

# Lecture 17

## Segment 2

Mixed factorial ANOVA:  
Example in R

# Example

## **Phonological similarity effects in simple and complex span tasks**

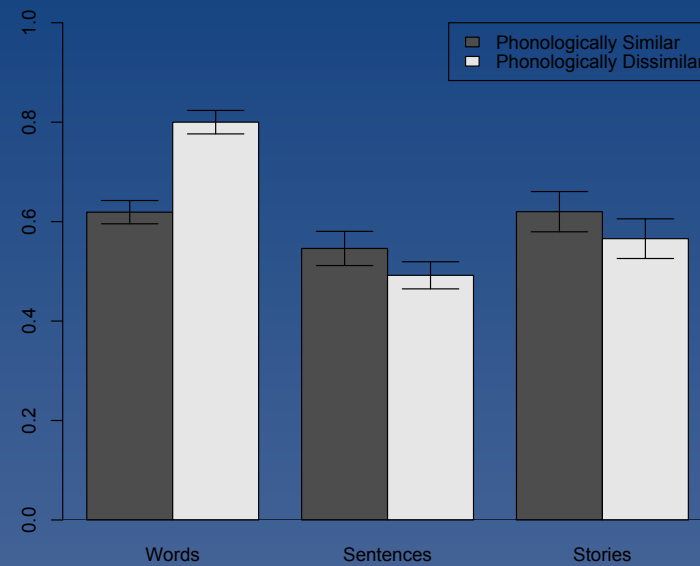
**Brooke N. Macnamara • Adam B. Moore •  
Andrew R. A. Conway**

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# Experimental design

- Mixed factorial design, 3x2, (Ax(BxS))
  - Between groups IV
    - Memory task (words, sentences, stories)
  - Within groups IV
    - Phonological similarity (similar, dissimilar)
  - DV
    - Percentage of words recalled

# Results ~ Preview



# Hypotheses

- Mixed factorial design,  $3 \times 2$ , ( $A \times (B \times S)$ )
  - Main effect of memory task?
    - Memory task (words, sentences, stories)
  - Main effect of similarity?
    - Phonological similarity (similar, dissimilar)
  - Interaction?

# Hypotheses

- Three F-ratios
  - $F_A$
  - $F_B$
  - $F_{A \times B}$

# R script

- This script is available on the course website
  - STATS1.EX.08.R
  - STATS1.EX.08.txt

# R script

```
#####  
# Macnamara, Moore, & Conway (2011), Experiment 1, Serial recall  
#####  
  
library(psych)  
library(car)  
source(file="eta_squared.R")  
  
e1sr <- read.table("STATS1.EX.08.txt", header = T)
```



# R script ~ Omnibus ANOVA

```
# Omnibus analysis is a 3x2 mixed factorial with task and stimuli as the independent variables and serial recall as the dependent variable.  
The three levels of task are word span, reading span, and story span. The two levels of stimuli are phonologically similar and  
phonologically dissimilar.
```

```
stim = factor(e1sr$stim, levels=c("S", "D")) #reverse levels (for graphs like the article)
```

```
aov.e1sr = aov(e1sr$recall ~ (e1sr$task*e1sr$stim) + Error(factor(e1sr$subject)/e1sr$stim))  
summary(aov.e1sr)  
eta.2(aov.e1sr, ret.labels=TRUE)
```

```
# Levene's test  
leveneTest(e1sr$recall, e1sr$task, center="mean")
```

# R script ~ Simple effects

```
# Simple effects analysis for simple span (i.e., word span)
aov.e1srw = aov(e1sr$recall[task=="W"] ~ e1sr$stim[task=="W"] +Error(factor(e1sr$subject[task=="W"])/e1sr$stim[task=="W"]))
summary(aov.e1srw)
eta.2(aov.e1srw, ret.labels=TRUE)
```

# R script ~ Simple effects

```
# Simple effects analysis for complex span (this is a 2x2 mixed factorial)
aov.e1srnw = aov(e1sr$recall[task!="W"] ~ e1sr$task[task!="W"]*e1sr$stim[task!="W"] +
  Error(factor(e1sr$subject[task!="W"]) / e1sr$stim[task!="W"]))
summary(aov.e1srnw)
eta.2(aov.e1srnw, ret.labels=TRUE)
```

