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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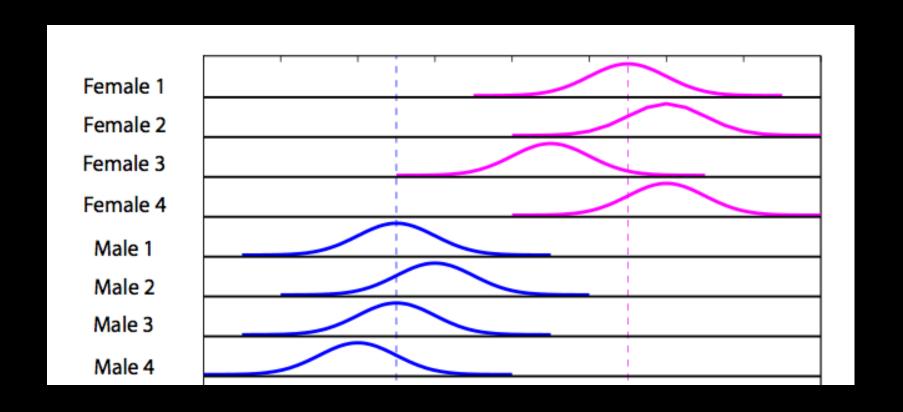
Mixed Model Motivation

- 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

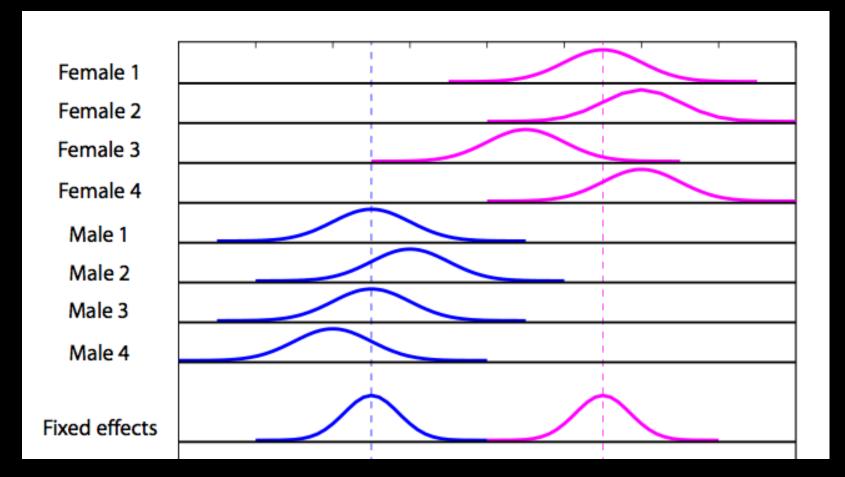


Each distribution has a variance of 1

Fixed effects analysis

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

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

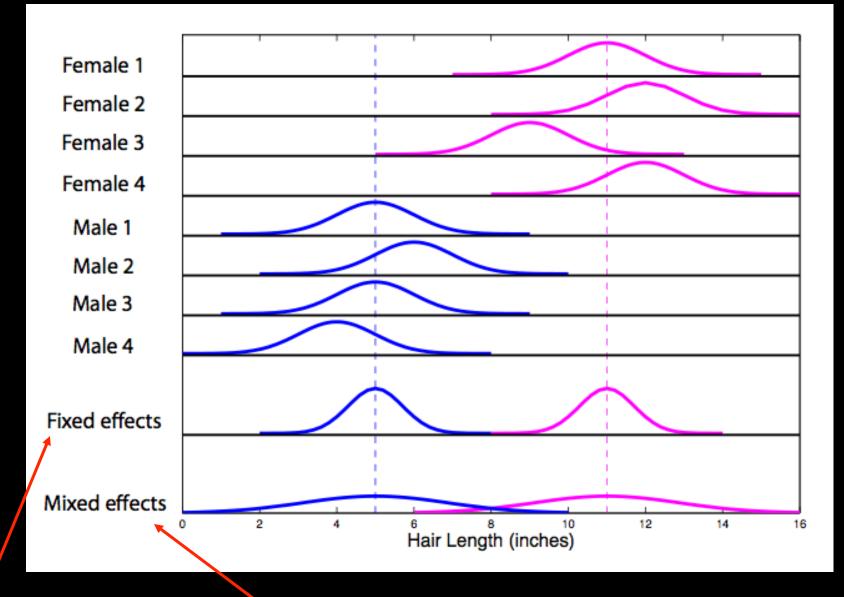


$$\sigma_{\scriptscriptstyle ext{FFX}}^2 = rac{1}{4}\sigma_{\scriptscriptstyle ext{W}}^2 = 0.25$$

