

One Way Within-participant or Repeated-measures Analysis of Variance, Balanced Designs

For Psychology 610, University of Wisconsin--Madison.

This tutorial uses data in Table 16.3, Keppel & Wickens, p. 355.

Install the package "car"

Contents of this handout:

- I. Basic data setup and analysis, including sphericity and H-F adjusted p-values, means and estimated standard errors
- II. Contrasts with partitioned error
- III. Line graphs and bar graph with error bars for one-way within design. Note: you need to calculate the estimated standard error from the residual of the anova model.

'Quick Look' Summary of R code for One-way within:

```
> library(car)
> multmodel=lm(cbind(A1,A2,A3) ~ 1) # make a multivariate linear model with
only the intercept as a predictor for your within-participants observations
> Trials=factor(c("A1","A2","A3"), ordered=F) # create a factor for your repeated-
measures variable.
> model1=Anova(multmodel, idata=data.frame(Trials), idesign=~Trials, type="III") #
use the repeated-measures factor as the 'internal' part of the design using 'Anova'
(with a capital A)
```

I. One-way within-subjects anova -- basic data setup and analysis.

A. Start R and bring in the data.

```
> library(car) # bring the 'car' package into the environment
> options(contrasts=c("contr.sum","contr.poly")) # set options for contrasts. Not necessary here, but a good practice if
you will be analyzing any unbalanced designs
> your.data=read.table(pipe("pbpaste"),header=T) # I copied the data to the clipboard from the excel sheet
> your.data
  A1  A2  A3 subj
1 745 764 774 s1
2 777 786 788 s2
3 734 733 763 s3
4 779 801 797 s4
5 756 786 785 s5
6 721 732 740 s6

> attach(your.data) # some warn against attaching data.
```

B. Carry out the analysis using 'Anova' in the 'car' package in two steps.

By using the ‘car’ package we obtain the sphericity test and p-values adjusted by both the Huynh-Feldt or Greenhouse-Geisser method.

Step 1. Use ‘lm’ on all the repeated-measures variables together

```
> multmodel=lm(cbind(A1,A2,A3) ~ 1) # we column bind the 3 response columns together, and use only the intercept
as our predictor variable.
```

Step 2. Construct the repeated measures variable and finish the analysis using ‘Anova’ (capital A) in the ‘car’ package.

```
> Trials=factor(c("A1","A2","A3"), ordered=F) # create a factor called "Trials" with labels A1 to A3.
```

```
> Trials
[1] A1 A2 A3
Levels: A1 A2 A3
> model1=Anova(multmodel, idata=data.frame(Trials), idesign=~Trials, type="III") # We name the model we
established in Step 1. Then the ‘idata’ parameter is for the repeated-measures part of the data, the ‘idesign’ is where
you specify the repeated part of the design. This is a one-way within design, so we have only the ‘Trials’ variable.
> summary(model1, multivariate=F) # omit the ‘multivariate=F’ part, and you will get the MANOVA results as well.
```

```
Univariate Type III Repeated-Measures ANOVA Assuming Sphericity

              SS num Df Error SS den Df          F      Pr(>F)
(Intercept) 10520285     1    8548     5 6153.773 6.378e-09 ***
Trials        1575      2     546    10  14.432  0.001128 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Mauchly Tests for Sphericity

```
          Test statistic p-value
Trials      0.75802 0.57459
```

Greenhouse-Geisser and Huynh-Feldt Corrections
for Departure from Sphericity

```
          GG eps Pr(>F[GG])
Trials 0.80516  0.002860 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
          HF eps Pr(>F[HF])
Trials 1.1302  0.001128 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Warning message:
In summary.Anova.mlm(model1, multivariate = F) : HF eps > 1 treated as 1
```

***** Check that the df's are correct!** ‘Trials’ has 3 levels, and should have 2 df. The error for Trials should be Trials x Subjects interaction, $df = (t-1)(n-1)$ in Keppel’s notation. We have 6 people, so df error should be 10.

II. Contrasts with partitioned error.

A. Pairwise test of means, without post-hoc alpha-adjustment

This is easiest to do with the t-test function.

