

## Random effects ANOVA

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ANOVA limitations

Hierarchical normal model

Estimation and inference

## Classical data analysis and estimation

The “classical” testing and estimation procedure is as follows:

If the  $p$ -value  $< 0.05$ ,

- reject  $H_0$ , and conclude there are group differences,
- estimate  $\theta_j$  with  $\bar{y}_{\cdot j}$ .

$$\hat{\theta}_j = \bar{y}_{\cdot j}$$

If the  $p$ -value  $> 0.05$ ,

- accept  $H_0$ , and conclude there is no evidence of group differences,
- estimate  $\theta_j$  with  $\bar{y}_{\cdot\cdot}$ .

$$\hat{\theta}_j = \bar{y}_{\cdot\cdot}$$

Note that the estimator of  $\theta_j$  can be written as

$$\hat{\theta}_j = w\bar{y}_j + (1 - w)\bar{y}_{\cdot\cdot}$$

## Classical data analysis and estimation

### Advantages of classical procedure:

- controls the type I error rate of rejecting  $H_0$ ;
- is easy to implement and report.

### Disadvantages:

- rejecting  $H_0$  doesn't mean no **similarities** across groups  
⇒  $\bar{y}_{\cdot j}$  is an inefficient estimate of  $\theta_j$
- accepting  $H_0$  doesn't mean no **differences** between groups  
⇒  $\bar{y}_{\cdot \cdot}$  is an inaccurate estimate of  $\theta_j$ .

## An alternative strategy

$$\hat{\theta}_j = w\bar{y}_j + (1 - w)\bar{y}_{..}$$

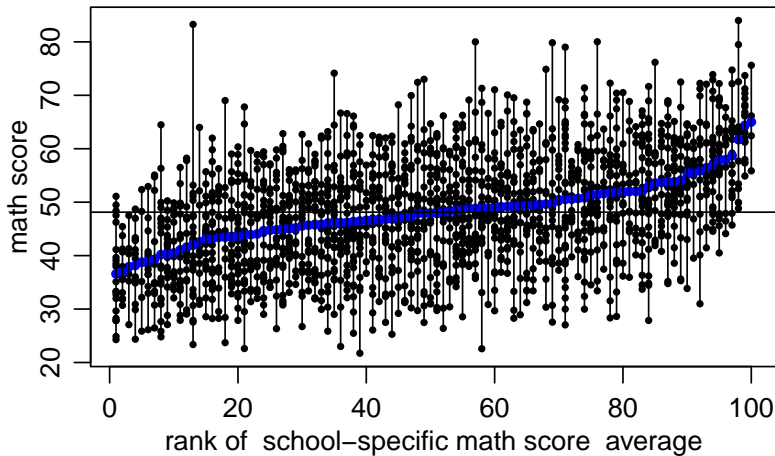
**Classical approach:**  $w$  is the indicator of rejecting  $H_0$ .

**Multilevel approach:**  $w = \frac{n/\hat{\sigma}^2}{n/\hat{\sigma}^2 + 1/\hat{\tau}^2}$

The multilevel approach will allow for

- sharing of information across groups,
- the amount of sharing to be estimated from the data.

## Example: Test scores



## Example: Test scores

```
y.3122<-ndat$mathscore[ndat$school=="3122"]
y.2832<-ndat$mathscore[ndat$school=="2832"]

y.3122
## [1] 75.62 55.86 66.16 62.43

y.2832
## [1] 66.26 66.12 71.22 54.90 61.98 69.42 61.22 62.99 57.99 61.33 66.85 67.87
## [13] 63.94 73.70 70.36 64.01 57.35 68.25 57.39

mean(ndat$mathscore)
## [1] 48.07446

mean(y.3122)
## [1] 65.0175

mean(y.2832)
## [1] 64.37632
```

## Example: Test scores

$$\begin{array}{ccccc} 48.0744556 & < & 64.3763158 & < & 65.0175 \\ \bar{y}_{..} & < & \bar{y}_{2832} & < & \bar{y}_{3122} \end{array}$$

but

$$n_{3122} = 4 < 19 = n_{2832}$$

Based on the data  $\{y_{i,j}\}$ , how would you estimate  $\theta_{3122}$  and  $\theta_{2832}$ ?