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

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

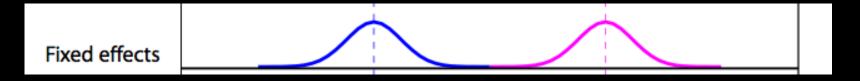
Mixed effects variance

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

Between subject variance typically dominates

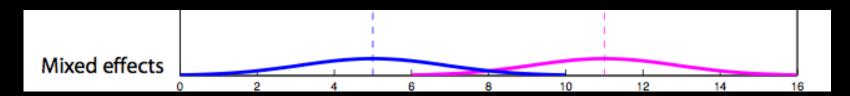
Wrong model leads to wrong conclusion

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



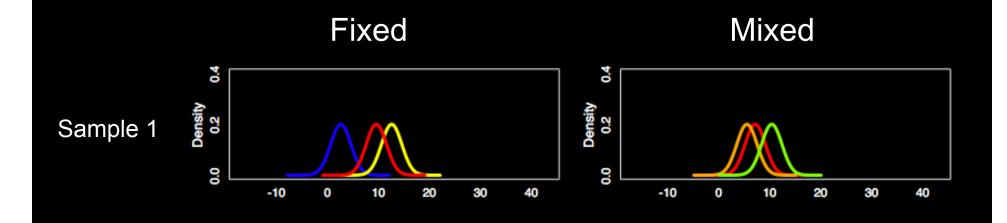
Wrong model leads to wrong conclusion

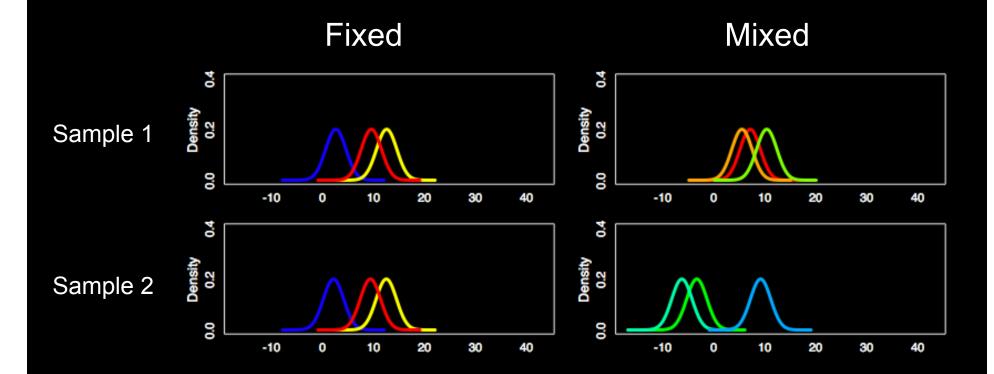
- Scenario 2: Mixed effects model
 - Cannot conclude there is a difference in hair length

