# Modeling Group fMRI Data

#### Overview

- What is a mixed effects model
  - Fixed effects
  - Random effects
- 2-stage summary statistics approach
- How do different software packages work?
- Overview FSL modeling options

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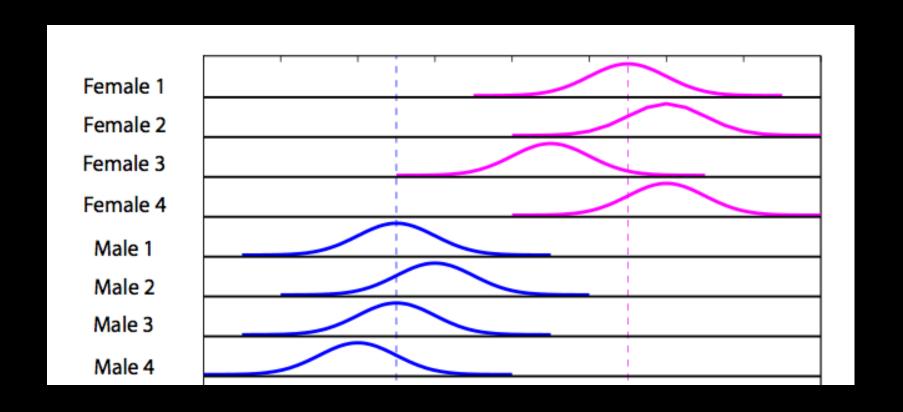
### Mixed Model Motivation

- Start with a simple ANOVA example
- Study: Is hair length different between males and females?

# Start: 1 hair per person

- Two sources of variability
  - Variance of hair length within person
  - Variance of hair length between people

Assume within-subject variance is 1

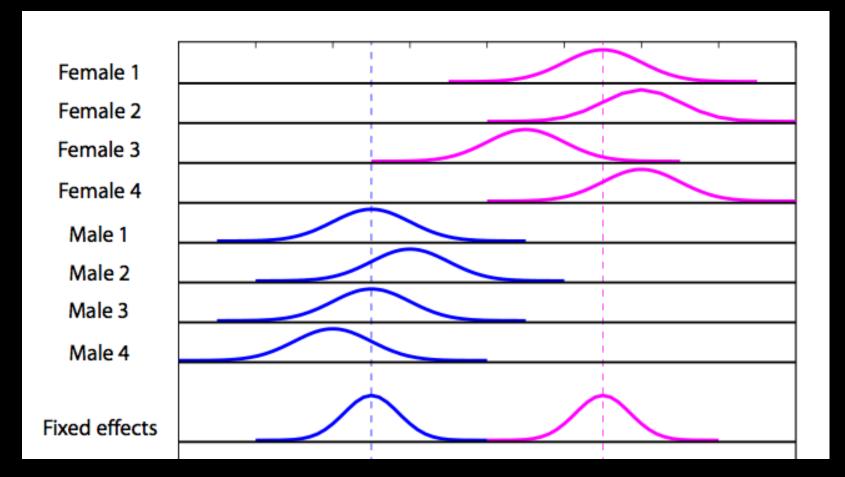


Each distribution has a variance of 1

# Fixed effects analysis

 We're only interested in these exact 4 men and 4 women

$$\sigma_{\scriptscriptstyle ext{FFX}}^2=rac{1}{4}\sigma_{\scriptscriptstyle ext{W}}^2=0.25$$

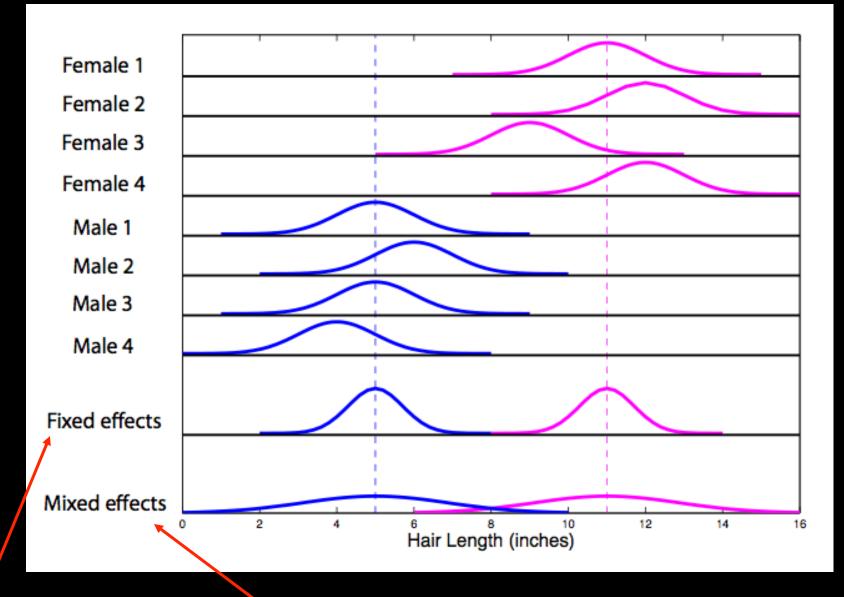


$$\sigma_{\scriptscriptstyle ext{FFX}}^2 = rac{1}{4}\sigma_{\scriptscriptstyle ext{W}}^2 = 0.25$$

#### Mixed effects

- Include both within and between subject variances
- Adding a between subject means subject is random
  - Anything with a variance is random!

$$\sigma_{ ext{MFX}}^2 = \sigma_{ ext{W}}^2/4 + \sigma_{ ext{B}}^2/4 = 1/4 + 49/4 = 12.5$$



$$\sigma_{\scriptscriptstyle \mathrm{FFX}}^2 = \frac{1}{4} \sigma_{\scriptscriptstyle \mathrm{W}}^2 = 0.25$$

$$\sigma_{ ext{MFX}}^2 = \sigma_{ ext{W}}^2/4 + \sigma_{ ext{B}}^2/4 = 1/4 + 49/4 = 12.5$$

# Multiple hairs per subject

Fixed effects variance

$$\sigma_{\text{FFX}}^2 = \frac{1}{4}\sigma_{\text{W}}^2/25 = 0.01$$

Mixed effects variance

$$\sigma_{\text{MFX}}^2 = \frac{1}{4}\sigma_{\text{W}}^2/25 + \frac{1}{4}\sigma_{\text{B}}^2 = 12.26$$

# Multiple hairs per subject

Fixed effects variance

$$\sigma_{\text{FFX}}^2 = \frac{1}{4}\sigma_{\text{W}}^2/25 = 0.01$$

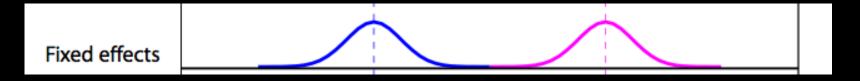
Mixed effects variance

$$\sigma_{\text{MFX}}^2 = \frac{1}{4}\sigma_{\text{W}}^2/25 + \frac{1}{4}\sigma_{\text{B}}^2 = 12.26$$

there is an innate tradeoff between collecting more data per subject, Between subject variance and collecting more subjects. typically dominates