**Ignoring across-group information :**

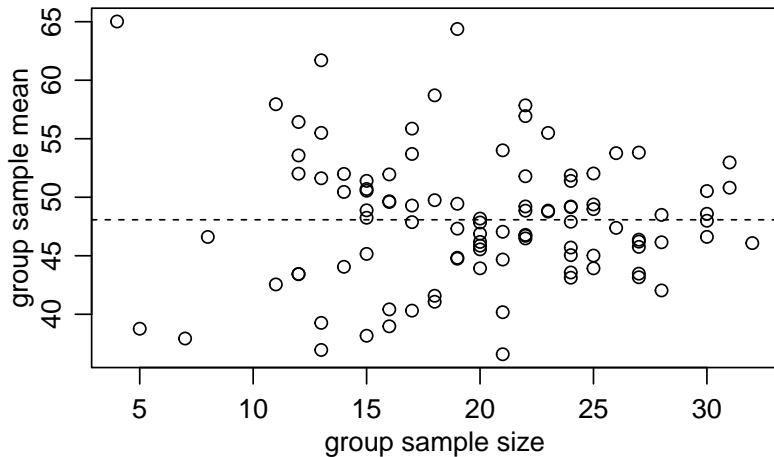
- $\hat{\theta}_{2832} = \bar{y}_{2832} = 64.3763158$
- $\hat{\theta}_{3122} = \bar{y}_{3122} = 65.0175$
- $\hat{\theta}_{2832} < \hat{\theta}_{3122}$

**Considering across-group information and sample size:**  $\bar{y}_{..} = 48.0744556$ .

- $\bar{y}_{..} < \hat{\theta}_{2832} < \bar{y}_{2832} = 64.3763158$
- $\bar{y}_{..} < \hat{\theta}_{3122} < \bar{y}_{3122} = 65.0175$
- $\hat{\theta}_{2832} \geq \hat{\theta}_{3122} ?$



## Example: Test scores



## Example: Test scores

### Possible explanations for $\bar{y}_{3122}$ :

- $\bar{y}_{3122}$  is large because  $\theta_{3122}$  is large;
- $\bar{y}_{3122}$  is large because  $\text{sd}(\bar{y}_{3122})$  is large.

### Possible explanations for $\bar{y}_{2832}$ :

- $\bar{y}_{2832}$  is large because  $\theta_{2832}$  is large;
- $\bar{y}_{2832}$  is large because  $\text{sd}(\bar{y}_{2832})$  is large (but is smaller than  $\text{sd}(\bar{y}_{3122})$ ).

The plausibility of the explanations will depend on

- the group specific sample sizes,  $n_1, \dots, n_m$ ;
- the observed across-group heterogeneity.

## Example: Free throws

```
ftdat[1:20,]
```

| ##    | PLAYER1  | PLAYER2    | TEAM | MIN  | FTM | FTA | FT.   |
|-------|----------|------------|------|------|-----|-----|-------|
| ## 1  | Sam      | Jacobson   | LAL  | 12   | 2   | 2   | 1.000 |
| ## 2  | Steve    | Henson     | DET  | 25   | 2   | 2   | 1.000 |
| ## 3  | Radoslav | Nesterovic | MIN  | 30   | 2   | 2   | 1.000 |
| ## 4  | Bryce    | Drew       | HOU  | 441  | 8   | 8   | 1.000 |
| ## 5  | Charles  | O'bannon   | DET  | 165  | 8   | 8   | 1.000 |
| ## 6  | Marty    | Conlon     | MIA  | 35   | 2   | 2   | 1.000 |
| ## 7  | Mikki    | Moore      | DET  | 6    | 2   | 2   | 1.000 |
| ## 8  | John     | Crotty     | POR  | 19   | 3   | 3   | 1.000 |
| ## 9  | Gerald   | Wilkins    | ORL  | 28   | 2   | 2   | 1.000 |
| ## 10 | Korleone | Young      | DET  | 15   | 2   | 2   | 1.000 |
| ## 11 | Brian    | Evans      | MIN  | 145  | 4   | 4   | 1.000 |
| ## 12 | Pooh     | Richardson | LAC  | 130  | 4   | 4   | 1.000 |
| ## 13 | Michael  | Hawkins    | SAC  | 203  | 3   | 3   | 1.000 |
| ## 14 | Randy    | Livingston | PHO  | 22   | 2   | 2   | 1.000 |
| ## 15 | Rusty    | Larue      | CHI  | 732  | 17  | 17  | 1.000 |
| ## 16 | Fred     | Hoiberg    | IND  | 87   | 6   | 6   | 1.000 |
| ## 17 | Herb     | Williams   | NYK  | 34   | 2   | 2   | 1.000 |
| ## 18 | Ryan     | Stack      | CLE  | 199  | 19  | 20  | 0.950 |
| ## 19 | Sam      | Cassell    | MIL  | 199  | 47  | 50  | 0.940 |
| ## 20 | Reggie   | Miller     | IND  | 1787 | 226 | 247 | 0.915 |