```
> c1=t.test(A1,A2,alternative="two.sided",mu=0,paired=T) # A1 and A2 are the variable names to test. We say  
'paired=True' so that R will do a paired t-test because this is a within-S design.
```

```
> c1  
  
Paired t-test  
  
data: A1 and A2  
t = -3.3597, df = 5, p-value = 0.02011  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 -26.476815 -3.523185  
sample estimates:  
mean of the differences  
 -15
```

```
> c2=t.test(A1,A3,alternative="two.sided",mu=0,paired=T)
```

```
> c2  
  
Paired t-test  
  
data: A1 and A3  
t = -7.2181, df = 5, p-value = 0.0007958  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 -30.51291 -14.48709  
sample estimates:  
mean of the differences  
 -22.5
```

```
> c3=t.test(A2,A3,alternative="two.sided",mu=0,paired=T)
```

```
> c3  
  
Paired t-test  
  
data: A2 and A3  
t = -1.5025, df = 5, p-value = 0.1933  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 -20.33147 5.33147  
sample estimates:  
mean of the differences  
 -7.5
```

B. Adjust the p-values using Holm.

```
> pvec=c(0.02011, 0.0007958, 0.1933) # make a vector of p-values from the pairwise tests  
> p.adjust(pvec,method="holm",n=3) # adjust p's by Holm method. Both A1 vs A2 and A1 vs A3 remain significant  
[1] 0.0402200 0.0023874 0.1933000
```

C. Calculate partitioned error contrasts from the data matrix, and then test H_0 that ψ equals zero with ‘aov’ or with a t-test. However, we have headings on our data, so we have to bind the data together column-wise (‘cbind’) and then operate on that.

1. Repeat one of the pairwise contrasts we did above and make sure it matches the t-test results.

```
> contr1=c(1,-1) # make a vector of contrast values
```

```
> contr1
[1] 1 -1
```

```
> xc1=cbind(A1,A2)%*% contr1 # ‘%*%’ is the symbol for matrix multiplication. So this multiplies A1 by 1 and A2 by -1, and puts the result in the vector that I named ‘xc1’. ‘xc1’ is just a table of psi-hats for each individual. Once we have that, we can test it with a t-test or by anova of the grand mean.
```

```
> xc1
```

```
      [,1]
[1,] -19
[2,]  -9
[3,]   1
[4,] -22
[5,] -30
[6,] -11
```

```
> aov1=aov(xc1~1) # ask R to do the anova. The ‘~1’ says to use only the intercept
```

```
> summary(aov1,intercept=T) # ask for the anova with intercept
```

```
      Df Sum Sq Mean Sq F value Pr(>F)
(Intercept)  1 1350.0  1350.0  11.288 0.02011 *
Residuals    5  598.0   119.6
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1
```

Does this agree with the t-test we did above? $F = t\text{-squared}$, or $t = \sqrt{F}$. Here’s a hand calculation to verify:

```
> sqrt(11.288)
```

```
[1] 3.359762
```

2. A more complex contrast.

```
> contr4=c(.5,.5,-1) # test average of A1 & A2 vs A3. First make contrast vector
```

```
> contr4
[1] 0.5 0.5 -1.0
```

```
> xc4 = cbind(A1,A2,A3)%*% contr4 # multiply contrast vector by a matrix made up of A1,A2,A3. Put the results in a vector called ‘xc4’.
```

```
> xc4 # here is the vector of psi-hats
```

```
      [,1]
[1,] -19.5
[2,]  -6.5
[3,] -29.5
[4,]  -7.0
[5,] -14.0
[6,] -13.5
```

```
> summary (aov(xc4 ~ 1), intercept=T) # do the anova & get summary table in one step
              Df Sum Sq Mean Sq F value Pr(>F)
(Intercept)  1   1350    1350  18.243 0.00793 **
Residuals    5    370     74
```

Show the mean of the psi-hats. We put the psi-hats are in the vector called 'xc4'.