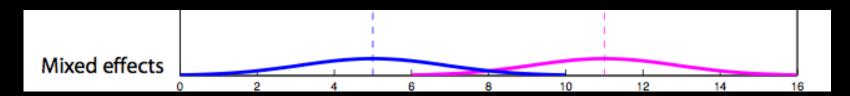
# Wrong model leads to wrong conclusion

- Scenario 1: Fixed effects model
  - Significant difference in hair length
  - Result only applies to these 8 subjects



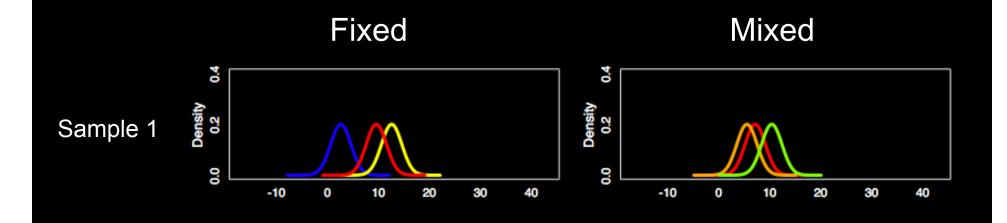
# Wrong model leads to wrong conclusion

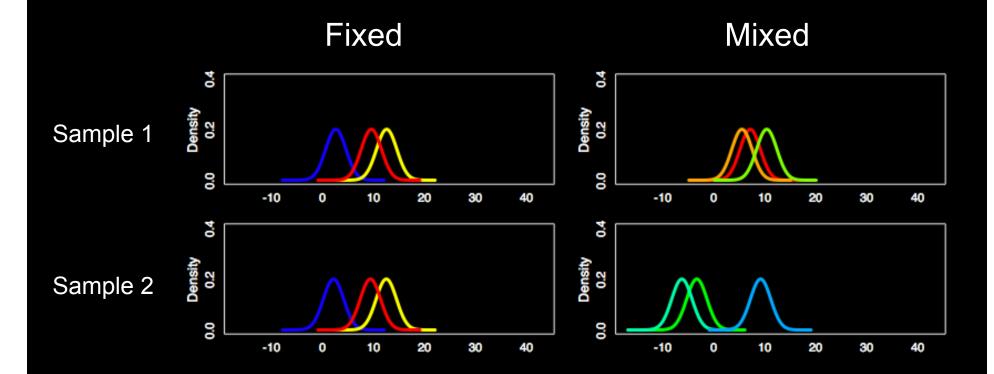
- Scenario 2: Mixed effects model
  - Cannot conclude there is a difference in hair length

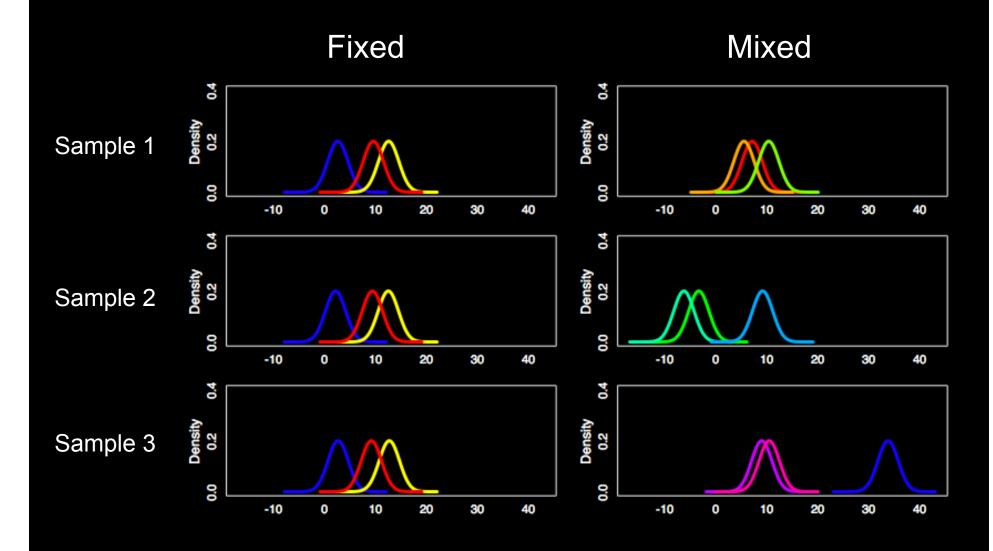


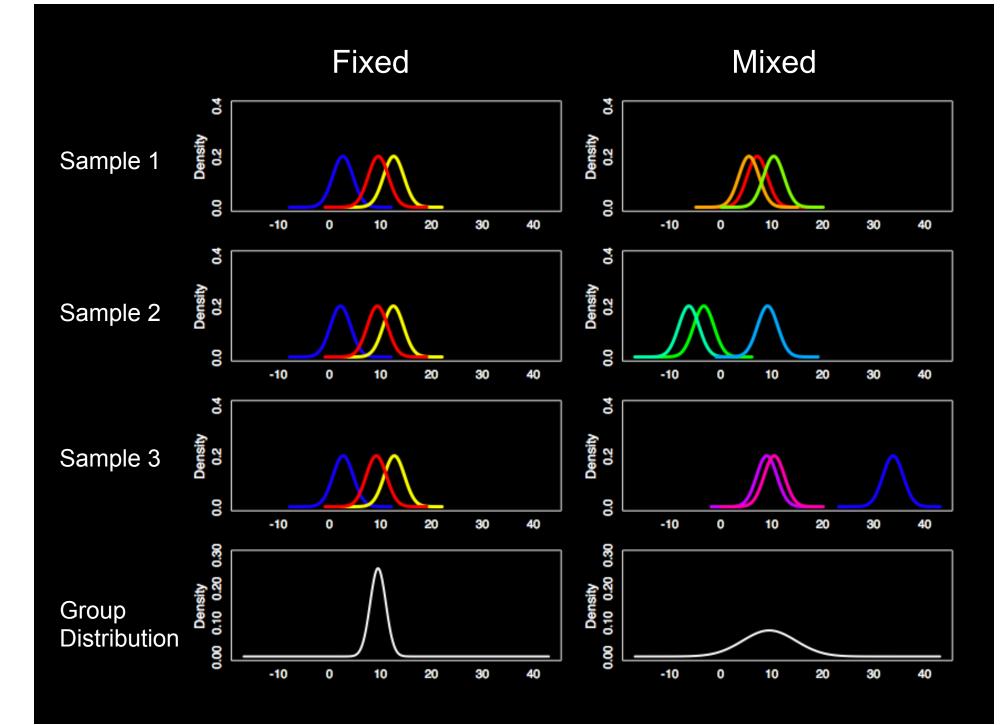
#### Mixed Model Comments

- If you fail to include a random effect when there is one
  - Results only apply to that data sample
  - P-values are smaller than mixed model pvalues









# Important points so far...

- Ignoring random subject effect means that you're ignoring the fact that these subjects were randomly sampled
  - Inference only applies to sample you collected
- Including random subject effect always increases your variance

## Important points so far...

- What has a bigger impact in reducing variance?
  - Adding more hairs per subject?
  - Adding more subjects?