Who does Indiana pick to shoot its technical foul free throws?

## Further limitations of ANOVA

In the wheat yield example we might be interested in

- (1) what the yield might be in other plots of land in these 10 regions, or
- (2) what the yield might be in other regions.

For general hierarchical data, these questions translate into

- (1) making inference about units within groups in our study;
- (2) making inference about groups that weren't in our study.

Inference for (1) can be obtained with ANOVA.

Inference for (2) requires

- treating the  $m$  groups as a sample from a larger population;
- a statistical model for this larger population.

## The hierarchical normal model

$$y_{i,j} = \mu + a_j + \epsilon_{i,j} \quad (1)$$

$$\{\epsilon_{1,1}, \dots, \epsilon_{n_1,1}\}, \dots, \{\epsilon_{1,m}, \dots, \epsilon_{n_m,m}\} \sim \text{i.i.d. normal}(0, \sigma^2) \quad (2)$$

$$a_1, \dots, a_m \sim \text{i.i.d. normal}(0, \tau^2) \quad (3)$$

The classical ANOVA model consists of (1) and (2).

The HNM assumes the sampling model (3) for the groups.

- $\{a_1, \dots, a_m\}$  represent differences across groups
- $\{\epsilon_{i,j}\}$  represent differences within groups

The HNM represents this heterogeneity in terms of population variances:

$$\text{Var}[a] = \tau^2 = \text{across-group variance}$$

$$\text{Var}[\epsilon] = \sigma^2 = \text{within-group variance}$$

## Marginal and conditional variation

Two levels of heterogeneity require two versions of variance and covariance:

### Within-group variance:

- The variance of  $y_{i,j}$  around  $\theta_j$ ;
- Describes heterogeneity/variance within a particular group;
- Mathematically, is calculated *conditionally* on group-level parameters.

### Population-level variance:

- Variance of  $y_{i,j}$  around  $\mu$ ;
- Describes heterogeneity/variance across the population;
- Mathematically, is calculated *marginally* over group-level parameters.

## Conditional variance and covariance

For a fixed group  $j$ ,

$$\{y_{1,j}, \dots, y_{n_j,j}\} | \mu, a_j, \sigma^2 \sim \text{i.i.d. normal}(\mu + a_j, \sigma^2)$$

$$\{y_{1,j}, \dots, y_{n_j,j}\} | \theta_j, \sigma^2 \sim \text{i.i.d. normal}(\theta_j, \sigma^2)$$

Variation *around the group mean*  $\theta_j$  is as follows

$$E[y_{i,j} | \mu, a_j] = \mu + a_j = \theta_j$$

$$\text{Var}[y_{i,j} | \mu, a_j] = \sigma^2,$$

$$\text{Cov}[y_{i_1,j}, y_{i_2,j} | \mu, a_j] = 0.$$

In words,

- sample observations *from the group* are centered around  $\theta_j$ ;
- the variation of the sample *around*  $\theta_j$  is  $\sigma^2$ ;
- the observations within a group are uncorrelated *around*  $\theta_j$ .

**Regarding correlation:** Knowing how far  $y_{1,j}$  is from  $\theta_j$  doesn't inform you about about how far  $y_{2,j}$  is from  $\theta_j$ .

