

# Lab 4 – Contrasts, Estimation, and Power Analysis

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FANR 6750

Richard Chandler and Bob Cooper  
University of Georgia

① CONTRASTS

② ESTIMATION

③ POWER

## CHAIN SAW DATA

```
sawData <- read.csv("sawData.csv")
sawData
```

```
##      y Brand
## 1  42    A
## 2  17    A
## 3  24    A
## 4  39    A
## 5  43    A
## 6  28    B
## 7  50    B
## 8  44    B
## 9  32    B
## 10 61    B
## 11 57    C
## 12 45    C
## 13 48    C
## 14 41    C
## 15 54    C
## 16 29    D
## 17 40    D
## 18 22    D
## 19 34    D
## 20 30    D
```

## CONTRASTS

Suppose we want to make 3 *a priori* comparisons:

- (1) Groups A&D vs B&C
- (2) Groups A vs D
- (3) Groups B vs C

	Comparison	Null hypothesis
1	AD vs BC	$\frac{\mu_A + \mu_D}{2} - \frac{\mu_B + \mu_C}{2} = 0$
2	A vs D	$\mu_A - \mu_D = 0$
3	B vs C	$\mu_B - \mu_C = 0$

## Coefficients

```
ADvBC <- c(1/2, -1/2, -1/2, 1/2)
AvD <- c(1, 0, 0, -1)
BvC <- c(0, 1, -1, 0)
```

## Are they orthogonal?

```
sum(ADvBC)
```

```
## [1] 0
```

```
sum(AvD)
```

```
## [1] 0
```

```
sum(BvC)
```

```
## [1] 0
```

```
sum(ADvBC * AvD)
```

```
## [1] 0
```

```
sum(ADvBC * BvC)
```

```
## [1] 0
```

```
sum(AvD * BvC)
```

```
## [1] 0
```

Yes, they are.

To use contrasts in R, each set of coefficients must be formatted as a column in a matrix.

We can use `cbind` for this:

```
contrast.mat <- cbind(ADvBC, AvD, BvC)
contrast.mat
```

```
##      ADvBC AvD BvC
## [1,]  0.5  1  0
## [2,] -0.5  0  1
## [3,] -0.5  0 -1
## [4,]  0.5 -1  0
```

## CONTRASTS

## Fit the model with contrasts

```
aov.out <- aov(y ~ Brand, data=sawData,
              contrasts=list(Brand=contrast.mat))
```

## Now “split” apart the sum-of-squares

```
summary(aov.out, split = list(Brand =
                             list("ADvBC"=1, "AvD"=2, "BvC"=3)))
```

```
##              Df Sum Sq Mean Sq F value Pr(>F)
## Brand          3  1080   360.0   3.556 0.03823 *
## Brand: ADvBC   1    980   980.0   9.679 0.00672 **
## Brand: AvD     1     10    10.0   0.099 0.75738
## Brand: BvC     1     90    90.0   0.889 0.35980
## Residuals     16  1620   101.2
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

## DIFFERENCE IN MEANS FOR EACH CONTRAST

## Group means

```
(group.means <- tapply(sawData$y, sawData$Brand, mean))
```

```
##  A  B  C  D
## 33 43 49 31
```

## Difference in means for A vs D

```
group.means <- unname(group.means) # Drop names (optional)
group.means[1] - group.means[4]
```

```
## [1] 2
```

## Difference in means for B vs C

```
group.means[2] - group.means[3]
```

```
## [1] -6
```

## Difference in means for AD vs BC

```
mean(group.means[c(1,4)]) - mean(group.means[2:3])
```

```
## [1] -14
```

## SE for A vs D

```
se.contrast(aov.out, list(sawData$Brand=="A",
                          sawData$Brand=="D"))
```

```
## [1] 6.363961
```

## SE for B vs C

```
se.contrast(aov.out, list(sawData$Brand=="B",
                          sawData$Brand=="C"))
```

```
## [1] 6.363961
```

## SE for AD vs BC

```
se.contrast(aov.out, list(sawData$Brand=="A" |
                          sawData$Brand=="D",
                          sawData$Brand=="B" |
                          sawData$Brand=="C"))
```

```
## [1] 4.5
```

In an ANOVA context, confidence intervals can be constructed using the equation:

$$CI = \text{Point estimate} \pm t_{\alpha/2, a(n-1)} \times SE$$

As usual, the hard part is computing the SE<sup>1</sup>

<sup>1</sup>See page 300 of Dowdy et al. for SE formulas

## 1 CONTRASTS

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SE's for the effect sizes ( $\alpha$ 's)