# Important points so far...

- What has a bigger impact in reducing variance?
  - Adding more hairs per subject?
  - Adding more subjects?

1 hair per subject

$$\sigma_{ ext{MFX}}^2 = \sigma_{ ext{W}}^2/4 + \sigma_{ ext{B}}^2/4 = 12.5$$

25 hairs per subject

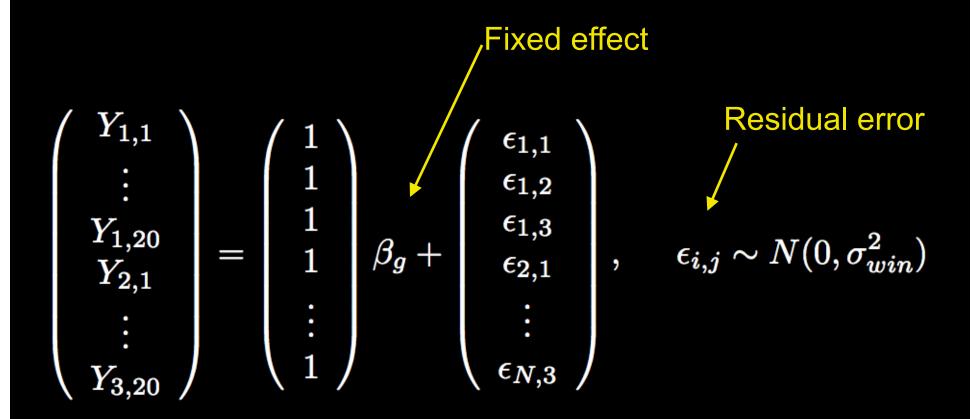
$$\sigma_{\text{MFX}}^2 = \sigma_{\text{W}}^2/4 + \sigma_{\text{B}}^2/4 = 12.5$$
  $\sigma_{\text{MFX}}^2 = \frac{1}{4}\sigma_{\text{W}}^2/25 + \frac{1}{4}\sigma_{\text{B}}^2 = 12.26$ 

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### Fixed effects model:

modeling the mean of 3 females, 20 hairs



#### Mixed Effects Model

Stage 1 
$$Y=Xeta$$

$$\epsilon$$

$$\begin{pmatrix} Y_{1,1} \\ \vdots \\ Y_{1,20} \\ Y_{2,1} \\ \vdots \\ Y_{3,20} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} + \begin{pmatrix} \epsilon_{1,1} \\ \epsilon_{1,2} \\ \epsilon_{1,3} \\ \epsilon_{2,1} \\ \vdots \\ \epsilon_{N,3} \end{pmatrix}, \epsilon_{i,j} \sim N(0, \sigma_{win}^2)$$

$$\left.egin{array}{ccc} \epsilon_{1,3} & & & \\ \epsilon_{2,1} & & , \epsilon_{i,j} \sim N(0,\sigma_{win}^2) \end{array}
ight.$$

$$eta = X_g eta_g + \eta$$
 Random effect

$$\eta$$
 Random effect

$$\left(egin{array}{c} eta_1 \ eta_2 \ eta_3 \end{array}
ight) = \left(egin{array}{c} 1 \ 1 \ 1 \end{array}
ight)eta_g + \left(egin{array}{c} \eta_1 \ \eta_2 \ \eta_3 \end{array}
ight), \quad \eta_i \sim N(0,\sigma_{btwn}^2)$$

#### Mixed Effects Model

Stage 1 
$$Y=Xeta+\epsilon$$
 
$$\begin{pmatrix} Y_{1,1} \\ \vdots \\ Y_{1,20} \\ Y_{2,1} \\ \vdots \\ Y_{3,20} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} + \begin{pmatrix} \epsilon_{1,1} \\ \epsilon_{1,2} \\ \epsilon_{1,3} \\ \epsilon_{2,1} \\ \vdots \\ \epsilon_{N,3} \end{pmatrix}, \epsilon_{i,j} \sim N(0, \sigma_{win}^2)$$
 Stage 2  $\beta=X_g\beta_g+\eta$  Random effect 
$$\begin{pmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} = \begin{pmatrix} 1 \\ 1 \\ 1 \end{pmatrix} \beta_g + \begin{pmatrix} \eta_1 \\ \eta_2 \\ \eta_3 \end{pmatrix}, \quad \eta_i \sim N(0, \sigma_{btwn}^2)$$

# Mixed Effects Model: All-In-One

$$Y = XX_{g}\beta_{g} + X\eta + \epsilon$$

$$\begin{pmatrix} Y_{1,1} \\ \vdots \\ Y_{1,20} \\ Y_{2,1} \\ \vdots \\ Y_{3,20} \end{pmatrix} = \begin{pmatrix} 1 \\ 1 \\ 1 \\ 1 \\ \vdots \\ 1 \end{pmatrix} \beta_{g} + \begin{pmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} \eta_{1} \\ \eta_{2} \\ \eta_{3} \end{pmatrix} + \begin{pmatrix} \epsilon_{1,1} \\ \epsilon_{1,2} \\ \epsilon_{1,3} \\ \epsilon_{2,1} \\ \vdots \\ \epsilon_{N,3} \end{pmatrix}$$

$$Variance Terms$$

#### How does this relate to fMRI?

Subject 1

Subject 2

Subject N

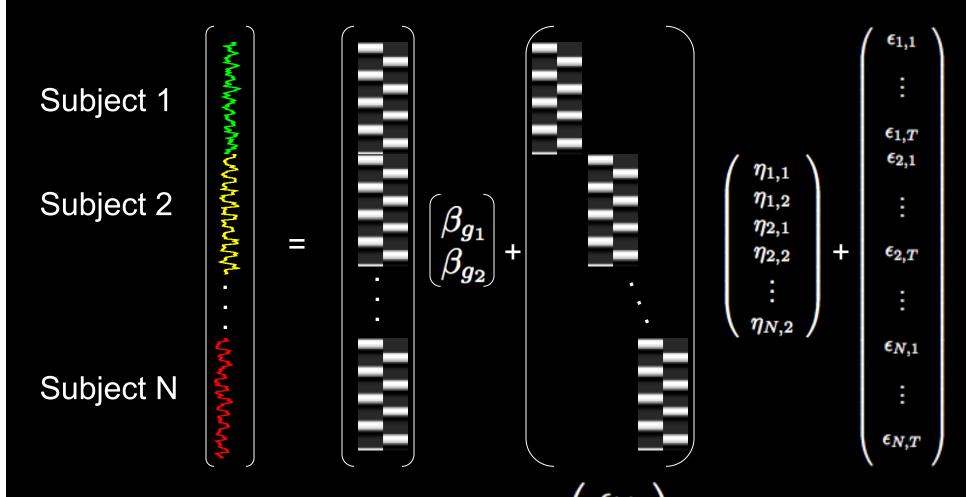
Subject N

Each time series is a collection of data grouped by subject

A random subject effect is necessary to apply inference to total population

#### Mixed Model for fMRI Data

- fMRI data are more complicated than the hair length example
  - Not typically estimating an intercept
  - Time series are temporally autocorrelated
  - Time series can be quite long
- Let's take a look at the model!
  - A study with 2 stimuli of interest