## Within-group variance and covariance

$$y_{i,j} = \mu + \alpha_j + \epsilon_{i,j}$$

$$y_{i,j} = \theta_j + \epsilon_{i,j}$$

$$\begin{aligned}\text{Var}[y_{i,j}|\theta_j] &\equiv \text{E}[(y_{i,j} - \text{E}[y_{i,j}|\theta_j])^2|\theta_j] \\ &= \text{E}[(y_{i,j} - \theta_j)^2|\theta_j] \\ &= \text{E}[(\theta_j + \epsilon_{i,j} - \theta_j)^2|\theta_j] \\ &= \text{E}[\epsilon_{i,j}^2|\theta_j] = \sigma^2\end{aligned}$$

$$\begin{aligned}\text{Cov}[y_{i_1,j}, y_{i_2,j}|\theta_j] &\equiv \text{E}[(y_{i_1,j} - \text{E}[y_{i_1,j}|\theta_j]) \times (y_{i_2,j} - \text{E}[y_{i_2,j}|\theta_j])|\theta_j] \\ &= \text{E}[(y_{i_1,j} - \theta_j) \times (y_{i_2,j} - \theta_j)|\theta_j] \\ &= \text{E}[\epsilon_{i_1,j}\epsilon_{i_2,j}|\theta_j] = 0\end{aligned}$$



## Population level variance and covariance

Across all groups,

$$\begin{aligned}a_1, \dots, a_m &\sim \text{i.i.d. normal}(0, \tau^2) \\ \{y_{1,j}, \dots, y_{n_j,j}\} &\sim \text{i.i.d. normal}(\mu + a_j, \sigma^2)\end{aligned}$$

For a randomly sampled observation  $i$  from a randomly sampled group  $j$ ,

$$\begin{aligned}\mathbb{E}[y_{i,j}|\mu] &= \mathbb{E}[\mu + a_j + \epsilon_{i,j}|\mu] \\ &= \mathbb{E}[\mu|\mu] + \mathbb{E}[a_j|\mu] + \mathbb{E}[\epsilon_{i,j}|\mu] \\ &= \mu + 0 + 0 = \mu\end{aligned}$$

This is the *population mean*.

## Population level variance and covariance

Variation *around the population mean  $\mu$*  is as follows:

$$\begin{aligned}E[y_{i,j}|\mu] &= E[\mu + a_j|\mu] = \mu + 0 = \mu, \\ \text{Var}[y_{i,j}|\mu] &= \sigma^2 + \tau^2, \\ \text{Cov}[y_{i_1,j}, y_{i_2,j}|\mu] &= \tau^2.\end{aligned}$$

In words,

- sampled observations *across groups* are centered around  $\mu$ ;
- the variation of the sample *around  $\mu$*  is  $\sigma^2 + \tau^2$ ;
- the observations within a group are correlated *around  $\mu$* .

**Regarding correlation:** Knowing how far  $y_{1,j}$  is from  $\mu$  *does* inform you about how far  $y_{2,j}$  is from  $\mu$ .

## Population level variance

$$\begin{aligned}\text{Var}[y_{i,j}|\mu] &\equiv \text{E}[(y_{i,j} - \text{E}[y_{i,j}|\mu])^2|\mu] \\ &= \text{E}[(y_{i,j} - \mu)^2|\mu] \\ &= \text{E}[(\mu + a_j + \epsilon_{i,j} - \mu)^2|\mu] \\ &= \text{E}[(a_j + \epsilon_{i,j})^2|\mu] \\ &= \text{E}[a_j^2 + 2a_j\epsilon_{i,j} + \epsilon_{i,j}^2|\mu] \\ &= \tau^2 + 0 + \sigma^2 = \sigma^2 + \tau^2\end{aligned}$$

Exercise: Draw a picture of within and across group sampling.

## Population level covariance and correlation

$$\begin{aligned}\text{Cov}[y_{i_1,j}, y_{i_2,j} | \mu] &\equiv \text{E}[(y_{i_1,j} - \text{E}[y_{i_1,j} | \mu]) \times (y_{i_2,j} - \text{E}[y_{i_2,j} | \mu]) | \mu] \\ &= \text{E}[(y_{i_1,j} - \mu) \times (y_{i_2,j} - \mu) | \mu] \\ &= \tau^2\end{aligned}$$

$$\begin{aligned}\text{Cor}[y_{i_1,j}, y_{i_2,j} | \mu] &\equiv \frac{\text{Cov}[y_{i_1,j}, y_{i_2,j} | \mu]}{\sqrt{\text{Var}[y_{i_1,j} | \mu] \text{Var}[y_{i_2,j} | \mu]}} \\ &= \frac{\tau^2}{\tau^2 + \sigma^2} \equiv \rho\end{aligned}$$

The correlation  $\rho$  is the *intraclass correlation coefficient*.