```
effects.SE <- model.tables(aov.out, type="effects",
                          se=TRUE)
```

```
effects.SE
```

```
## Tables of effects
```

```
##
```

```
## Brand
```

```
## Brand
```

```
## A B C D
```

```
## -6 4 10 -8
```

```
##
```

```
## Standard errors of effects
```

```
## Brand
```

```
## 4.5
```

```
## replic. 5
```

## Extract the $\alpha$ 's and the SEs

```
# str(effects.SE)
alpha.i <- as.numeric(effects.SE$tables$Brand)
SE <- as.numeric(effects.SE$se)
```

## Compute confidence intervals

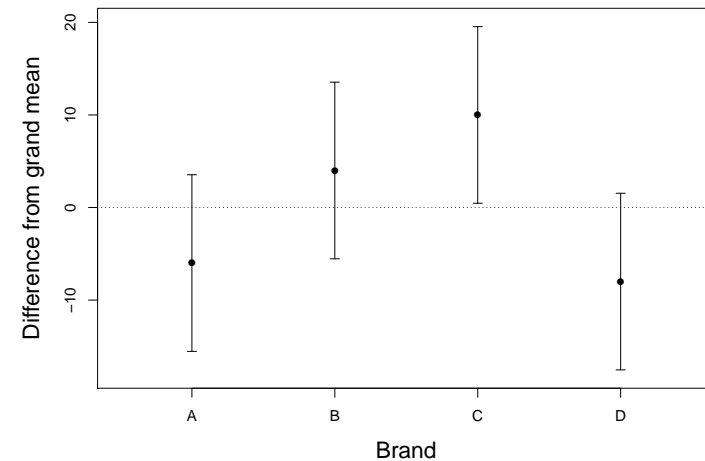
```
tc <- qt(0.975, 4*(5-1))
lowerCI <- alpha.i - tc * SE
upperCI <- alpha.i + tc * SE
```

## Put results into a data.frame

```
CI <- data.frame(effect.size=alpha.i, SE,
                 lowerCI, upperCI)
round(CI, 2)
```

```
##   effect.size  SE lowerCI upperCI
## 1         -6 4.5  -15.54    3.54
## 2          4 4.5   -5.54   13.54
## 3         10 4.5    0.46   19.54
## 4         -8 4.5  -17.54    1.54
```

```
plot(1:4, CI$effect.size, xlim=c(0.5, 4.5), ylim=c(-18, 20), xaxt="n",
     xlab="Brand", ylab="Difference from grand mean", pch=16, cex.lab=1.5)
axis(1, at=1:4, labels=c("A", "B", "C", "D"))
abline(h=0, lty=3)
arrows(1:4, CI$lowerCI, 1:4, CI$upperCI, code=3, angle=90, length=0.05)
```



1 CONTRASTS

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```
power.t.test(n=NULL, delta=3, sd=2, sig.level=0.05,
             power=0.8)
```

```
##
##   Two-sample t test power calculation
##
##           n = 8.06031
##          delta = 3
##           sd = 2
##   sig.level = 0.05
##          power = 0.8
##   alternative = two.sided
##
## NOTE: n is number in *each* group
```

```
power.anova.test(groups=4, n=5, between.var=360.0,
                 within.var=101.2, power=NULL)

##
##      Balanced one-way analysis of variance power calculation
##
##      groups = 4
##      n = 5
##      between.var = 360
##      within.var = 101.2
##      sig.level = 0.05
##      power = 0.9999359
##
## NOTE: n is number in each group
```

Researchers wish to know if food supplementation affects the growth of nestling Canada warblers. The treatment groups are: (A) No supplementation control, (B) low, (C) medium, (D) high, and (E) very high. The response variable is the weight of a 10 day old nestling.

- (1) The researchers are interested in the following contrasts. Are they orthogonal?
  - ▶ Groups A,B vs C,D,E
  - ▶ Groups A vs B
  - ▶ Groups C vs D,E
  - ▶ Groups D vs E
- (2) Using the warblerWeight data, test the null hypothesis of each contrast by constructing an ANOVA table in **R**.
- (3) For each contrast:
  - ▶ Compute the difference in the means
  - ▶ The SE of the difference in means
  - ▶ The 95% CI for the difference in means
- (4) Suppose you wanted to replicate the study with a smaller sample size of  $n = 2$  per treatment group? What would be your power?

Submit your answers in a self-contained script before the next lab.