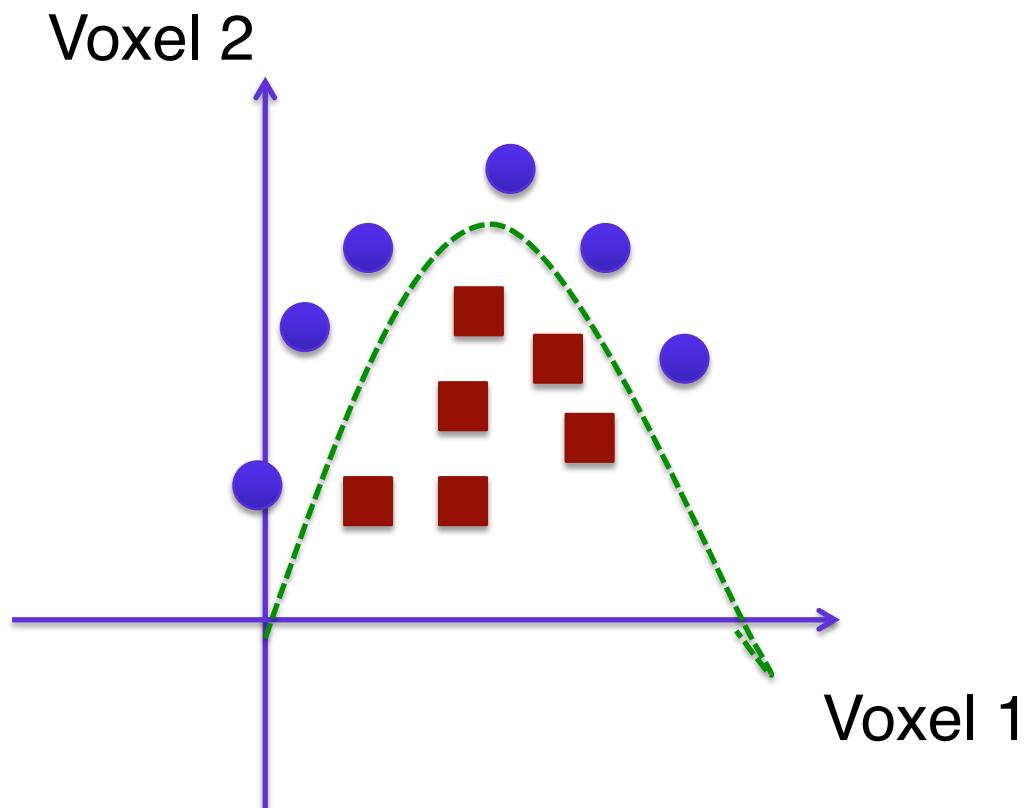
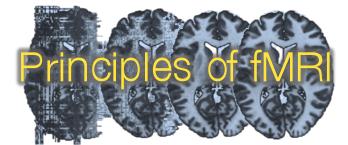
Multivariate Analysis



Multivariate Analysis



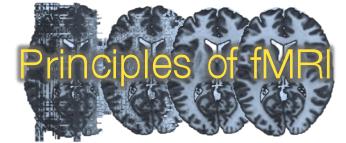
Multivariate Analysis



MVPA vs GLM

- In MVPA the goal is to determine the model parameters that allow for the most accurate prediction of new observations.
 - Seek to create rules that can be used to categorize new observations.
- In contrast, the GLM seeks to determine the model parameters that best fit the data at hand.

Classifiers



- A classifier is a function $f(\cdot)$ that takes the values of observed **features** (e.g., voxels) and predicts to which **class** the observation belongs (e.g., disease state).
- Let us denote the set of features $\underline{x} = (x_1, \dots, x_v)$ and the class label y .
- Predicted class: $\hat{y} = f(\underline{x})$

Training Data

- A classifier has a number of parameters w that needed to be estimated, or learned.
- The learning is typically performed on a subset of the observations called the [training data](#).
- The learned classifier models the relationship between the features and class labels in the training data set.



Test Data

- Once trained, the classifier is evaluated using an independent set of observations called the **test data**.
- If the classifier truly captures the relationship between features and classes, it should be able to predict the class label for data it hasn't seen before.
- The accuracy of the classifier measures the fraction of observations in the test data for which the correct label was predicted.