## Estimation of $\tau^2$ and $\rho$

The easiest way to estimate  $\tau^2$  is using the method-of-moments. Recall,

$$\begin{aligned}MSA &= \frac{1}{m-1} \sum_j \sum_i (\bar{y}_j - \bar{y}_{..})^2 \\&= \frac{n}{m-1} \sum (\bar{y}_j - \bar{y}_{..})^2 \\E[MSA|a_1, \dots, a_m] &= \frac{n}{m-1} \left( \frac{m-1}{n} \sigma^2 + \sum a_j^2 \right) \\&= \sigma^2 + n \times \frac{1}{m-1} \sum a_j^2.\end{aligned}$$

The expectation of  $MSA$  over samples *and* groups is given by

$$\begin{aligned}E[E[MSA|a_1, \dots, a_m]] &= E[\sigma^2 + n \times \frac{1}{m-1} \sum a_j^2] \\&= \sigma^2 + n \times E[\frac{1}{m-1} \sum a_j^2] \\&= \sigma^2 + n\tau^2.\end{aligned}$$

(In the ANOVA parameterization,  $\sum a_j^2 = \sum (a_j - \bar{a})^2$  because  $\bar{a} = 0$ )

## Estimation of $\tau^2$ and $\rho$

The result suggests

$$\widehat{\sigma^2 + n\tau^2} = MSA.$$

How to estimate  $\tau^2$ ? Recall  $E[MSW] = \sigma^2$ , so we can use

$$\hat{\sigma}^2 = MSW.$$

This suggests

$$\begin{aligned}\widehat{n\tau^2} &= MSA - MSW \\ \hat{\tau}^2 &= (MSA - MSW)/n.\end{aligned}$$

### Comments:

- $MSA - MSW$  could be negative. If so, it is standard to set  $\hat{\tau}^2 = 0$ .
- If sample sizes are unequal, the formula must be modified slightly:

$$\hat{\tau}^2 = (MSA - MSW)/\tilde{n}$$

where there is a horrible formula for  $\tilde{n}$ .

## Unequal sample sizes

$$\hat{\tau}^2 = (MSA - MSW)/\tilde{n}$$

$$\tilde{n} = \bar{n} - \frac{\text{sample variance}(n_1, \dots, n_m)}{m\bar{n}}$$

where  $\bar{n} = \sum_j n_j / m = \text{sample mean}(n_1, \dots, m_m)$ .

## Estimation of $\tau^2$ and $\rho$

It is common to use a “plug-in” estimate of  $\rho$ :

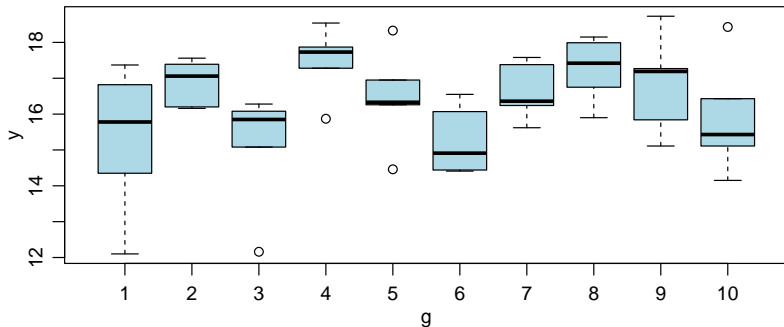
$$\hat{\rho} = \frac{\widehat{\tau^2}}{\tau^2 + \sigma^2} = \frac{\hat{\tau}^2}{\hat{\tau}^2 + \hat{\sigma}^2}.$$

A standard error for  $\rho$  (with which we can get a CI) is

$$\text{se}(\hat{\rho}) = (1 - \hat{\rho}) \times (1 + (n - 1)\hat{\rho}) \sqrt{\frac{2}{n(n - 1)(m - 1)}}.$$



## Example: Wheat



```
anova(lm(y~as.factor(g)))
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: y
```

```
##           Df Sum Sq Mean Sq F value Pr(>F)
```

```
## as.factor(g)  9 33.368   3.7076   2.0745 0.0555 .
```

```
## Residuals    40 71.488   1.7872
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## Example: Wheat

