STAT 425 Assignment 7

Due Tuesday, May 4, 11:59 pm. Submit through Moodle.

Name: (insert your name here)

Netid: (insert)

Submit your computational work both as an R markdown (*.Rmd) document and as a pdf, along with any files needed to run the code. Embed your answers to each problem in the document below after the question statement. If you have hand-written work, please scan or take pictures of it and include in a pdf file, ideally combined with your pdf output file from R Markdown.

Most relevant class notes: 10.1.TwoWayAnova1, 10.2.TwoWayAnova2, 11.1.ExperimentalDesign1, 11.2.ExperimentalDesign2

Problem 1

The data set ctsib in faraway comes from an experiment described in the documentation as follows:

An experiment was conducted to study the effects of surface and vision on balance. The balance of subjects were observed for two different surfaces and for restricted and unrestricted vision. Balance was assessed qualitatively on an ordinal four-point scale based on observation by the experimenter. Forty subjects were studied ... while standing on foam or a normal surface and with their eyes closed or open or with a dome placed over their head. Each subject was tested twice in each of the surface and eye combinations for a total of 12 measures per subject.

In this problem, we will ignore the subject effects, and perform the two-way ANOVA analysis of the response score CTSIB as a function of Surface and Vision as if the experiment had completely randomized design. In Problem 2, for comparison, we will perform the more proper analysis that accounts for the subject as a blocking variable.

a) Here is a way to count how many observations fall in each combination of the two treatment factors:

```
library(faraway)
xtabs(~Surface+Vision, data=ctsib)
```

```
## Vision
## Surface closed dome open
```

```
## foam 80 80 80
## norm 80 80 80
```

Because the design is balanced across all 6 factor combinations, we can do the balanced two-way ANOVA analysis. Here is the ANOVA table for the model with main effects and interactions.

```
modCRD = lm(CTSIB~Surface*Vision, data=ctsib)
summary(aov(modCRD))
```

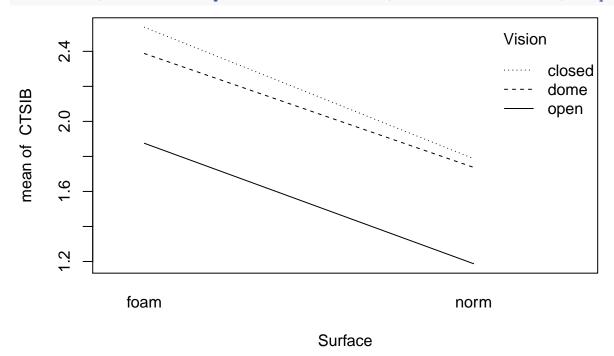
```
##
                   Df Sum Sq Mean Sq F value Pr(>F)
## Surface
                        58.10
                                58.10 297.132 <2e-16 ***
## Vision
                     2
                        36.84
                                18.42
                                       94.193 <2e-16 ***
                    2
                         0.20
                                        0.522 0.594
## Surface: Vision
                                 0.10
## Residuals
                  474
                        92.69
                                 0.20
## ---
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
```

Are there significant interactions? Explain.

Answer:

b) Here is how we can get one of the two possible interaction plots for these data.

with(ctsib, interaction.plot(x.factor=Surface, trace.factor=Vision, response=CTSIB))



Provide the other interaction plot, with Vision on the x-axis, and describe what the plots suggest about how the mean balance scores vary across the different combinations of Vision and Surface.

Answer:

c) Fit the additive model (main effects only for Surface and Vision), and show the corresponding analysis of variance table. Are the main effects significant?

Answer:

d) Display the additive model summary that includes the coefficient estimates and standard errors. What are the reference categories for Surface and Vision?

Answer:

e) Perform the Tukey honest significant difference analysis of the two treatment factors, based on the additive model. Which Surface and Vision level differences are statistically significant, controlling the family-wise false postive rate at 0.05?

Answer:

Problem 2

The Problem 1 analysis of the ctsib data is technically incorrect, because we expect the 12 measurements from each subject to be more closely related (correlated) than observations between subjects. In other words, Subject is a blocking variable. The subject id is a number, but we should treat it as a factor (categorical label) variable. Therefore, in this experiment we have 40 blocks (the subjects), and 2 observations for each of the 6 treatment combinations within each block. Here are the numbers of observations per cell for the first 3 subjects (blocks), so you can see the design:

```
xtabs(~Surface+Vision+as.factor(Subject), data=ctsib)[,,1:3]
```

```
, , as.factor(Subject) = 1
##
##
           Vision
## Surface closed dome open
##
      foam
                  2
                       2
                 2
                       2
                             2
##
      norm
##
   , , as.factor(Subject) = 2
##
##
##
           Vision
## Surface closed dome open
##
      foam
                  2
                       2
                 2
                       2
##
                             2
      norm
##
   , , as.factor(Subject) = 3
##
##
##
           Vision
## Surface closed dome open
##
      foam
                  2
                       2
                             2
                 2
                       2
                             2
##
      norm
```