Illustration

Features (voxels)



Class Labels

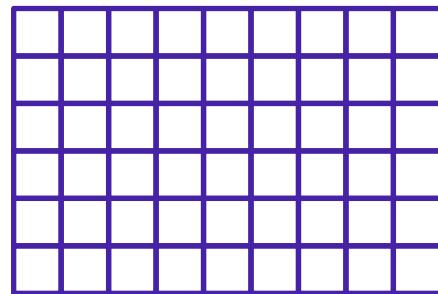


$$\underline{x} = (x_1, \dots, x_v)$$

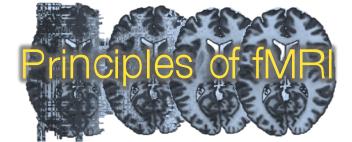
y

Data

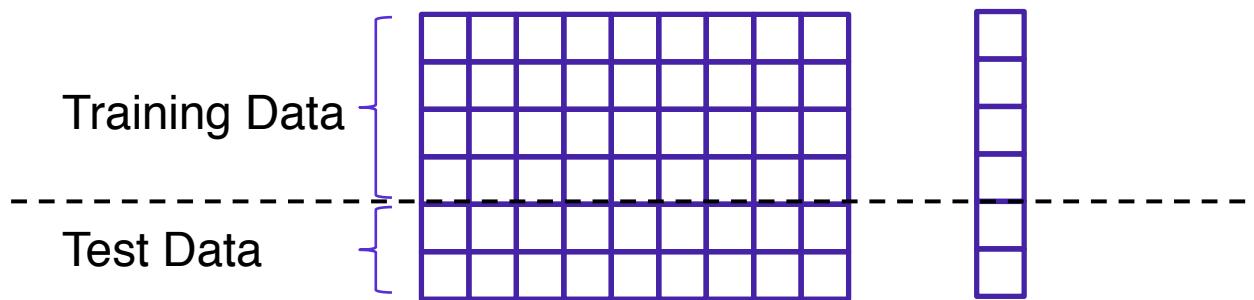
Observations



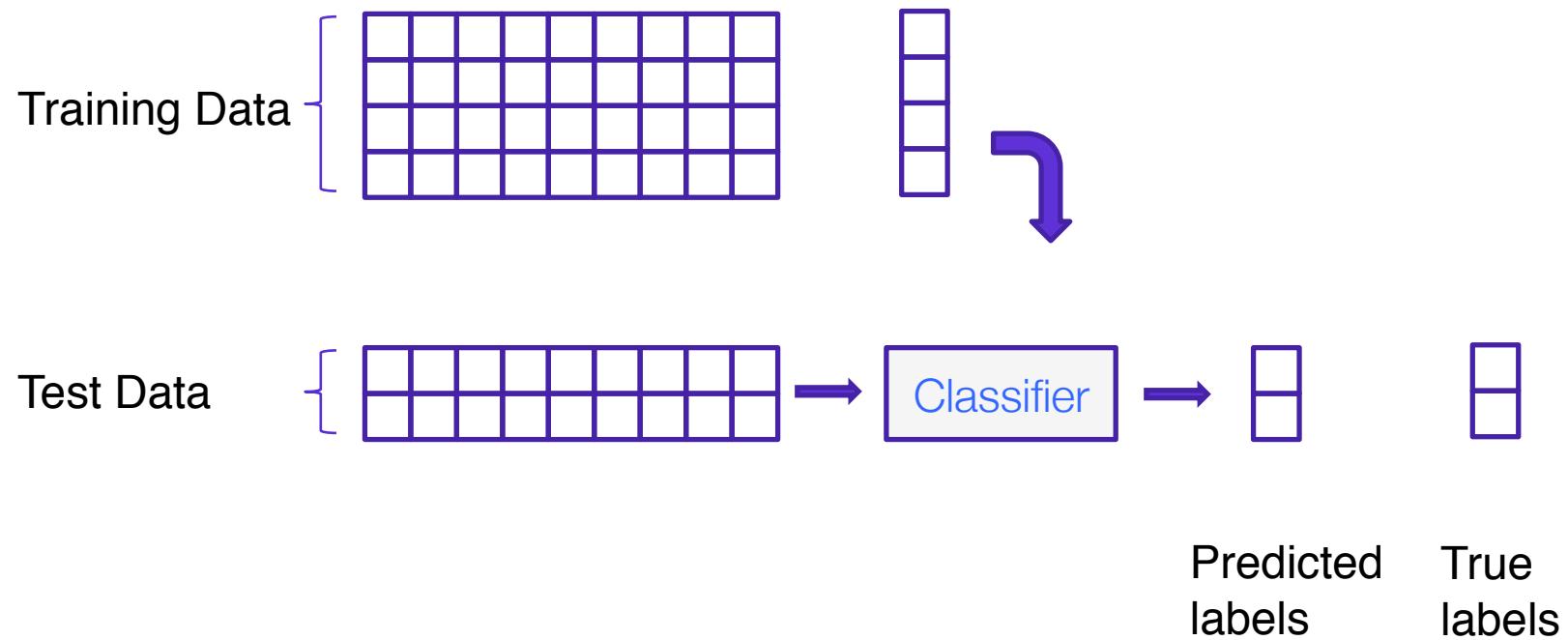
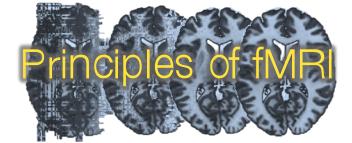
Illustration