$$\operatorname{Var}(\eta_{i,1}) = \sigma_{btwn_1}^2 \qquad \operatorname{Cov} \left( egin{array}{c} \epsilon_{i,1} \\ \epsilon_{i,2} \\ \vdots \\ \epsilon_{i,T} \end{array} 
ight) = \sigma_{win_i}^2 V_i$$

#### Yuck!

- Computationally intensive
  - Large matrices that need to be inverted
- What if we add another subject?
  - Must estimate whole model for all subjects

### Recall the two stages

Stage 1 
$$Y=X\beta+\epsilon$$
 
$$\begin{pmatrix} Y_{1,1} \\ \vdots \\ Y_{1,20} \\ Y_{2,1} \\ \vdots \\ Y_{3,20} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \end{pmatrix} \begin{pmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} + \begin{pmatrix} \epsilon_{1,1} \\ \epsilon_{1,2} \\ \epsilon_{1,3} \\ \epsilon_{2,1} \\ \vdots \\ \epsilon_{N,3} \end{pmatrix}, \epsilon_{i,j} \sim N(0, \sigma_{win}^2)$$
 Stage 2  $\beta=X_g\beta_g+\eta$  
$$\begin{pmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \end{pmatrix} = \begin{pmatrix} 1 \\ 1 \\ \beta_2 \\ \beta_3 \end{pmatrix} = \begin{pmatrix} 1 \\ 1 \\ 1 \end{pmatrix} \beta_g + \begin{pmatrix} \eta_1 \\ \eta_2 \\ \eta_3 \end{pmatrix}, \quad \eta_i \sim N(0, \sigma_{btwn}^2)$$

# Two-Stage Summary Statistics

Stage 1

$$\begin{pmatrix} Y_{1,1} \\ \vdots \\ Y_{1,20} \end{pmatrix} = \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} \beta_1 + \begin{pmatrix} \epsilon_{1,1} \\ \vdots \\ \epsilon_{1,20} \end{pmatrix}$$

$$\begin{pmatrix} Y_{2,1} \\ \vdots \\ Y_{2,20} \end{pmatrix} = \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} \beta_2 + \begin{pmatrix} \epsilon_{2,1} \\ \vdots \\ \epsilon_{2,20} \end{pmatrix} \qquad \epsilon_{i,j} \sim N(0, \sigma_{win}^2)$$

$$\begin{pmatrix} Y_{3,1} \\ \vdots \\ Y_{3,20} \end{pmatrix} = \begin{pmatrix} 1 \\ \vdots \\ 1 \end{pmatrix} \beta_3 + \begin{pmatrix} \epsilon_{3,1} \\ \vdots \\ \epsilon_{3,20} \end{pmatrix}$$

breaking apart the fixed-effects model for 3 participants into 3 separate regressions

# Two-Stage Summary Statistics

Stage 1

$$\left(egin{array}{c} Y_{1,1} \\ dots \\ Y_{1,20} \end{array}
ight) = \left(egin{array}{c} 1 \\ dots \\ 1 \end{array}
ight) eta_1 + \left(egin{array}{c} \epsilon_{1,1} \\ dots \\ \epsilon_{1,20} \end{array}
ight) \ \left(egin{array}{c} Y_{2,1} \\ dots \\ Y_{2,20} \end{array}
ight) = \left(egin{array}{c} 1 \\ dots \\ 1 \end{array}
ight) eta_2 + \left(egin{array}{c} \epsilon_{2,1} \\ dots \\ \epsilon_{2,20} \end{array}
ight) \ \left(egin{array}{c} \epsilon_{i,j} \sim N(0, \sigma_{win}^2) \\ \varepsilon_{i,j} \sim N(0, \sigma_{win}^2) \\ \varepsilon_{i,j} \sim N(0, \sigma_{win}^2) \\ \varepsilon_{i,j} \sim N(0, \sigma_{win}^2) \end{array}
ight) \ \left(egin{array}{c} Y_{3,1} \\ dots \\ Y_{3,20} \end{array}
ight) = \left(egin{array}{c} 1 \\ dots \\ 1 \end{array}
ight) eta_3 + \left(egin{array}{c} \epsilon_{3,1} \\ dots \\ \epsilon_{3,20} \end{array}
ight) \ \left(egin{array}{c} \epsilon_{1,1} \\ dots \\ \varepsilon_{3,20} \end{array}
ight) \ \left(egin{array}{c} \epsilon_{2,1} \\ dots \\ \varepsilon_{3,20} \end{array}
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ight) \ \left(egin{array}{c} \epsilon_{3,1} \\ dots \\ \varepsilon_{3,20} \end{array}
ight) \ \left(egin{array}{c} \epsilon_{3,20} \\ \varepsilon_{3,20} \end{array}
ight) \ \left(egin{array}{c} \epsilon_{3,1} \\ dots \\ \varepsilon_{3,20} \end{array}
ight) \ \left(egin{array}{c} \epsilon_{3,20} \\ \varepsilon_{3,20} \\$$

Stage 2

Use first stage estimates

$$\left( \left( \begin{array}{c} \hat{\beta}_1 \\ \hat{\beta}_2 \\ \hat{\beta}_3 \end{array} \right) = \left( \begin{array}{c} 1 \\ 1 \\ 1 \end{array} \right) \beta_g + \left( \begin{array}{c} \eta_1^* \\ \eta_2^* \\ \eta_3^* \end{array} \right), \quad \operatorname{Var}(\eta_i^*) = \frac{\sigma_{win}^2}{W} + \sigma_{btwn}^2$$

# Two-Stage Summary Statistics

Stage 1

$$\left( egin{array}{c} Y_{1,1} \ dots \ Y_{1,20} \end{array} 
ight) = \left( egin{array}{c} 1 \ dots \ 1 \end{array} 
ight) eta_1 + \left( egin{array}{c} \epsilon_{1,1} \ dots \ \epsilon_{1,20} \end{array} 
ight)$$

$$\left(\begin{array}{c} Y_{2,1} \\ \vdots \\ Y_{2,20} \end{array}\right) = \left(\begin{array}{c} 1 \\ \vdots \\ 1 \end{array}\right) \beta_2 + \left(\begin{array}{c} \epsilon_{2,1} \\ \vdots \\ \epsilon_{2,20} \end{array}\right) \qquad \epsilon_{i,j} \sim N(0, \sigma_{win}^2)$$

$$\epsilon_{i,j} \sim N(0, \sigma_{win}^2)$$

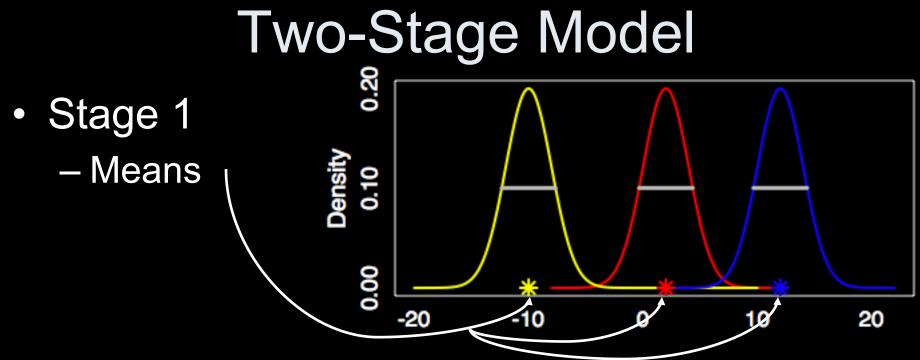
within

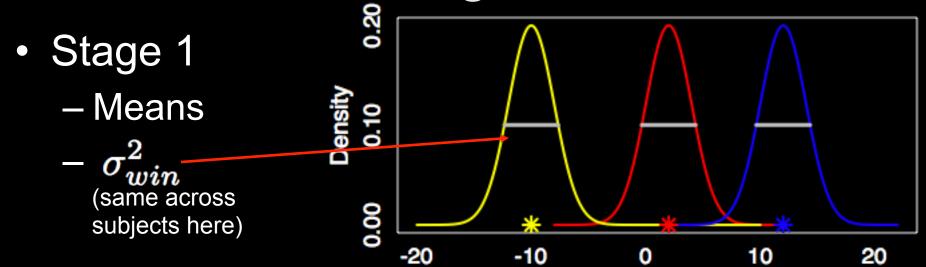
between

$$\left(\begin{array}{c} Y_{3,1} \\ \vdots \\ Y_{3,20} \end{array}\right) = \left(\begin{array}{c} 1 \\ \vdots \\ 1 \end{array}\right) \beta_3 + \left(\begin{array}{c} \epsilon_{3,1} \\ \vdots \\ \epsilon_{3,20} \end{array}\right)$$

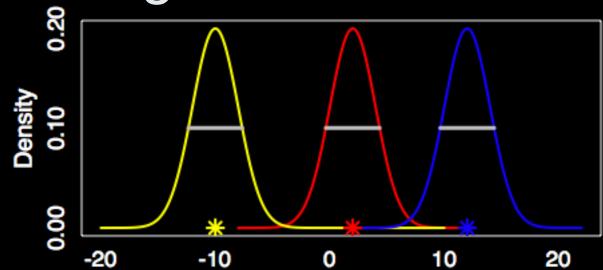
Stage 2

$$\left(egin{array}{c} \hat{eta}_1 \ \hat{eta}_2 \ \hat{eta}_3 \end{array}
ight) = \left(egin{array}{c} 1 \ 1 \ 1 \end{array}
ight)eta_g + \left(egin{array}{c} \eta_1^* \ \eta_2^* \ \eta_3^* \end{array}
ight), \quad ext{Var}(\eta_i^*) = \left(egin{array}{c} \sigma_{win}^2 \ W \end{array}
ight) + \left(\sigma_{btwn}^2 \ \eta_3^* \end{array}
ight)$$





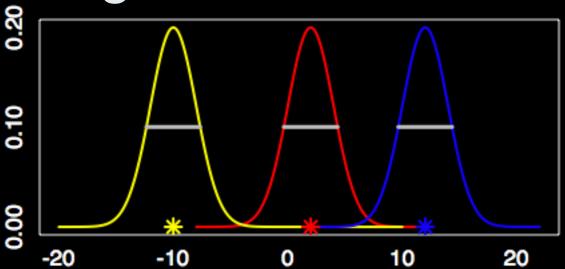
- Stage 1
  - Means
  - σ<sup>2</sup><sub>win</sub>
     (same across subjects here)



- Stage 2
  - $=\sigma_{btwn}^2$



- Stage 1
  - Means
  - $-\sigma_{win}^2$  (same across subjects here)

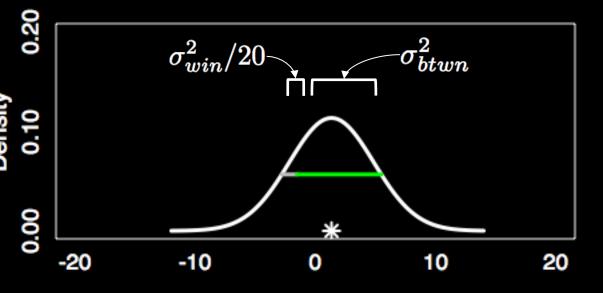


Stage 2

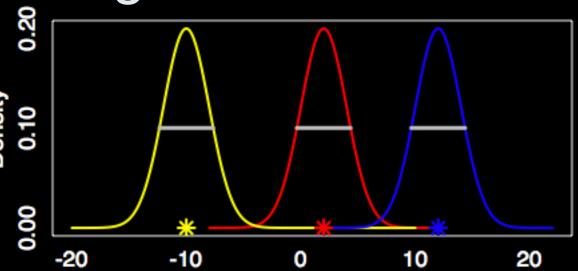
$$=\sigma_{btwn}^2$$

$$\sigma_{mix}^2 = rac{\sigma_{win}^2}{20} + \sigma_{btwn}^2$$

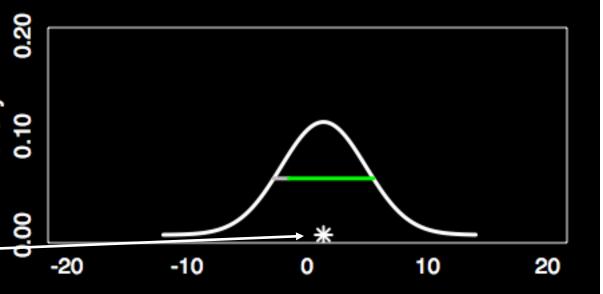
• 20 hairs/subject



- Stage 1
  - Means
  - $\sigma_{win}^2$  (same across subjects here)



