2/25/2010 36-402/608 ADA-II Breakout #13 Results H. Seltman

Remember the bran study with 20 subjects given three diets (baseline, hi-fiber, and lofiber) in random order to see how they affect cholesterol. (We saw that order appeared to have no effect, and will not use it as a variable.) One way to express the scientific hypotheses is that we are simultaneously interested in testing $\mu_B = (\mu_L + \mu_H)/2$ and $\mu_H = \mu_L$. Although we could do two separate paired t-tests, we will do a single overall one-sample test of the two hypotheses.

```
BHL = bran$hifiber-bran$lofiber
BF = (bran$lofiber+bran$hifiber)/2-bran$baseline
means = c(mean(BF),mean(BHL))
n = nrow(bran)
T2 = n * as.numeric(means %*% solve(cov(cbind(BF,BHL))) %*% means)
T2 # 19.70
F = (n-2)/2/(n-1)*T2
F # 9.33
1-pf(F, 2, n-2) # 0.00166
```

Question 1: How does the code relate to the formulas in the handout? What conclusion do you reach? What followup testing should be done?

Recall the flea beetle study in which two different measurements are taken on two similar species of beetles. The question of interest is whether the collection of p=2 measurements are a distinguishing feature between the species (though not necessarily useful for distinguishing individuals).

```
> anova(aov(beet[,1:2]~species, data=beet), test="Hotelling")
# Error in model.frame.default(formula = beet[, 1:2] ~ species, data = beet, :
    invalid type (list) for variable 'beet[, 1:2]'
#
anova(aov(as.matrix(beet[,1:2])~species, data=beet), test="Hotelling")
# Analysis of Variance Table
#
              Df Hotelling-Lawley approx F num Df den Df
                                                             Pr(>F)
# (Intercept)
                           344.15
                                     5678.4
                                                 2
                                                       33 < 2.2e-16 ***
               1
# species
               1
                             4.81
                                       79.4
                                                 2
                                                       33 2.455e-13 ***
# Residuals
              34
```

Question 2: Why did the first attempt fail? How does the code relate to the formulas in the handout? How does anova() "know" to do MANOVA? What conclusion do you reach? How can you get a small p-value and then perhaps find that these measurements are not very useful for categorizing individual beetles?

Recall the monkeys being tested for short and long term memory with and without brain surgery on the hippocampus.

```
mem$SL = with(mem,cbind(short=(week2+week4)/2,
                         long=(week8+week12+week16)/3))
anova(aov(SL<sup>treatment, data=mem), test="Hotelling")</sup>
# Analysis of Variance Table
#
              Df Hotelling-Lawley approx F num Df den Df
                                                               Pr(>F)
                           250.795
                                    1880.96
                                                  2
                                                         15 < 2.2e-16 ***
# (Intercept)
               1
# treatment
                             1.643
                                       12.32
                                                  2
                                                         15 0.0006831 ***
               1
# Residuals
              16
```

Question 3: What is the first statement doing? What are the values of p and K? What conclusion do you reach? What could you do to check assumptions?