# R script ~ Bar chart

```
# Bar plot

wspan = describe.by(recall[task=="W"], group = stim[task=="W"], mat = T)
rspan = describe.by(recall[task=="R"], group = stim[task=="R"], mat = T)
sspan = describe.by(recall[task=="S"], group = stim[task=="S"], mat = T)
graphme = cbind(Words = wspan$mean, Sentences = rspan$mean, Stories = sspan$mean)
rownames(graphme) = c("Phonologically Similar", "Phonologically Dissimilar")
se = cbind(wspan$se, rspan$se, sspan$se)

bp = barplot(graphme, beside = TRUE,
             ylim = c(0,1), space = c(0, .5), legend.text = TRUE,
             args.legend = c(x = "topright"))
abline(h=0)
for (ii in 1:3) {
  arrows(bp[1, ii], graphme[1,ii] - se[1, ii],
        y1 = graphme[1,ii] + se[1, ii], angle = 90, code = 3)
  arrows(bp[2, ii], graphme[2,ii] - se[2, ii],
        y1 = graphme[2,ii] + se[2, ii], angle = 90, code = 3)
}
```

# R script ~ Effect size function

```
eta.2 = function(aov.mdl, ret.labels = FALSE){  
  eta.2vector = c()  
  labels = c()  
  for (table in summary(aov.mdl)){ #each block of factors  
    SS.vector = table[[1]]$"Sum Sq" #table is a list with 1 entry, but you have to use [[1]] anyway  
    last = length(SS.vector)  
    labels = c(labels, row.names(table[[1]])[-last]) #all but last (error term)  
    for (SS in SS.vector[-last]) { #all but last entry (error term)  
      new.etaval = SS / (SS + SS.vector[last])  
      eta.2vector = c(eta.2vector, new.etaval)  
    }  
  }  
  if (ret.labels) return(data.frame(eta.2 = eta.2vector, row.names = labels))  
  return(eta.2vector)  
}
```

# R output ~ Omnibus analysis

```
Error: factor(e1sr$subject)
      Df Sum Sq Mean Sq F value    Pr(>F)
e1sr$task  2  0.7384   0.3692   10.54 0.000124 ***
Residuals 58  2.0309   0.0350
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Error: factor(e1sr$subject):e1sr$stim
      Df Sum Sq Mean Sq F value    Pr(>F)
e1sr$stim  1  0.0161  0.01607   1.956   0.167
e1sr$task:e1sr$stim  2  0.3716  0.18582  22.624 5.46e-08 ***
Residuals      58  0.4764  0.00821
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> eta.2(aov.e1sr, ret.labels=TRUE)
      eta.2
e1sr$task  0.26663954
e1sr$stim  0.03262359
e1sr$task:e1sr$stim 0.43824551
```

# R output ~ Levene's test

```
>
> # Levene's test
> leveneTest(e1sr$recall, e1sr$task, center="mean")
Levene's Test for Homogeneity of Variance (center = "mean")
      Df F value  Pr(>F)
group   2  3.4274 0.03571 *
      119
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
>
```

# R output ~ Simple effects

```
Error: factor(subject[task == "W"])
      Df Sum Sq Mean Sq F value Pr(>F)
Residuals 19 0.3404 0.01792

Error: factor(subject[task == "W"]):stim[task == "W"]
      Df Sum Sq Mean Sq F value    Pr(>F)
stim[task == "W"]  1 0.3276  0.3276    78.8 3.46e-08 ***
Residuals          19 0.0790  0.0042
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> eta.2(aov.e1srw, ret.labels=TRUE)
      eta.2
stim[task == "W"] 0.8057304
```



# R output ~ Simple effects

```
Error: factor(subject[task != "W"])
              Df Sum Sq Mean Sq F value Pr(>F)
task[task != "W"]  1 0.1118 0.11176   2.578  0.116
Residuals        39 1.6905 0.04335

Error: factor(subject[task != "W"]):stim[task != "W"]
              Df Sum Sq Mean Sq F value Pr(>F)
stim[task != "W"]      1 0.0601 0.06010   5.898 0.0199 *
task[task != "W"]:stim[task != "W"]  1 0.0000 0.00000   0.000 0.9949
Residuals            39 0.3974 0.01019

---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> eta.2(aov.e1srnw, ret.labels=TRUE)
              eta.2
task[task != "W"]  6.200964e-02
stim[task != "W"]  1.313716e-01
task[task != "W"]:stim[task != "W"] 1.052142e-06
```

# Results