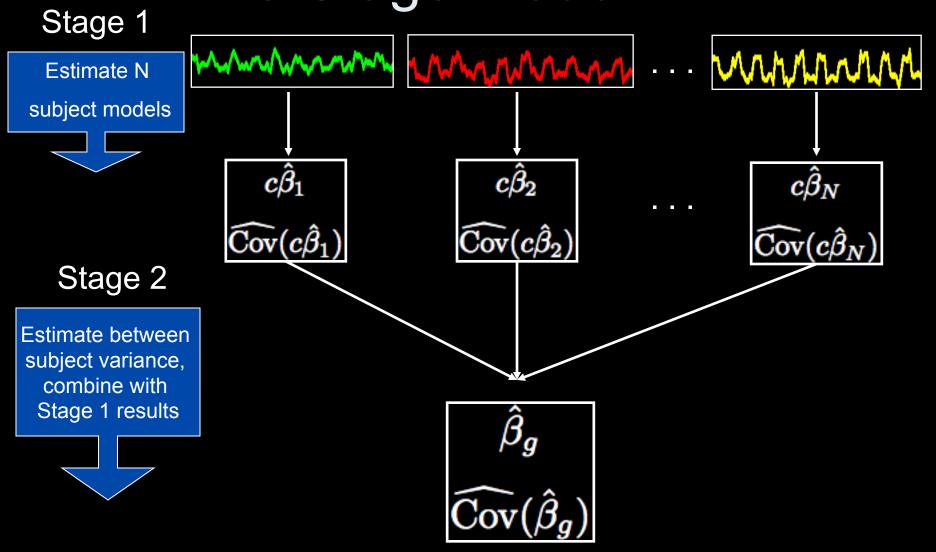
- Stage 2
  - $-\sigma_{btwn}^2$
  - $\begin{array}{l} \quad \quad \sigma_{mix}^2 = \\ \quad \frac{\sigma_{win}^2}{20} + \sigma_{btwn}^2 \end{array}$ 
    - 20 hairs/subject
- Pop mean



• 
$$T = rac{\sqrt{N}\hat{eta}}{\sqrt{\sigma_{win}^2/W + \sigma_{btwn}^2}}$$

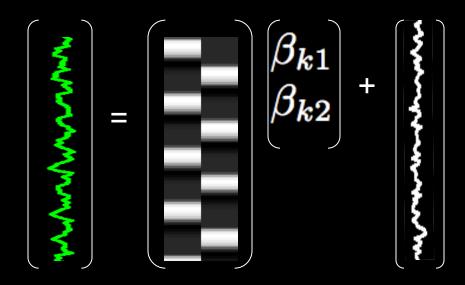
- -N = # subjects
- W = # measures within subject
- If new data are added, only run first stage for new data

## Two Stage Model fMRI



## Stage 1: Subject Model

$$Y_k = X_k \beta_k + \epsilon_k$$



$$Cov(\epsilon_k) = \sigma_k^2 V_k$$
$$H_0: \beta_{k1} - \beta_{k2} = 0$$

•  $W_k$  such that  $W_k V_k W_k' = I_T$ 

- $W_k$  such that  $W_k V_k W_k' = I_T$
- Whitened model

$$-W_k Y_k = W_k X_k \beta_k + W_k \epsilon_k$$

$$-Y_k^* = X_k^* \beta_k + \epsilon_k^*$$

- $W_k$  such that  $W_k V_k W_k' = I_T$
- Whitened model

$$-W_k Y_k = W_k X_k \beta_k + W_k \epsilon_k -Y_k^* = X_k^* \beta_k + \epsilon_k^*$$

Use OLS on whitened model

$$egin{aligned} &-c\hat{eta}_k = \left(X_k^{*'}X_k^*
ight)^{-1}X_k^{*'}Y_k^* \ &-\widehat{Cov}(c\hat{eta}_k) = \hat{\sigma}_k^2\left(X_k^{*'}X_k^*
ight)^{-1} \end{aligned}$$

## Stage 2: Group Model

$$\hat{eta}_{cont} = X_g eta_g + \epsilon_g$$
 $c\hat{eta}_1$ 
 $= \begin{bmatrix} 1 \\ 1 \\ 1 \\ 1 \\ 1 \\ 1 \\ 1 \end{bmatrix}$ 
 $eta_g$  +

$$egin{aligned} \operatorname{Cov}(\epsilon_g) &= V_g = \ & \left( egin{array}{cccc} \sigma_1^2 c(X_1^{*'}X_1^*)^{-1}c' & & & & \\ & \ddots & & & & \\ & & \sigma_N^2 c(X_N^{*'}X_N^*)^{-1}c' & & \\ & & & & & \wedge & \\ & & & & \wedge & \\ \end{aligned} 
ight) + \sigma_g^2 I_N \ & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & \\ & & & \\ & & \\ & & & \\ & &$$

•  $W_g$  such that  $W_g V_g W_g' = I_N$ 

- $W_g$  such that  $W_g V_g W_g' = I_N$
- $W_g \hat{\beta}_{cont} = W_g X_g \beta_g + W_g \epsilon_g$  $\hat{\beta}_{cont}^* = X_g^* \beta_g + \epsilon_g^*$

- $W_g$  such that  $W_g V_g W'_g = I_N$
- $W_g \hat{\beta}_{cont} = W_g X_g \beta_g + W_g \epsilon_g$  $\hat{\beta}_{cont}^* = X_g^* \beta_g + \epsilon_g^*$
- $\hat{\beta}_g = \left(X_g^{*'} X_g^*\right)^{-1} X_g^{*'} \hat{\beta}_{cont}^*$   $\widehat{Cov}(\hat{\beta}_g) = \left(X_g^{*'} X_g^*\right)^{-1}$

- $W_g$  such that  $W_g V_g W'_g = I_N$
- $W_g \hat{\beta}_{cont} = W_g X_g \beta_g + W_g \epsilon_g$  $\hat{\beta}_{cont}^* = X_g^* \beta_g + \epsilon_g^*$

$$\hat{\beta}_g = \left(X_g^{*'} X_g^*\right)^{-1} X_g^{*'} \hat{\beta}_{cont}^*$$

$$\widehat{Cov}(\hat{\beta}_g) = \left(X_g^{*'} X_g^*\right)^{-1}$$

• 
$$T = \hat{\beta}_g / \sqrt{\widehat{\mathrm{Cov}}(\hat{\beta}_g)}$$

Using a similar process to pre-whitening, multiplying by a matrix such that subject variances are equivalent. This is a weighted regression that will allow the assumptins of Gauss-Markov to hold.

### Overview so far...

- The mixed model can be thought of as a two stage process
  - Stats software estimates both stages simultaneously
  - In fMRI it is computationally easier and more convenient to keep the stages separate
    - Easier to add new subjects
    - Not exactly the same as the stats software results

### Overview so far

 When the model is estimated how is a subject with a high mfx variance treated differently than a subject with a low mfx variance?

### Overview so far

- When the model is estimated how is a subject with a high mfx variance treated differently than a subject with a low mfx variance?
  - Subjects with higher variability are downweighted in the analysis!