```
fit<-anova( lm(y~as.factor(g)) )  
  
MSA<-fit[1,3]  
MSW<-fit[2,3]  
  
MSA  
## [1] 3.70759  
  
MSW  
## [1] 1.787206  
  
t2<-(MSA-MSW)/n  
  
t2  
  
##          1  
## 0.3840768
```

## Example: Wheat

```
rho<-t2/(t2+MSW)

rho

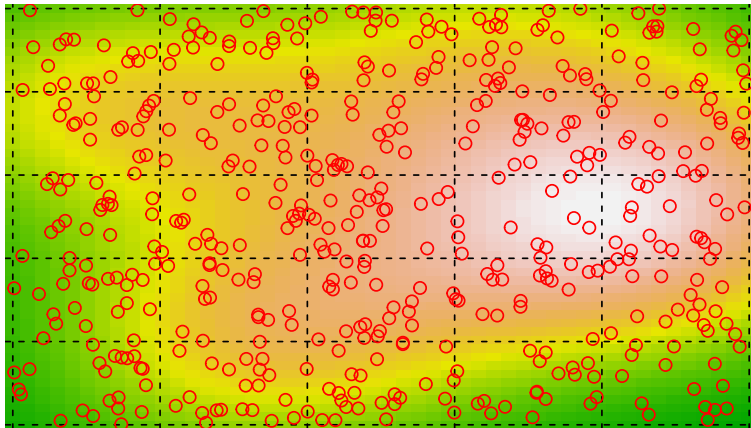
##           1
## 0.1768894

se.rho<- (1-rho)*(1+(n-1)*rho)*sqrt( 2/( n*(n-1)*(m-1)))