The full data set is split into two parts: training and test data



Illustration



End of Module



@fMRIstats

Criteria for a good prediction algorithm

- **Accuracy:** How accurate are predictions, and are they good enough to be useful?
- **Specificity:** How well can a pattern of brain activity discriminate among alternative outcomes or processes?
- **Generalizability:** Does a pattern predict other, dissimilar variants of the outcome of interest?
- **Interpretability:** Are the brain maps neuroscientifically interpretable?
- **Transparency:** Can scientists understand the algorithm and its results, and thus know why it might succeed or fail?

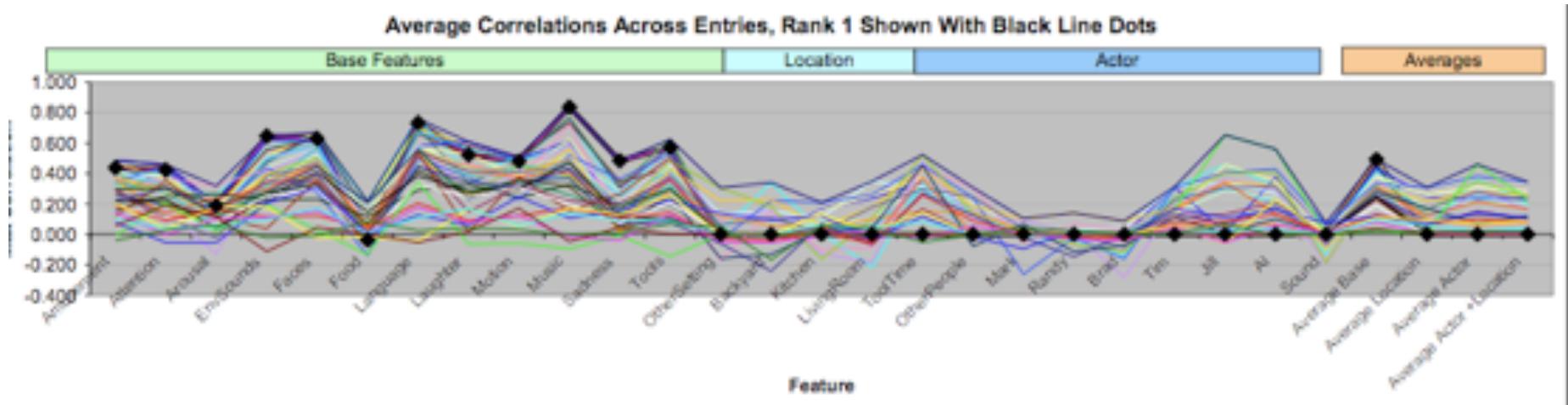
Interpretability: Example



Olivetti et al., 2006

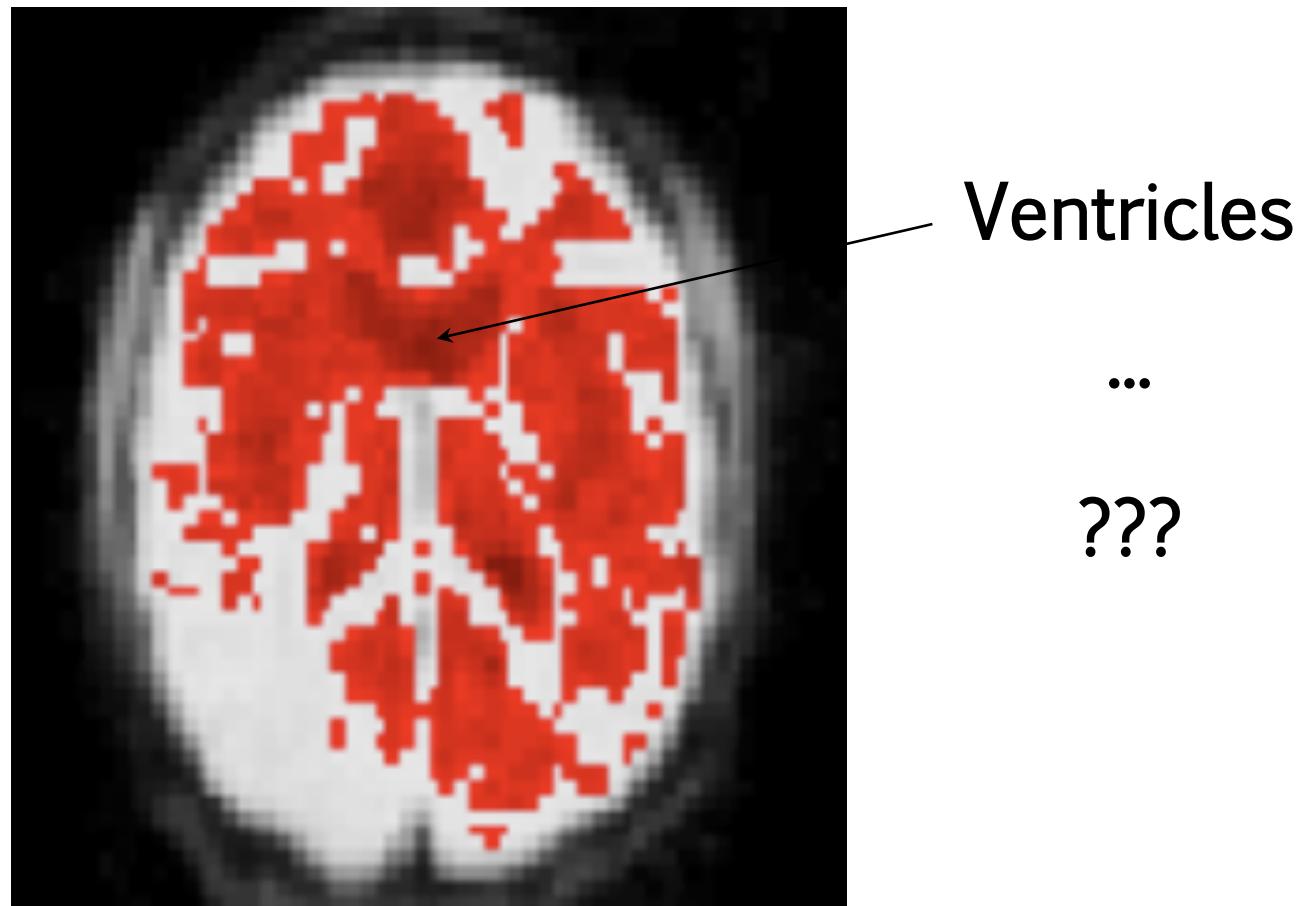
Winner: Pittsburgh Brain Activity Interpretation
Competition