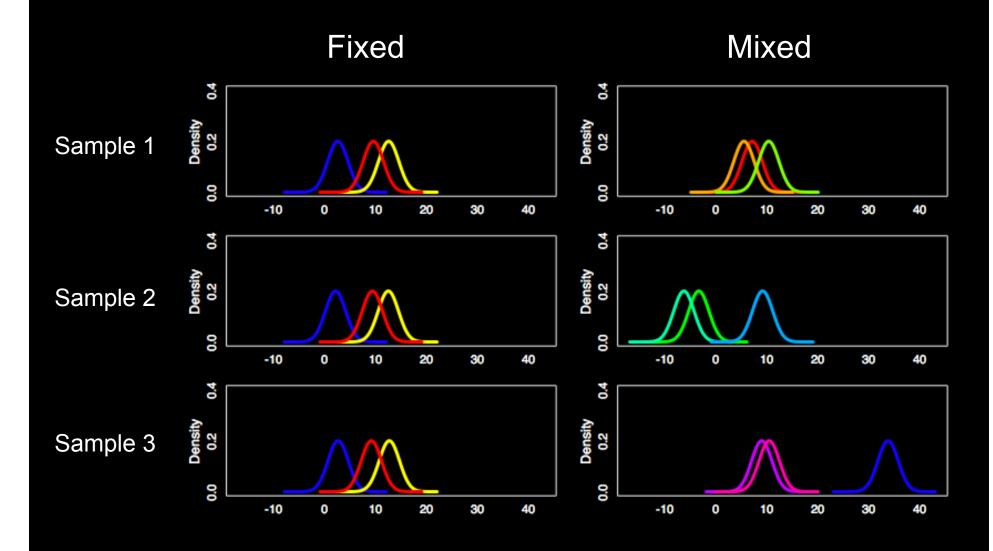


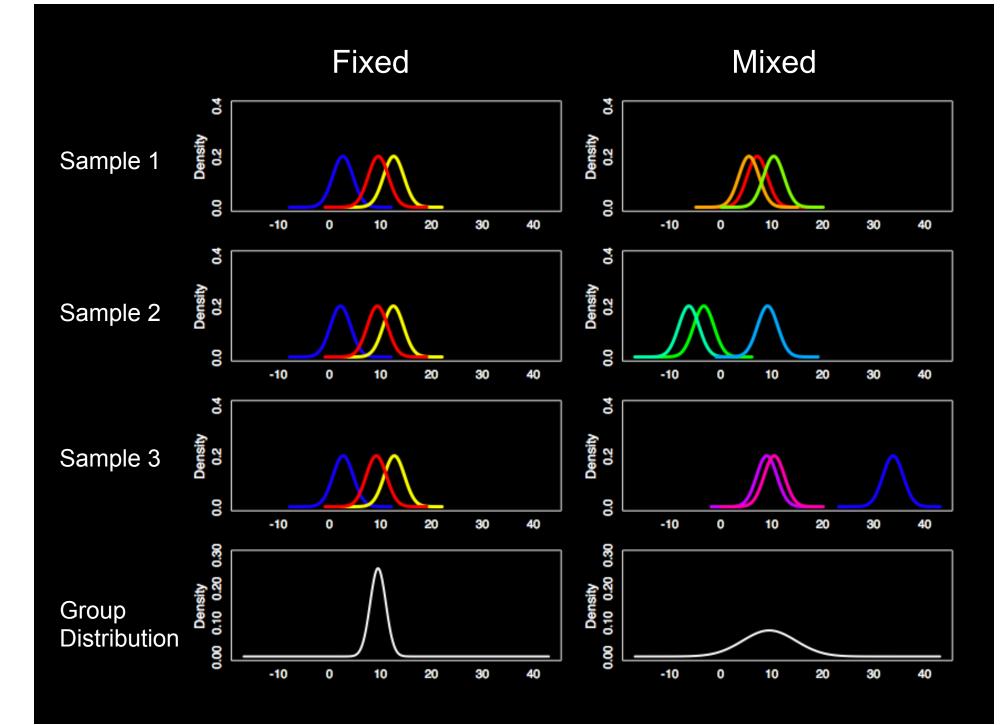
Mixed Model Comments

- 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?
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1 hair per subject

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

25 hairs per subject

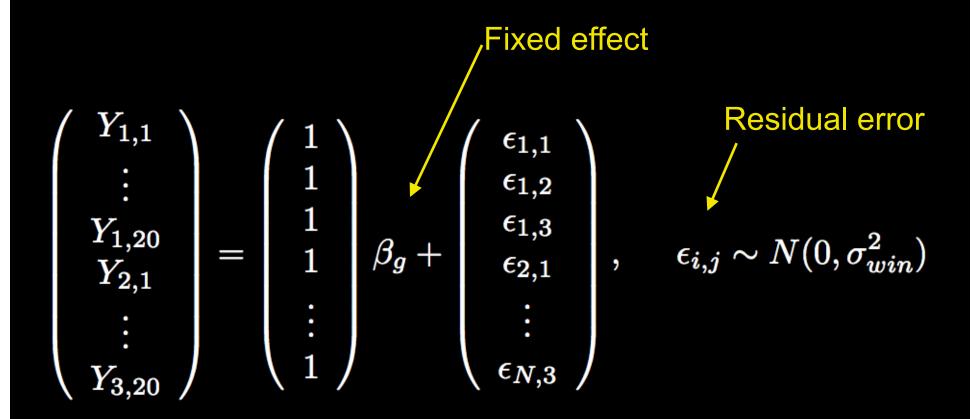
$$\sigma_{\text{MFX}}^2 = \sigma_{\text{W}}^2/4 + \sigma_{\text{B}}^2/4 = 12.5$$
 $\sigma_{\text{MFX}}^2 = \frac{1}{4}\sigma_{\text{W}}^2/25 + \frac{1}{4}\sigma_{\text{B}}^2 = 12.26$