rho + c(-2,2)*se.rho

## [1] -0.1194179  0.4731966
```

## Two-stage sampling



$$\mu=2.1124814$$

## Ignoring across-group heterogeneity

**Task:** Construct a 95% CI for the population mean.

***t*-interval for SRS:**

If  $y_1, \dots, y_n$  is an iid sample with  $E[y_i] = \mu$  and  $\text{Var}[y_i] = \sigma^2$ ,

$$E[\bar{y}] = \mu, \text{Var}[\bar{y}] = \sigma^2/n.$$

By the central limit theorem,

$$\bar{y} \dot{\sim} N(\mu, \sigma^2/n), \quad \frac{\bar{y} - \mu}{\sigma/\sqrt{n}} \dot{\sim} N(0, 1).$$

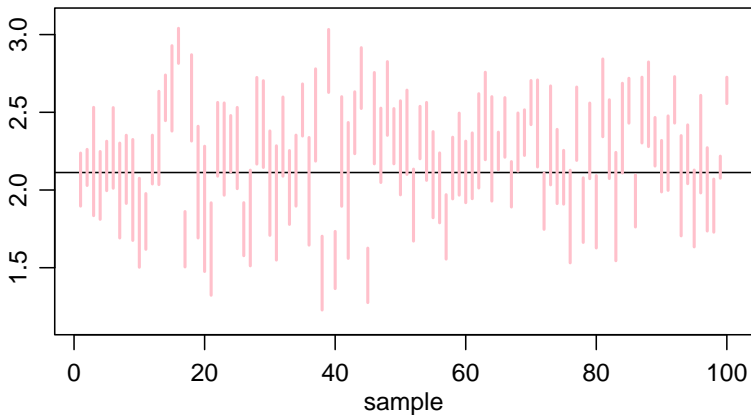
As  $\sigma^2$  is generally unknown, we use

$$\frac{\bar{y} - \mu}{s/\sqrt{n}} \dot{\sim} t_{n-1}, \quad \text{where } s^2 = \frac{1}{n-1} \sum (y_i - \bar{y})^2.$$

From this, we have

$$\bar{y} \pm t_{n-1, .975} \times s/\sqrt{n} \text{ is a 95\% CI for } \mu.$$

## Ignoring across-group heterogeneity



## Building an accurate $t$ -interval

Recall that an approximate 95% CI for  $\mu$  is given by

$$\bar{y} \pm 2 \times \text{se}(\bar{y}),$$

where  $\text{se}(\bar{y})$  is an approximation to the standard deviation of  $\bar{y}$ .

### How to find $\text{se}(\bar{y})$ :

1. compute the variance  $v$  of  $\bar{y}$  based on the model;
2. find an estimate  $\hat{v}$  of  $v$ ;
3. let  $\text{se}(\bar{y}) = \sqrt{\hat{v}}$ .

So the first step is to find  $\text{Var}[\bar{y}]$ :

## Variance of the grand mean around population mean

$$\begin{aligned}\text{Var}[\bar{y}] &= \text{Var}\left[\frac{1}{mn} \sum_j \sum_i y_{i,j}\right] \\ &= \text{Var}\left[\frac{1}{m} \sum_j \frac{1}{n} \sum_i y_{i,j}\right] \\ &= \text{Var}\left[\frac{1}{m} \sum_j \bar{y}_j\right] \\ &= \frac{1}{m^2} \text{Var}\left[\sum_j \bar{y}_j\right] \\ &= \frac{1}{m^2} \sum_j \text{Var}[\bar{y}_j] \\ &= \frac{1}{m^2} m \text{Var}[\bar{y}_1] \\ &= \frac{1}{m} \text{Var}[\bar{y}_1]\end{aligned}$$



## Variance of a group mean around population mean

What is  $\text{Var}[\bar{y}_1]$ ? We've shown

$$\text{Var}[y_{i,1}] = \sigma^2 + \tau^2,$$

but generally,

$$\text{Var}[\bar{y}_1] \neq [\sigma^2 + \tau^2]/n.$$

**Quiz:** What is the smallest that  $\text{Var}[\bar{y}_1]$  could be for fixed  $\sigma^2$  and  $n$ ? Recall

$$\text{Cor}[y_{i,1}, y_{i,2}] = \frac{\tau^2}{\tau^2 + \sigma^2}$$

**Answer:** When  $\tau^2$  is zero the within group samples are independent and so

$$\text{Var}[\bar{y}_1] \geq \sigma^2/n$$

## Variance of a group mean around population mean

**Quiz:** what is the smallest that  $\text{Var}[\bar{y}_1]$  could be for fixed  $\sigma^2$  and  $\tau^2$ ?

**Answer:** Increasing  $n$  reduces variation of  $\bar{y}_1$  around  $\theta_1$ , but across group heterogeneity remains:

for large  $n$ ,  $\bar{y}_1 \approx \theta_1$

$$\text{Var}[\theta_1] = \tau^2$$

$$\text{Var}[\bar{y}_1] \geq \tau^2$$

## Variance of a group mean around population mean

Let's compute  $\text{Var}[\bar{y}_1]$ . For notational convenience, we'll drop the group index, and assume  $\mu = 0$ , so

$$E[y_i] = 0, \quad E[y_i^2] = \sigma^2 + \tau^2, \quad E[y_i y_k] = \tau^2$$

In this case,

$$\begin{aligned}\text{Var}[\bar{y}] &= E[\bar{y}^2] \\ &= E\left[\frac{1}{n^2} \left(\sum y_i\right)^2\right] \\ &= \frac{1}{n^2} E\left[\sum y_i^2 + \sum_{i \neq k} y_i y_k\right] \\ &= \frac{1}{n^2} (n[\sigma^2 + \tau^2] + n(n-1)\tau^2) \\ &= \frac{\sigma^2}{n} + \frac{1}{n}\tau^2 + \frac{n-1}{n}\tau^2 \\ &= \frac{\sigma^2}{n} + \tau^2\end{aligned}$$

**Exercise:** Make sure the answer makes sense to you intuitively.

## Variance of the sample grand mean

$$\text{Var}[\bar{y}_{..}] = \frac{1}{m} \text{Var}[\bar{y}_j]$$

$$\text{Var}[\bar{y}_j] = \frac{1}{n} \sigma^2 + \tau^2$$

$$\text{Var}[\bar{y}_{..}] = \frac{1}{nm} \sigma^2 + \frac{1}{m} \tau^2$$

What happens as

- $n \rightarrow \infty$  and  $m$  stays fixed?
- $m \rightarrow \infty$  and  $n$  stays fixed?

In this sense,  $m$  is the “sample size” for the population-level parameter  $\mu$ .

## Standard error and CI

$$\widehat{\text{Var}}[\bar{y}_{..}] = \frac{1}{nm} \hat{\sigma}^2 + \frac{1}{m} \tau^2$$

- $\hat{\sigma}^2 = MSW$
- $\hat{\tau}^2 = (MSA - MSW)/n$

$$\widehat{\text{Var}}[\bar{y}_{..}] = \frac{1}{mn} MSA$$

This should make sense, because previously we claimed

$$E[MSA] = \sigma^2 + n \times \tau^2,$$

so

$$E\left[\frac{1}{mn} MSA\right] = \frac{1}{mn} \sigma^2 + \frac{1}{m} \tau^2 = \text{Var}[\bar{y}_{..}]$$

## Confidence interval

$$\bar{y}_{..} \pm 2 \times \sqrt{MSA/mn}$$