### Overview

- What is a mixed effects model
  - Fixed effects
  - Random effects
- 2-stage summary statistics approach
- How do different software packages work?
- Overview FSL modeling options

### How is the model estimated?

- Depends on software
  - -SPM: Does not estimate  $\sigma_g^2$ 
    - Due to a set of assumptions, estimation of is unnecessary
  - FSL: Bayesian approach to estimating  $\sigma_g^2$

### SPM2

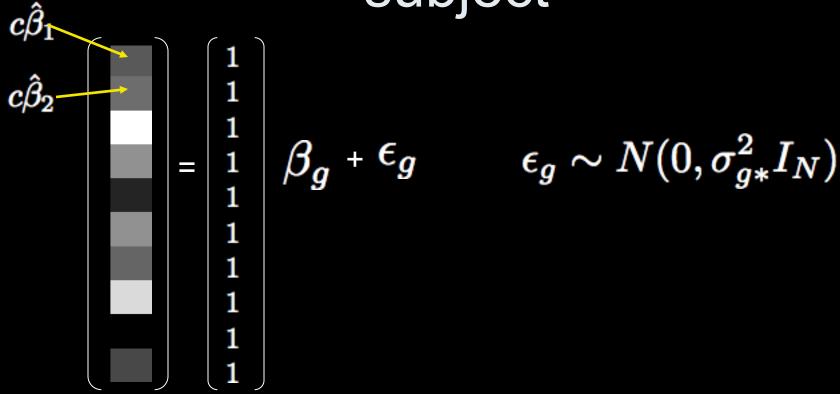
- Does not estimate  $\sigma_g^2$ 
  - Assumes homogeneous variance across subjects
  - Assumes first level design is same across subjects

$$\hat{\sigma}_{win_{all}}^2 = \hat{\sigma}_1^2 c \left( X_1^{*'} X_1^* \right)^{-1} c' = \ldots = \hat{\sigma}_N^2 c \left( X_N^{*'} X_N^* \right)^{-1} c'$$

$$V_g = \sigma_{win_{all}}^2 I_N + \sigma_g^2 I_N = \sigma_{g*}^2 I_N$$

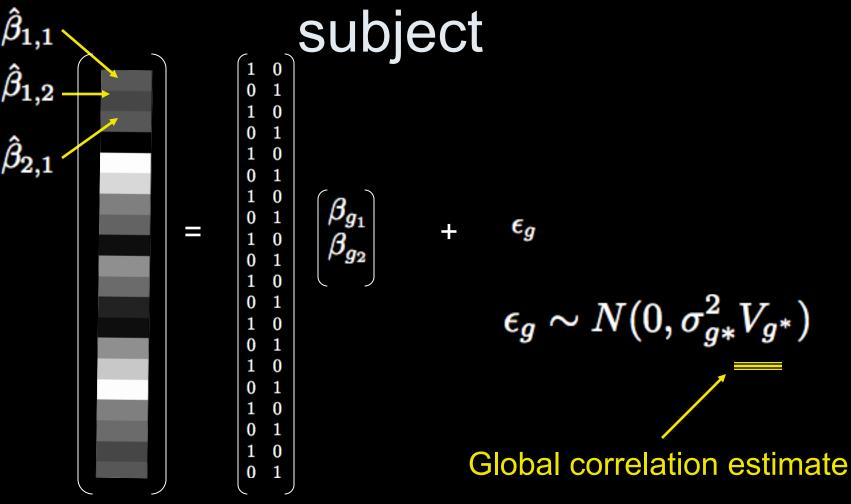
OLS can be used

# SPM2 : Single contrast per subject



A one-sample T-test!

## SPM2 : Multiple contrasts per



### SPM2: Summary

- Multiple contrasts per subject can enter second level
  - Contrasts can be correlated
  - T and F-tests are possible
- Special case
  - One contrast per subject…Reduces to Ttest!

### SPM2

### Pros

This is a good model for ROI analyses

- Model is easy to estimate
- Model is easy to understand
- Multiple contrasts can enter the group model and are not considered independent

### Cons

- Global covariance estimate (same across voxels)
- Assumes variance is homogeneous across subjects

Type I error rates remain stable, but power will take a hit if the data contain high levels of heteroskedasicity

## **FSL: FMRIB Software Library**

- Bayesian approach to estimating model
- Inference is based on posterior distribution of the data
  - $-P(\beta_g,\sigma_g^2,\nu_g|Y)$
  - Parameters of interest are treated as random

### FSL: Second Level Estimation

- Flame 1: Maximum a posteriori (MAP) estimate of  $\sigma_g^2$  found iteratively
  - Assumes degrees of freedom,  $\nu_g=N-p$
- Flame 2: Slower MCMC method of estimation
  - Applied to voxels close to threshold in step 1
  - Fine tunes estimates of  $eta_g, \sigma_g^2, 
    u_g$
- Details
  - Woolrich et al. NI (2004) 1732-47

### FSL

### Pros

- When single contrast is taken to the second level, equivalent to all-in-one model
- Within-subject variances are carried to the second level
  - Heterogeneity across subjects is modeled

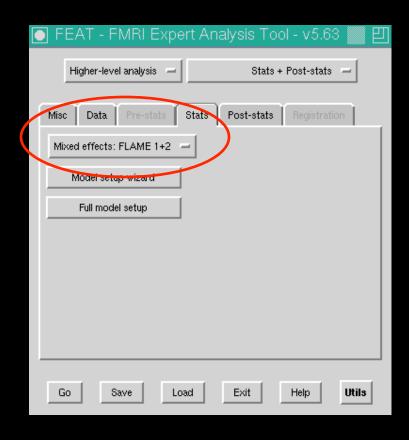
### Cons

 Multiple contrasts in the group model are assumed to be independent

### Which software?

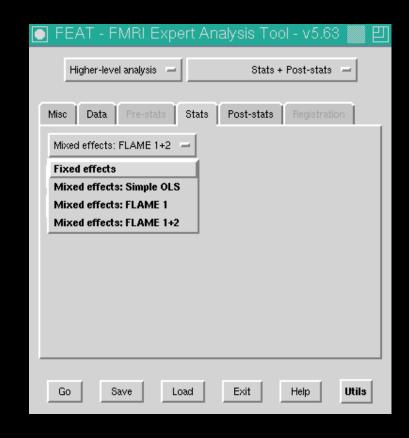
- FSL and FMRIstat best for heteroscedastic variances
  - Different number of trials per subject
- SPM best for multiple correlated contrasts at group level
- Other differences in first level modeling may sway users one way or another

## FSL Group Model Options



## FSL Group Model Options

- Fixed effects
  - Only uses w/in sub variance
- Simple OLS
  - Assumes w/in sub variances are equal
- Flame 1& 2
  - w/in sub var and btwn sub var