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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

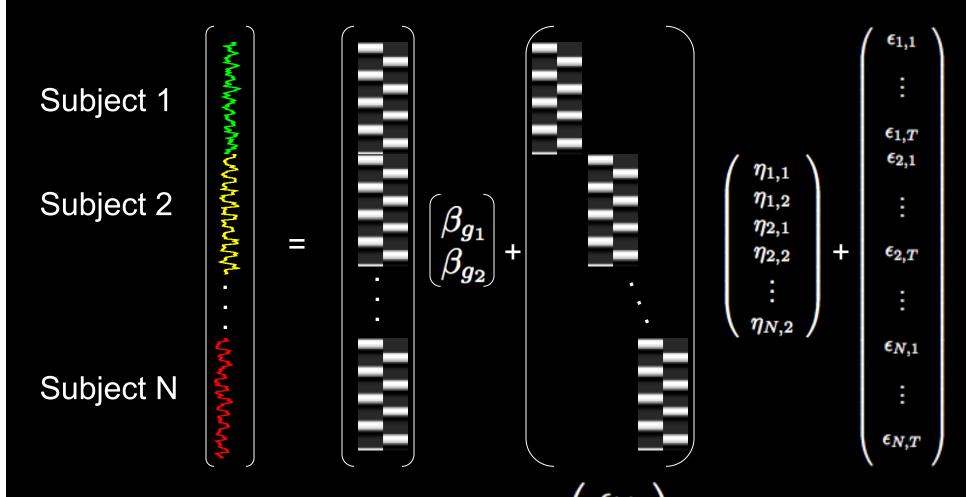
Subject N

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

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

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

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

Two-Stage Summary Statistics

Stage 1

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

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

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

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

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

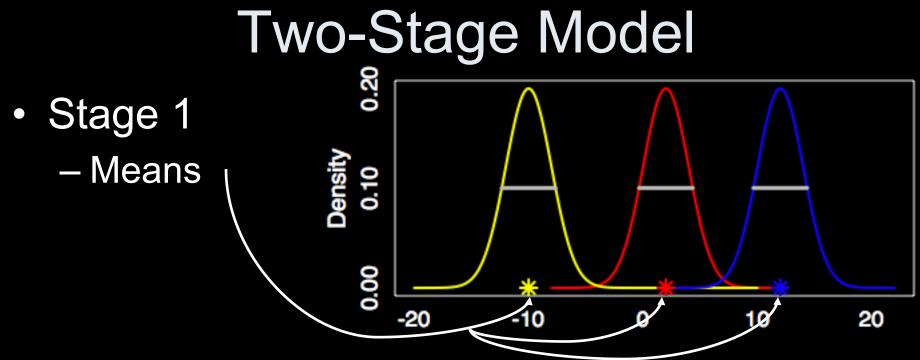