```
> mean(xc4) # 'mean' is a built-in function
[1] -15
> sd(xc4) # sd is also built-in
[1] 8.602325
```

Square this sd and it equals MS-residual in the test of the intercept above.

Can also do a t-test on the vector of psi-hats:

```
> t.test(xc4)

One Sample t-test

data: xc4
t = -4.2712, df = 5, p-value = 0.00793
alternative hypothesis: true mean is not equal to 0
95 percent confidence interval:
 -24.027587 -5.972413
sample estimates:
mean of x
      -15
```

III Graphs in a repeated measures design.

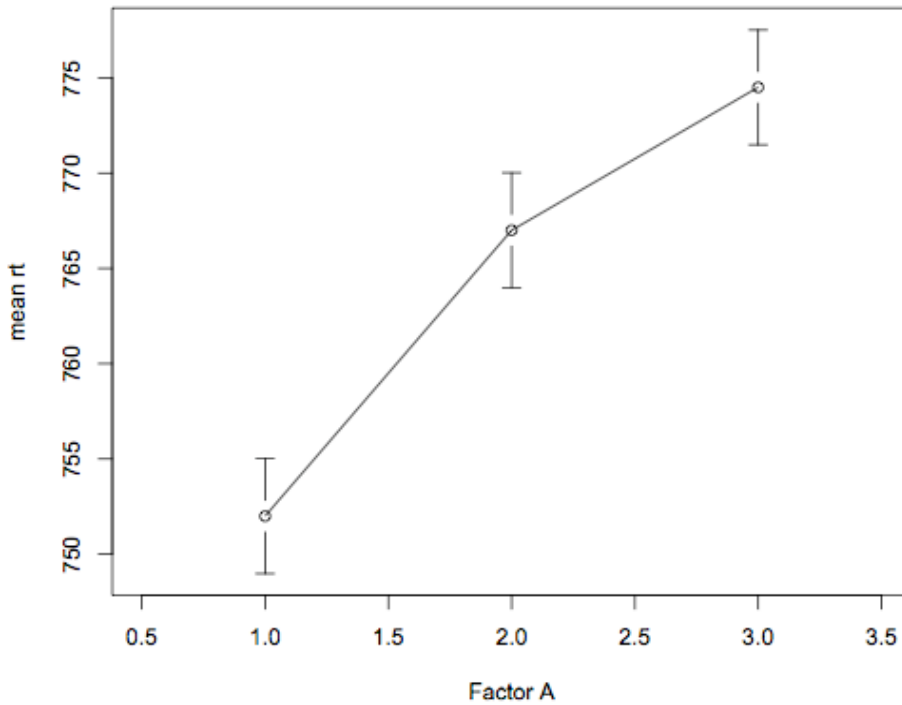
The functions in 'sciplot' (lineplot.CI, and barplot.CI) requires you to stack your data, and then calculate the standard errors as though it is a between-group design. We can easily build the bar plot by hand, or we can use the 'gplots' package to make a line graph with the standard error bars.

A. Line graph with error bars, using 'gplots' package.

```
> means=c(mean(A1),mean(A2),mean(A3)) # make a vector of means called 'means'.
> means
[1] 752.0 767.0 774.5
> se=sqrt(54.6 / 6); se # make a variable called 'se', and use it to calculate estimated standard error. I used the MS
error from the overall anova.
[1] 3.016621

> library (gplots)
> plotCI(x = means, uiw = se, ylab="mean rt", type="l", xlab="Factor A", main="Keppel Table 16.3",xlim=c(.5,3.5)) #
After looking at the first graph, I changed 'xlim' to move the bars in off the edges of the graph. I am not sure how to
get "plotCI" to treat the x-axis as categorical
```

Keppel Table 16.3



B. Bar plot with error bars.

We calculated the means and standard errors above. Set up the 'superpose' function (from the website of Raoul Grasman: <http://users.fmg.uva.nl/rgrasman/rpages/2005/09/error-bars-in-plots.html>

```
> superpose.eb = function (x, y, ebl, ebu = ebl, length = 0.08, ...) arrows(x, y + ebu, x, y - ebl, angle = 90, code = 3,  
  length = length, ...)
```

```
> Keppelbars = barplot(means, beside=T, ylim=c(750,780), space=c(.1,.8), main="Keppel Table 16.3", xlab="Level of  
A", ylab="mean reaction time", legend=T, axis.lty=1, xpd=F) # 'xpd = F' tells R not to plot outside of the x and y  
limits you set. 'means' is our vector of data.
```

```
> superpose.eb(x=Keppelbars, y=means, ebl=c(se,se,se), col="black", lwd=1) # superpose wants a value for each error  
bar. For this design our estimated standard errors are model-based, so they are identical across the 3 levels of factor A.  
'se' contains our estimated standard error, so we make a vector that repeats it 3 times.
```

```
> box() # add a box around the graph
```

```
> axis(4,labels=F) # add tick marks on the right-hand side
```

Keppel Table 16.3