```
round(y,2)

## [1] 0.55 0.56 0.48 0.85 0.81 2.76 2.71 2.47 2.43 2.43 2.68 2.52 2.97 2.92 2.60
## [16] 2.42 1.90 1.99 2.37 1.87

g

## [1] 1 1 1 1 1 2 2 2 2 2 3 3 3 3 3 4 4 4 4 4

anova(lm(y~as.factor(g)))

## Analysis of Variance Table
##
## Response: y
##          Df Sum Sq Mean Sq F value    Pr(>F)
## as.factor(g)  3 13.4751   4.4917   110.5 6.603e-11 ***
## Residuals    16  0.6504   0.0406
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

MSA<-anova(lm(y~as.factor(g)))[1,3]

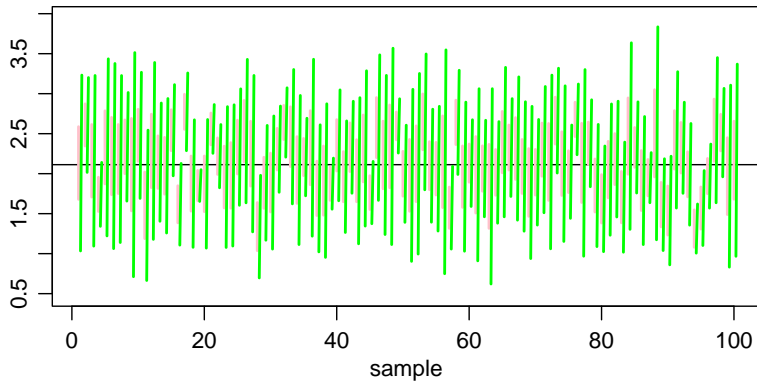
mean(y) + c(-2,2)*sqrt( MSA/(m*n) )

## [1] 1.066935 2.962551

mean(y) + c(-2,2)*sqrt( var(y)/(m*n) )

## [1] 1.629141 2.400345
```

## Accounting for across-group heterogeneity



```
mean( CI.tss0[,1] < mu & mu < CI.tss0[,2] )  
## [1] 0.729  
  
mean( CI.tss1[,1] < mu & mu < CI.tss1[,2] )  
## [1] 0.933
```

## Summary

$$y_{i,j} = \mu + a_j + \epsilon_{i,j}$$

$$\text{Var}[\epsilon_{i,j}] = \sigma^2$$

$$\text{Var}[a_j] = \tau^2$$

Variation around the group mean:  $\theta_j = \mu + a_j$

- $\text{Var}[y_{i,j}|\theta_j] = \sigma^2$
- $\text{Cov}[y_{i_1,j}, y_{i_2,j}|\theta_j] = 0$
- $\text{Exp} \bar{y}_j | \theta_j = \theta_j, \text{Var}[\bar{y}_j | \theta_j] = \sigma^2/n$

Variation around the grand mean:

- $\text{Var}[y_{i,j}|\mu] = \sigma^2 + \tau^2$
- $\text{Cov}[y_{i_1,j}, y_{i_2,j}|\mu] = \tau^2$
- $\text{E}[\bar{y}_j|\mu] = \mu, \text{Var}[\bar{y}_j|\mu] = \sigma^2/n + \tau^2$
- $\text{E}[\bar{y}_{..}|\mu] = \mu, \text{Var}[\bar{y}_{..}|\mu] = \sigma^2/(mn) + \tau^2/m$