within

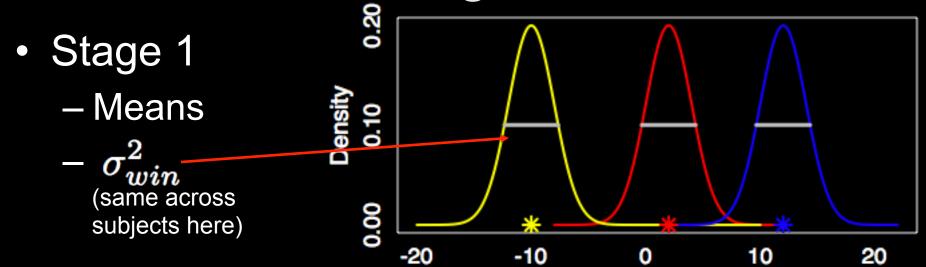
between

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

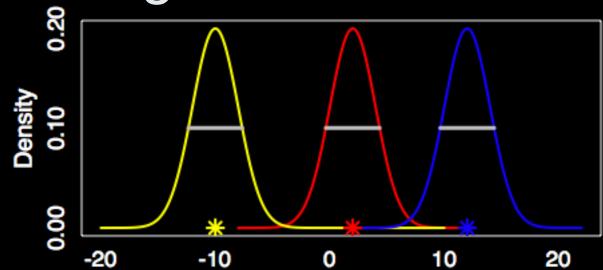
Stage 2

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





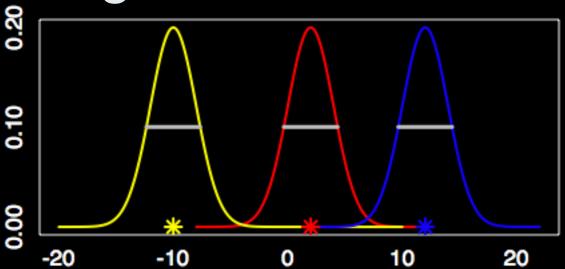
- Stage 1
 - Means
 - σ²_{win}
 (same across subjects here)



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



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

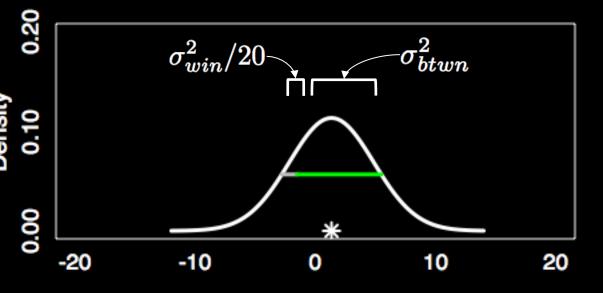


Stage 2

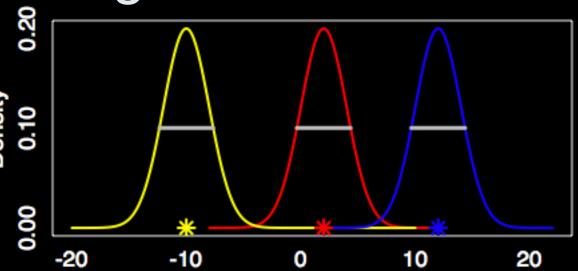
$$=\sigma_{btwn}^2$$

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

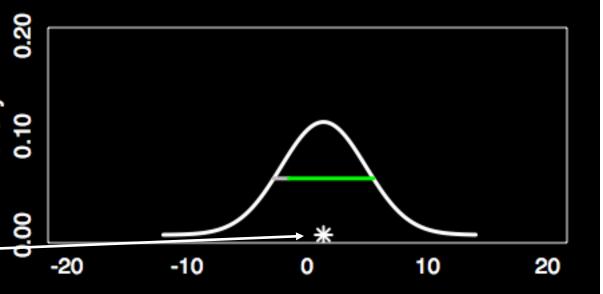
• 20 hairs/subject



- Stage 1
 - Means
 - σ_{win}^2 (same across subjects here)