## W<sub>g</sub> Matrix

 Recall we pre-multiply by W<sub>g</sub> so our errors are uncorrelated and constant variance

$$W_g \hat{eta}_{cont} = W_g X_g \beta_g + W_g \epsilon_g$$

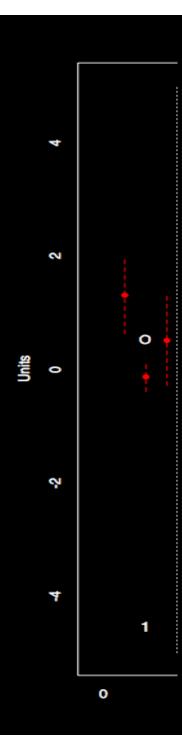
Act as

$$V_g = \left( egin{array}{ccc} \sigma_{win_1}^2 + \sigma_g^2 & & 0 \ & \ddots & & 0 \ & & \sigma_{win_N}^2 + \sigma_g^2 \end{array} 
ight) 
ightarrow W_g = \left( egin{array}{ccc} rac{1}{\sqrt{\sigma_{win_1}^2 + \sigma_g^2}} & & 0 \ & & \ddots & & 0 \ & & & rac{1}{\sqrt{\sigma_{win_N}^2 + \sigma_g^2}} \end{array} 
ight)$$

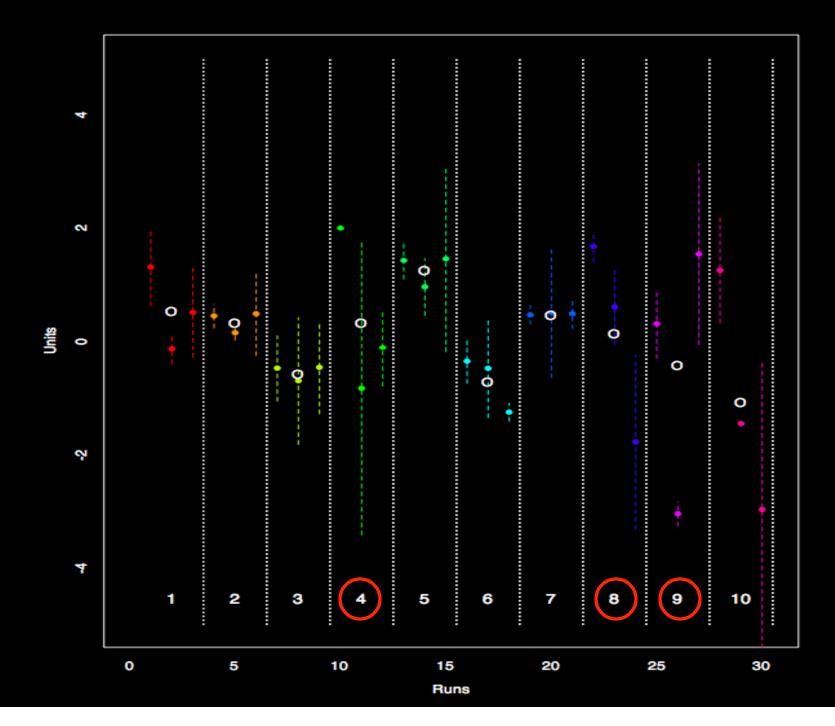
## V<sub>g</sub> Matrix Assumptions

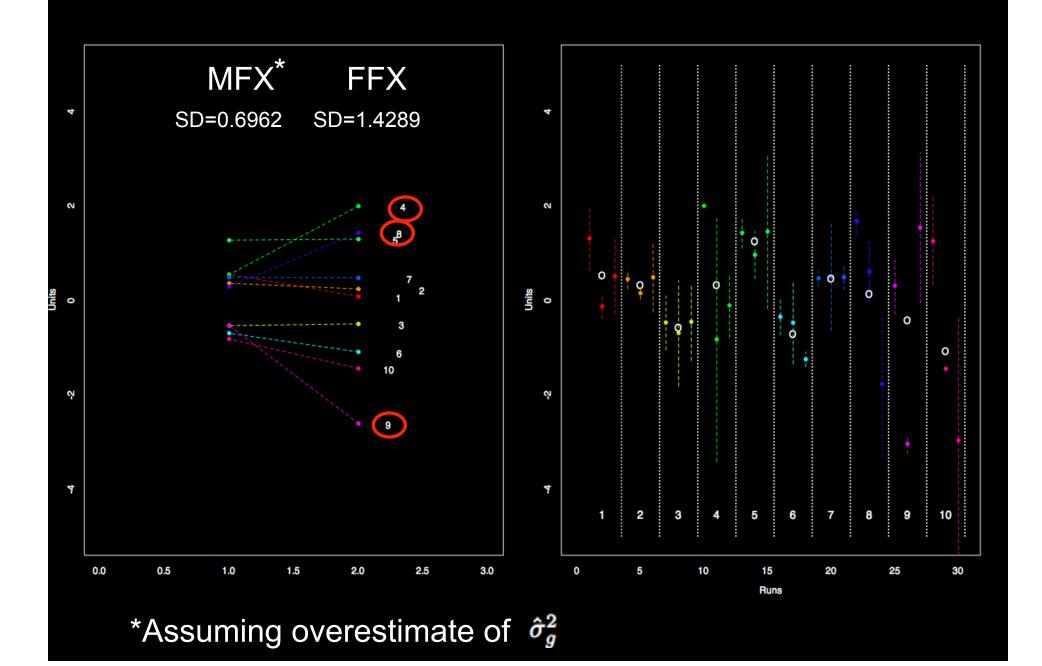
- Fixed effects analysis
  - Only appropriate for intermediate levels

- $V_g = \left( egin{array}{ccc} \sigma_{win_1}^2 & & 0 \\ & \ddots & \\ 0 & & \sigma_{win_N}^2 \end{array} 
  ight)$
- Assumes the between-run variability=0
- Why would we do this?
  - What if df are low?
    - $-\hat{\sigma}_g^2$  has high variance
    - If  $\hat{\sigma}_g^2$  is too large, it will override differences in  $\hat{\sigma}_{win_k}^2$



- 3 contrast estimates from 3 runs for 1 subject
- Dotted line indicates variance
- "o" marks unweighted mean





### **Fixed Effects**

- Use to improve your mean estimates
  - eg correct trials
- Since variance is underestimated, you \*must\* only run this at an intermediate level
  - Higher level analysis soaks up rest of variance

## Third Level Analysis

- Typically Flame and OLS have similar results
  - Flame is probably the best choice, since it adjusts for heterogeneous variance
  - OLS runs faster
  - OLS stats can be larger or smaller than Flame stats
- FE at level 3 is bad
  - Variance is underestimated
  - High risk of false positives

# What is the "group" column for?

- This is the first column in the 2<sup>nd</sup> level Feat analysis design
- Assigns groups for estimating different between subject variances
- Must have a separable design
- Typically best if left to all 1's

## Concluding Remarks

- Mixed models are appropriate for fMRI data
  - Include between-subject variance
  - Allows inference to be applied to entire population
- The two-stage summary statistics model
  - Computationally easier to estimate
  - Easier to add new subjects
- Software packages use the same basic model, but estimate  $\sigma_g^2$  differently
- Use FE at intermediate levels and Flame at the top level in FSL