Interpretability: Example



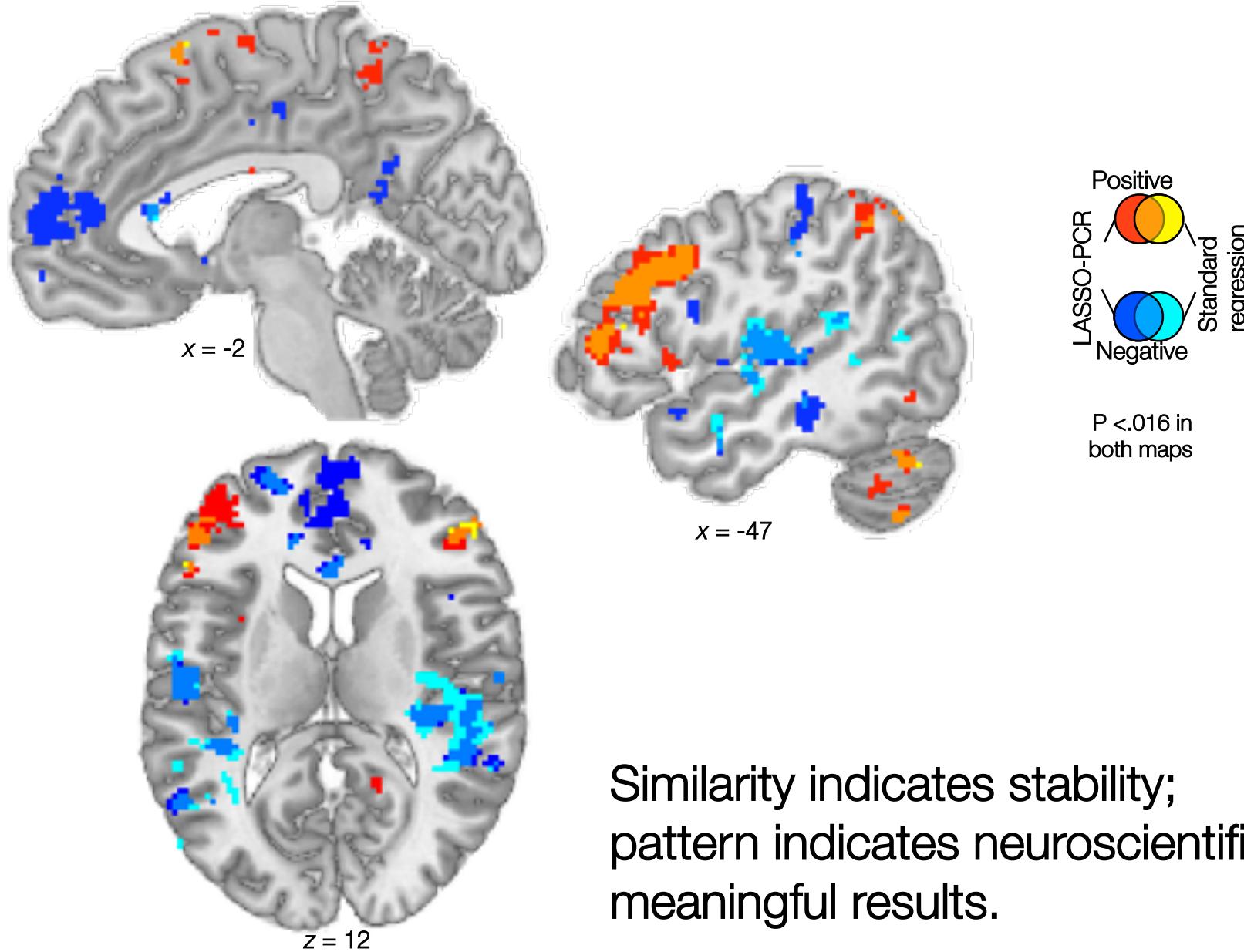
Olivetti et al., 2006

Interpretability: Example



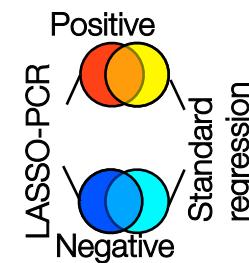
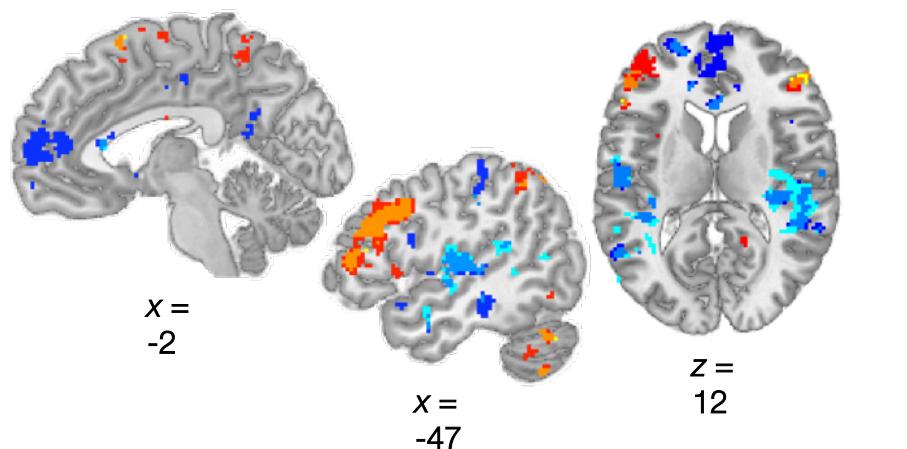
Language, Subject 2

Interpretability: Comparison between LASSO-PCR and standard regression weights



Similarity indicates stability;
pattern indicates neuroscientifically
meaningful results.

Comparison between LASSO-PCR and standard regression weights



$P < .016$ in both maps