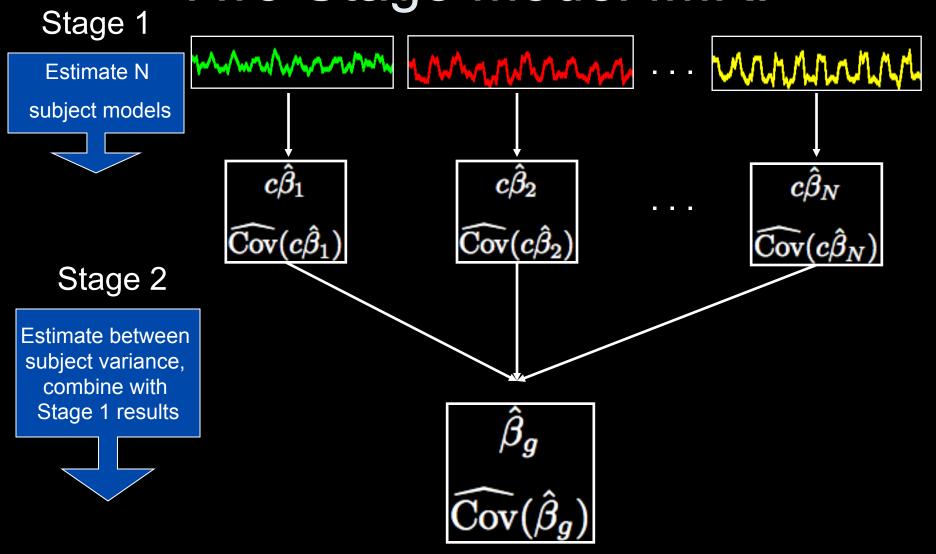
- Stage 2
 - $-\sigma_{btwn}^2$
 - $\begin{array}{l} \quad \quad \sigma_{mix}^2 = \\ \quad \frac{\sigma_{win}^2}{20} + \sigma_{btwn}^2 \end{array}$
 - 20 hairs/subject
- Pop mean



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

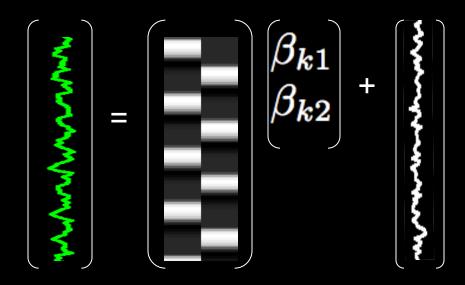
- -N = # subjects
- W = # measures within subject
- If new data are added, only run first stage for new data

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

$$eta_{cont} = X_g eta_g + \epsilon_g$$
 $c\hat{eta}_1$
 $c\hat{eta}_2$
 $= \begin{bmatrix} 1 \\ 1 \\ 1 \\ 1 \\ 1 \\ 1 \end{bmatrix}$
 eta_g
 eta_g
 $Cov(\epsilon_g) = V_g = \left(egin{array}{c} \sigma_1^2 c(X_1^{\star\prime} X_1^{\star\prime})^{-1} c' \\ & \ddots & \\ & & \sigma_N^2 c(X_N^{\star\prime} X_N^{\star\prime})^{-1} c' \end{array}
ight) + \sigma_g^2 I_N$

• 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 $W_g V_g W'_g = I_N$
- $W_g \hat{\beta}_{cont} = W_g X_g \beta_g + W_g \epsilon_g$ $\hat{\beta}_{cont}^* = X_g^* \beta_g + \epsilon_g^*$
- $\hat{\beta}_g = \left(X_g^{*'} X_g^*\right)^{-1} X_g^{*'} \hat{\beta}_{cont}^*$ $\widehat{Cov}(\hat{\beta}_g) = \left(X_g^{*'} X_g^*\right)^{-1}$
- $T = \hat{\beta}_g / \sqrt{\widehat{\mathrm{Cov}}(\hat{\beta}_g)}$

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 σ_g^2
 - Due to a set of assumptions, estimation of is unnecessary
 - FSL: Bayesian approach to estimating σ_g^2

SPM2

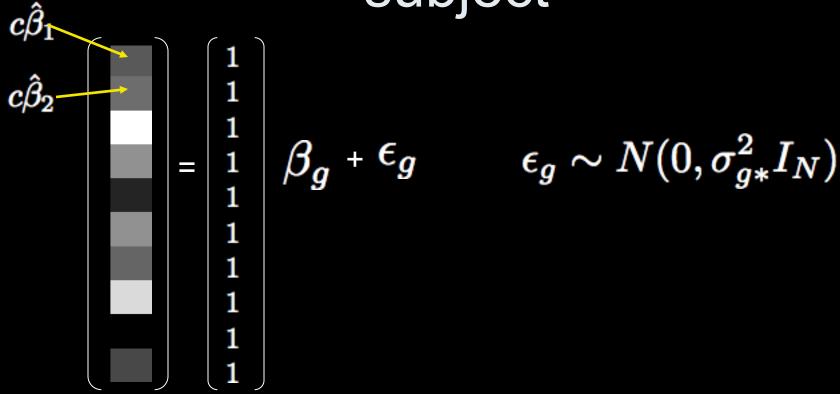
- Does not estimate σ_g^2
 - Assumes homogeneous variance across subjects
 - Assumes first level design is same across subjects

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

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

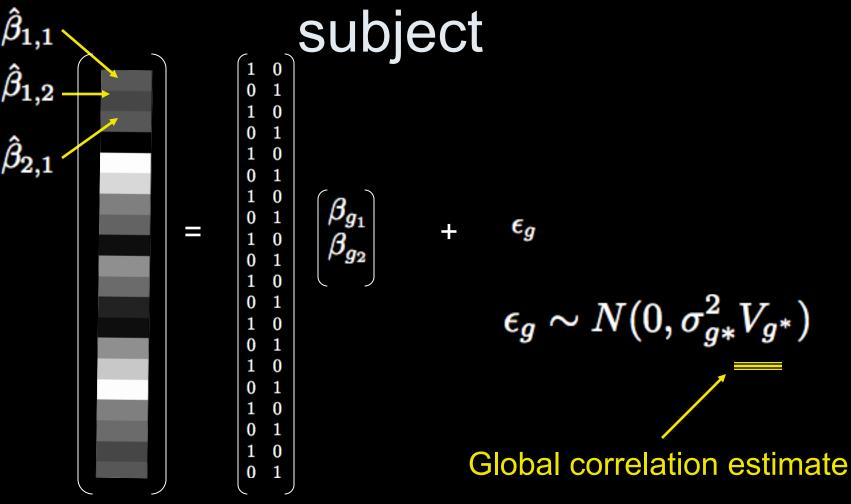
OLS can be used

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

- 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

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 σ_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

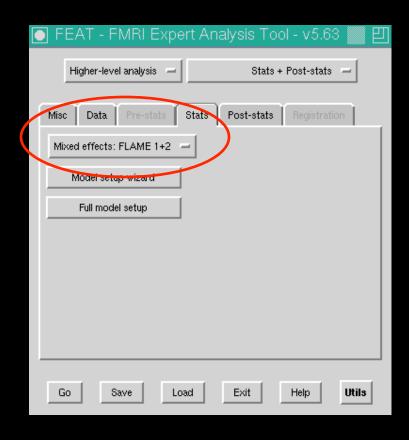
Cons

 Multiple contrasts in the group model are assumed to be independent

Which software?

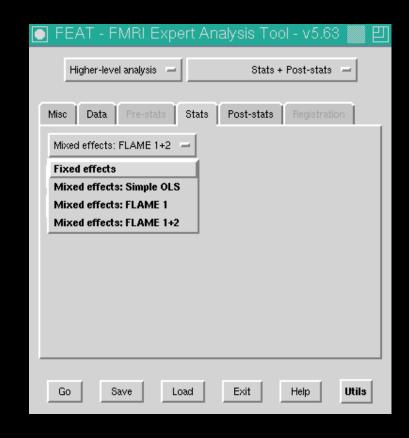
- FSL and FMRIstat best for heteroscedastic variances
 - Different number of trials per subject
- SPM best for multiple correlated contrasts at group level
- Other differences in first level modeling may sway users one way or another

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_g Matrix

 Recall we pre-multiply by W_g so our errors are uncorrelated and constant variance

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

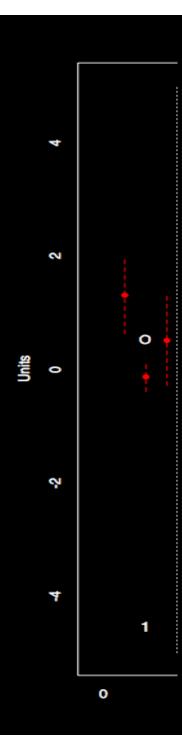
Act as

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

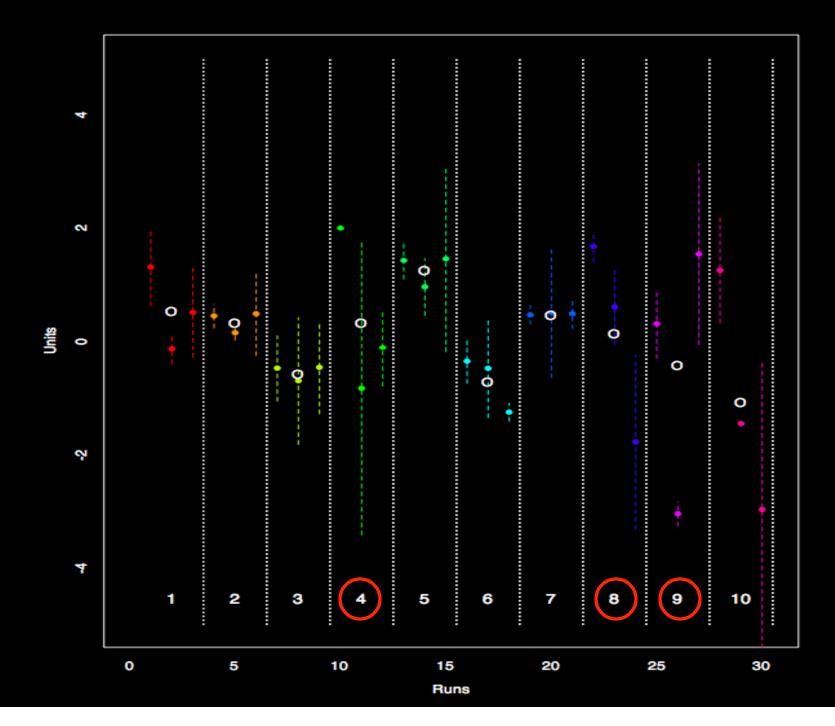
V_g Matrix Assumptions

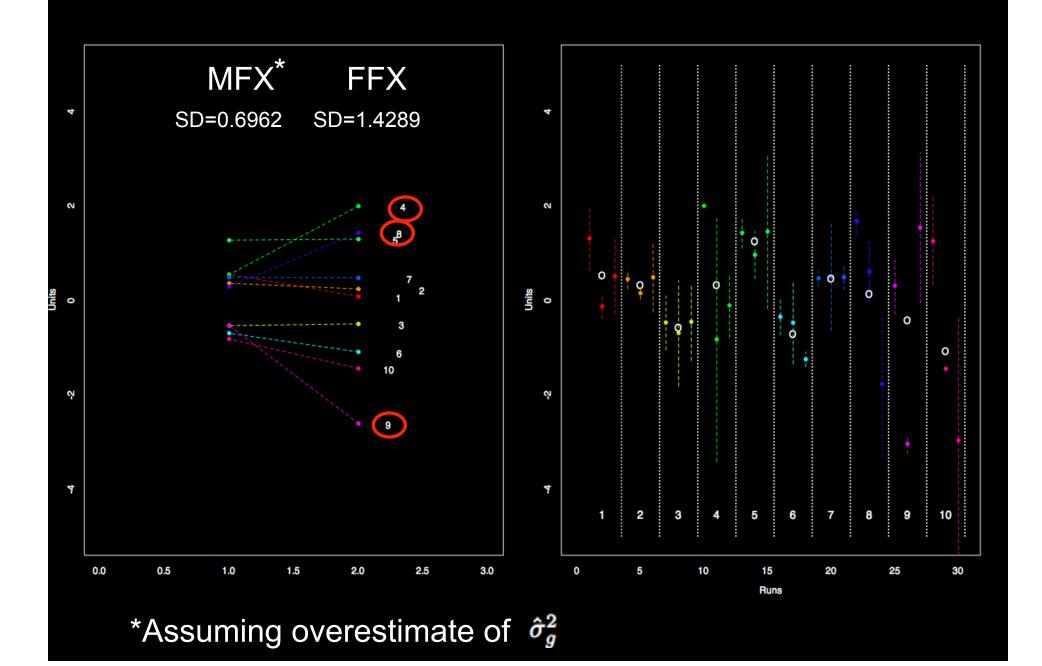
- Fixed effects analysis
 - Only appropriate for intermediate levels

- $V_g = \left(egin{array}{ccc} \sigma_{win_1}^2 & & 0 \\ & \ddots & \\ 0 & & \sigma_{win_N}^2 \end{array}
 ight)$
- 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 2nd 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 σ_g^2 differently
- Use FE at intermediate levels and Flame at the top level in FSL