a) Here is the ANOVA table for the model that adds the blocking variable as a main effect in addition to the main effects and interactions between the two treatment factors:

```
modRCBD = lm(CTSIB~as.factor(Subject)+Surface*Vision, data=ctsib)
summary(aov(modRCBD))
```

```
##
                        Df Sum Sq Mean Sq F value Pr(>F)
                        39
                            38.75
                                      0.99
                                             8.012 <2e-16 ***
## as.factor(Subject)
## Surface
                            58.10
                                     58.10 468.569 <2e-16 ***
                         1
## Vision
                         2
                                     18.42 148.539 <2e-16 ***
                            36.84
## Surface: Vision
                         2
                             0.20
                                      0.10
                                             0.823
                                                      0.44
## Residuals
                       435
                            53.94
                                      0.12
## ---
                    0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
```

Are there significant interactions? Explain.

Answer:

b) Comparing the ANOVA table part 2a) with the Anova table in Problem 1a), which sums of squares have changed and which ones remain the same? Same question for the different F values, how did each of them change (larger, smaller, unchanged)?

Answer:

c) Fit the additive model (main effects only for as.factor(Subject), Surface and Vision), and show the corresponding analysis of variance table. Are the main effects significant? How have the F values of the main effects for Vision and Surface changed compared with the results in Problem 1c)?

Answer:

d) Compute the ratio of the residual standard error $\hat{\sigma}$ for the model in Part 1c to the residual standard error for the model in Part 2c. What does this say about the relative efficiency of the completely randomized design compared with the randomized blocks design?

Answer:

e) Using the additive model that included the subject as a blocking variable, redo the Tukey honest significant difference analysis of the two treatment factors, Surface and Vision. Which Surface and Vision level differences are statistically significant, controlling the family-wise false postive rate at 0.05? Hint: if using the TukeyHSD function, include the argument which=c("Surface", "Vision") to obtain results for those factors only and not show the 40*39/2=780 blocking variable differences, which are not of interest.

Answer:

Problem 3

The alfalfa data in the faraway library are from an experiment to study yield from planted alfalfa seeds give different inoculum treatments. The experiment compared five treatments, A-E, where E was the control level. Two blocking variables were used: 1) shade, which is the distance of the location in the field from a tree line divided into 5 shade areas (1-5); and 2) irrigation, which is an irrigation measure divided into 5 levels. The experiment used an incomplete design such that each of the 5 treatments was given once for each level of shade and each level of irrigation, for a total of 25 observations.

a) To understand which combinations of variable levels were used in the experiment, display the result of running the cross-tabulation command

xtabs(yield
$$\sim$$
 inoculum + shade + irrigation, data=alfalfa)

as illustrated in the class notes on experimental design. Note that an entry in the cross-tab is 0 if that combination did not occur, and the entry is the response value if that combination did occur. Based on the information in the data write down the form of the 5×5 latin square used in this design. Rows should be levels of irrigation, columns should be levels of shade, and the entries in the table should be treatment labels.

Answer:

Block	Shade1	Shade2	Shade3	Shade4	Shade5
Irrig1	?	?	?	?	?
Irrig2	?	?	?	?	?
Irrig3	?	?	?	?	?
Irrig4	?	?	?	?	?
Irrig5	?	?	?	?	?

b) Fit the additive linear model and display the model summary. Informally, which of the treatments appear to increase the yield the most compared to the control, E?

Answer:

c) Display diagnostic plots for the fitted model. Are there any concerning patterns in the residuals versus fitted values, quantile-quantile plot of studentized residuals, or scale plot of absolute residuals versus fitted values?

Answer:

d) As illustrated in the notes, use the "drop1" method to test the perform F tests of the blocking and treatment factors. What do you conclude from the results?

Answer:

e) Perform a Tukey honest significant differences analysis of the differences between treatment means. Because of the lack of orthoginality between treatment and blocking factors, we can't use the aov and TukeyHSD functions to do the job. Instead we have to compute directly as illustrated in the notes on latin squares and balanced incomplete block designs. To get

started, observe that the margin of error (confidence interval half-width) for Tukey HSD paired differences has the form

qtukey(0.95, nmeans, df)
$$*se_{\mathrm{diff}}/\sqrt{2}$$

where nmeans is the number of different means being compared, df is the residual degrees of freedom for the model, and $se_{\rm diff}$ is the standard error treatment differences reported for the non-reference treatment coefficients in the model summary. Determine which pairs of treatments have significantly different mean yields.

Answer: