

# ***Models with Random Effects***

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- Levels are a random sample
- Variability between levels is of interest
- Nested vs. crossed factors

# One Random Factor

Serum measurements of blood samples  
Model:

$$Y_{ij} = \mu + a_i + \epsilon_{ij}, \quad i = 1, \dots, I; j = 1, \dots, J$$

$a_i$  random effect of sample  $i$ ,  $a_i \sim \mathcal{N}(0, \sigma_a^2)$ ,

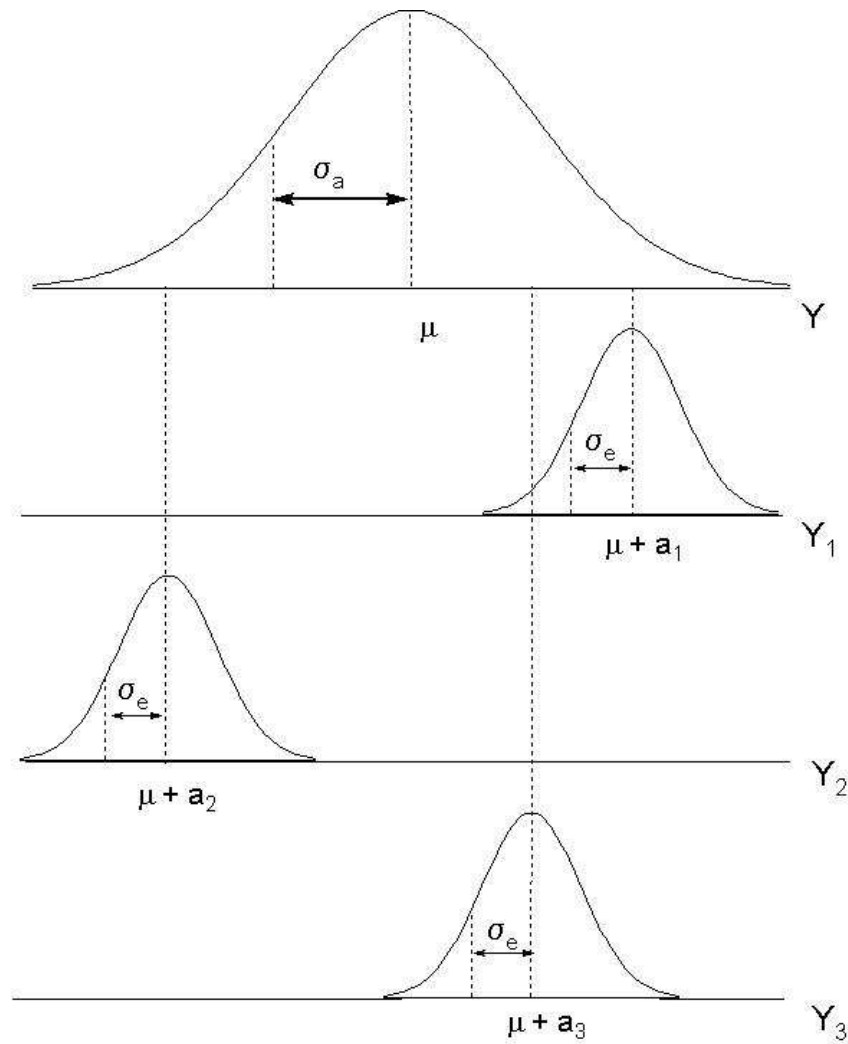
$\epsilon_{ij}$  error of  $j$ th measurement of sample  $i$ ,  $\epsilon_i \sim \mathcal{N}(0, \sigma_e^2)$ ,

$a_i$  and  $\epsilon_{ij}$  are all independent.

$$\text{Var}(Y_{ij}) = \text{Var}(a_i + \epsilon_{ij}) = \sigma_a^2 + \sigma_e^2, \quad \text{Cov}(Y_{ij}, Y_{ij'}) = \sigma_a^2$$

The variance of  $Y_{ij}$  consists of two components. Such models are also called **variance components models**.

# Illustration



# Anova Table

$$H_0 : \sigma_a^2 = 0 \quad H_A : \sigma_a^2 > 0$$

Source	SS	df	MS=SS/df
Sample	$SS_a = \sum \sum (y_{i.} - y_{..})^2$	$I - 1$	$MS_a$
Residual	$SS_{res} = \sum \sum (y_{ij} - y_{i.})^2$	$N - I$	$MS_{res}$
Total	$SS_{tot} = \sum \sum (y_{ij} - y_{..})^2$	$N - 1$	

# Parameter estimations

$$\hat{\sigma}_e^2 = MS_{res}$$

$$\hat{\sigma}_a^2 = (MS_a - MS_{res})/J \quad \text{can be negative!}$$

$$\hat{\mu} = y_{..} \quad \text{with } Var(\hat{\mu}) = \frac{1}{I}(\sigma_a^2 + \sigma_e^2/J)$$

Either Maximum Likelihood estimators or  $\hat{\sigma}_a^2 \geq 0$

# Variability between Laboratories

$$Y_{ijk} = \mu + a_i + b_j + \epsilon_{ijk}$$

$a_i$  random effect of lab  $i$ ,  $a_i \sim \mathcal{N}(0, \sigma_a^2)$ ,  
 $b_j$  random effect of sample  $j$ ,  $b_j \sim \mathcal{N}(0, \sigma_b^2)$ ,  
 $\epsilon_{ijk}$  measurement error,  $\epsilon_{ijk} \sim \mathcal{N}(0, \sigma_e^2)$ ,  
all random variables are independent.

Source	df	E(MS)	F
Lab	$I - 1$	$\sigma_e^2 + JK\sigma_a^2$	$MS_a/MS_{res}$
Sample	$J - 1$	$\sigma_e^2 + IK\sigma_b^2$	$MS_b/MS_{res}$
Residual	$\ll diff \gg$	$\sigma_e^2$	
Total	$IJK - 1$		

# Parameter Estimation

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$$\hat{\sigma}_e^2 = MS_{res}$$

$$\hat{\sigma}_a^2 = (MS_a - MS_{res})/JK$$

$$\hat{\sigma}_b^2 = (MS_b - MS_{res})/IK$$

# Model with Interaction Lab:Sample

Source	E(MS)	$H_0$	F
Lab	$\sigma_e^2 + JK\sigma_a^2 + K\sigma_{ab}^2$	$\sigma_a^2 = 0$	$MS_a/MS_{ab}$
Sample	$\sigma_e^2 + IK\sigma_b^2 + K\sigma_{ab}^2$	$\sigma_b^2 = 0$	$MS_b/MS_{ab}$
Lab : Sample	$\sigma_e^2 + K\sigma_{ab}^2$	$\sigma_{ab}^2 = 0$	$MS_{ab}/MS_{res}$
Residual	$\sigma_e^2$		

$H_0 : \sigma_a^2 = 0$       Test statistic:  $F = MS_a/MS_{ab}$

$H_0 : \sigma_a^2 = \sigma_{ab}^2 = 0$       Test statistic:  $F = MS_a/MS_{res}$



# Crossed factors

Factors A and B are called **crossed** if every level of B occurs with every level of A. A factorial design involves crossed factors.

	Factor A			
Factor B	1	2	3	4
1	xx	xx	xx	xx
2	xx	xx	xx	xx
3	xx	xx	xx	xx

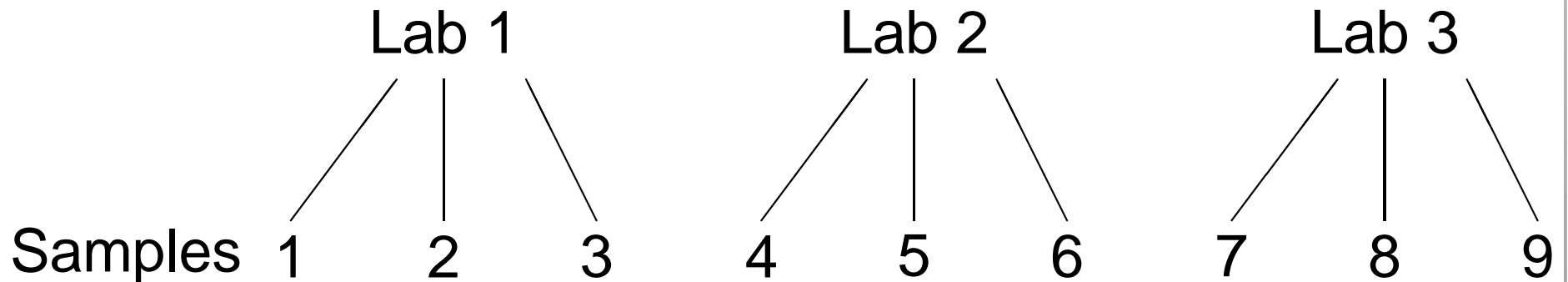
## ***Nested factors***

Factors A and B are called **nested** if there are different levels of B within each level of A. B is nested within A in the following layout.

A		1			2			3			4	
B	1	2	3	4	5	6	7	8	9	10	11	12
	XX	XX	XX	XX	XX	XX	XX	XX	XX	XX	XX	XX

Designs with nested factors are called **nested designs** or **hierarchical designs**.

# *Nested Designs*



Factors Lab and Sample are not **crossed**, but **nested**.

Model for a two-stage nested design:

$$Y_{ijk} = \mu + a_i + b_{j(i)} + \epsilon_{k(ij)}, \quad i = 1, \dots, I; j = 1, \dots, J; k = 1, \dots, K$$

The subscript  $j(i)$  indicates that the  $j$ th level of factor B is nested within the  $i$ th level of factor A.

# Anova table

Decomposition of sum of squares:

$$SS_{tot} = SS_A + SS_{B(A)} + SS_{res}.$$

Source	df	E(MS)
Lab	$I - 1$	$\sigma_e^2 + K\sigma_b^2 + JK\sigma_a^2$
Sample	$I(J - 1)$	$\sigma_e^2 + K\sigma_b^2$
Residual	"diff"	$\sigma_e^2$
Total	$IJK - 1$	

# Moisture Content of Cowpea

Effect of milling on moisture content. 3 samples of 100g from 5 batches were milled. From each sample 10g are measured.

batch	sample								
	1	2	3	4	5	6	7	8	9
1	9.3	9.2	8.8	8.6	8.7	9.9	8.9	8.7	8.5
2	8.0	8.2	9.2	9.7	9.4	8.2	9.3	9.5	9.4
3	11.0	10.7	9.9	9.3	13.9	9.2	9.2	10.9	9.7
4	10.1	10.2	9.9	8.6	9.4	8.3	8.3	9.9	9.5
5	12.0	9.3	10.8	12.2	9.6	11.7	11.4	9.8	12.4

# Anova Table

```
> mod1=aov(moisture~batch/sample)
```

```
> summary(mod1)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
batch	4	30.928	7.7320	7.0390	0.0004027	**
batch:sample	10	5.911	0.5911	0.5381	0.8491520	
Residuals	30	32.953	1.0984			

$$\hat{\sigma}_e^2 = 1.0984$$

$$\hat{\sigma}_e = 1.048$$

$$\hat{\sigma}_s^2 = (0.5911 - 1.0984)/3 = 0$$

$$\hat{\sigma}_s = 0$$

$$\hat{\sigma}_b^2 = (7.732 - 1.0984)/9 = 0.737$$

$$\hat{\sigma}_b = 0.858$$

# Linear mixed-effects model fit

```
> summary(lme(moisture~1,random=~1|batch/sample))
```

Random effects:

```
Formula: ~1 | batch
```

```
(Intercept)
```

```
StdDev: 0.8666916
```

```
Formula: ~1 | sample %in% batch
```

```
(Intercept) Residual
```

```
StdDev: 3.783493e-05 0.9857034
```

```
Number of Observations: 45
```

```
Number of Groups: batch sample %in% batch
```

```
5
```

```
15
```