HW3: Sections 2.3, 5.2, and 6.2

Solutions

The code below just loads some packages and makes it so that enough digits are printed that you won't get confused by rounding errors.

```
library(dplyr) # functions like summarize
library(ggplot2) # for making plots
library(mosaic) # convenient interface to t.test function
library(readr)
library(gmodels)
options("pillar.sigfig" = 10) # print 10 significant digits in summarize output
```

Problem 1: Sleuth3 2.12 (Marijuana use during pregnancy)

For the birth weights of babies in two groups, one born to mothers who used marijuana during pregnancy and the other born to mothers who did not, the difference in sample averages (mothers who did not use marijuana minus mothers who used marijuana) was 280 grams, and the standard error of the difference was 46.66 grams with 1,095 degrees of freedom.

(a) From the information above, find a 95% confidence interval for $\mu_2 - \mu_1$ (the difference in means for mothers who did not use marijuana and for mothers who did use marijuana) and interpret the interval in context. As part of your answer, explain what the phrase "95% confident" means.

```
c(280-qt(0.975, 1095)*46.66, 280+qt(0.975, 1095)*46.66)
```

```
## [1] 188.4469 371.5531
```

We are 95% confident that the true mean difference in birth weights between babies born to mothers who did not use marijuana and to mothers who did use marijuana is between 188.45 and 371.55 grams. By 95% confident, we mean that for 95% of samples from this population, confidence intervals constructed in this way will contain the true mean difference in birth weights.

(b) Find a 90% confidence for $\mu_2 - \mu_1$. No need to interpret the interval in context.

```
c(280-qt(0.95, 1095)*46.66, 280+qt(0.95, 1095)*46.66)
```

```
## [1] 203.1861 356.8139
```

(c) Calculate the t statistic for a test of the claim that there is no difference in the birth weights for the two groups. You don't need to find the p-value.

```
(280-0)/46.66
```

[1] 6.000857

Problem 2: Adapted from Sleuth3 2.21

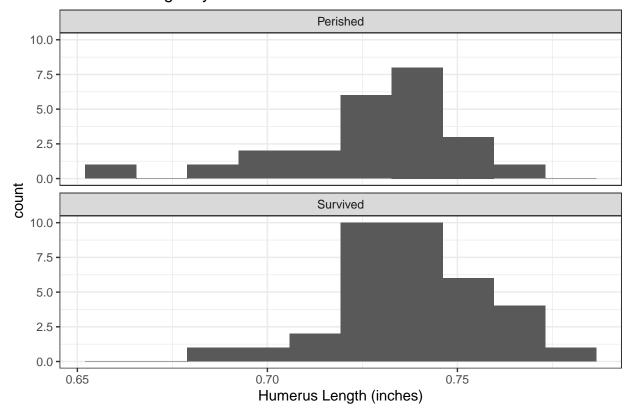
In 1899, biologist Hermon Bumpus presented as evidence of natural selection a comparison of numerical characteristics of moribund house sparrows that were collected after an uncommonly severe winter storm and which had either perished as a result of their injuries or survived. The following R code reads in a data set with the length of the humerus (arm bone) in inches for 59 of these sparrows, grouped according to whether they survived or perished. Analyze these data to summarize the evidence that the mean humerus length is different in the two populations.

```
sparrows <- read.csv("http://www.evanlray.com/data/sleuth3/ex0221_sparrows.csv")</pre>
```

(a) Make an appropriately labelled plot of the data.

```
ggplot(data=sparrows, aes(x=Humerus)) +
  geom_histogram(bins=10) +
  facet_wrap(~ Status, ncol=1) +
  xlab("Humerus Length (inches)") +
  ggtitle("Humerus Length by Survival Status") +
  theme_bw()
```

Humerus Length by Survival Status



(b) Fit a linear model to the data and print out the model summary.

```
lm_sparrows <- lm(Humerus ~ Status, data=sparrows)
summary(lm_sparrows)</pre>
```

```
##
## Call:
## lm(formula = Humerus ~ Status, data = sparrows)
##
## Residuals:
##
         Min
                    1Q
                          Median
                                        3Q
                                                 Max
  -0.068917 -0.010000 0.001083
                                 0.014000
                                            0.042000
##
##
## Coefficients:
##
                  Estimate Std. Error t value Pr(>|t|)
                  0.727917
                             0.004370 166.555
## (Intercept)
                                                <2e-16 ***
## StatusSurvived 0.010083
                             0.005674
                                        1.777
                                                0.0809 .
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.02141 on 57 degrees of freedom
## Multiple R-squared: 0.05249,
                                    Adjusted R-squared:
## F-statistic: 3.158 on 1 and 57 DF, p-value: 0.0809
```

(c) Write down the equation for the mean in terms of the linear model coefficients β_0 and β_1 .

```
\mu = \beta_0 + \beta_1 \times \mathtt{StatusSurvived}
```

(d) What is the interpretation of each of the parameters β_0 and β_1 ?

 β_0 is the mean humerus length in the population of birds that died in the storm.

 β_1 is the difference between the mean humerus length in the populations of birds that survived the storm and the mean humerus length in the population of birds that died in the storm.

(e) Using the output from the model summary in part (b), conduct a test of the claim that there is no difference in the mean length of the humerus in the populations of sparrows before and after the storm.

```
H_0: \beta_1 = 0 versus H_A: \beta_1 \neq 0.
```

The p-value for this test is 0.0809. This provides weak evidence against the null hypothesis that there is no difference in the mean length of the humerus in sparrows that survived the storm versus those that died.

(f) Find a confidence interval for the difference in the mean length of the humerus in the populations of sparrows before and after the storm and discuss what it means in context. You can do this either using the fit.contrast function as we've done in class, or by calling the confint function directly on your model fit from part (b) to obtain confidence intervals for the linear model coefficients.

```
fit.contrast(lm sparrows, "Status", c(-1,1), conf.int = 0.95)
```

```
## Estimate Std. Error t value Pr(>|t|) lower CI
## Status c=(-1 1) 0.01008333 0.005674365 1.776998 0.0809045 -0.001279386
## upper CI
## Status c=(-1 1) 0.02144605
## attr(,"class")
## [1] "fit_contrast"
```

We are 95% confident that the difference in mean humerus lengths in the populations of sparrows that survived in the storm and that died in the storm is between -0.001 and 0.021 inches.

Problem 3: Adapted from Sleuth3 6.22

Was Tyrannosaurus rex warm-blooded? The R code below reads in data with several measurements of the oxygen isotopic composition of bone phosphate in each of 123 bone specimens from a single Tyrannosaurus rex skeleton. It is known that the oxygen isotopic composition of vertebrate bone phosphate is related to the body temperature at which the bone forms. Differences in means at different bone sites would indicate nonconstant temperatures throughout the body. Minor temperature differences would be expected in warm-blooded animals.

The following R code reads in a data set with measurements of oxygen isotopic composition of vertebrate bone phosphate (per mil deviations from SMOW) in 12 bones of a singe Tyrannosaurus rex specimen. For each bone sample, multiple measurements were taken.

```
trex <- read_csv("http://www.evanlray.com/data/sleuth3/ex0523_trex.csv") %>%
  mutate(
    Bone = factor(Bone, levels = paste0("Bone", 1:12))
)

## Parsed with column specification:
## cols(
## Oxygen = col_double(),
```

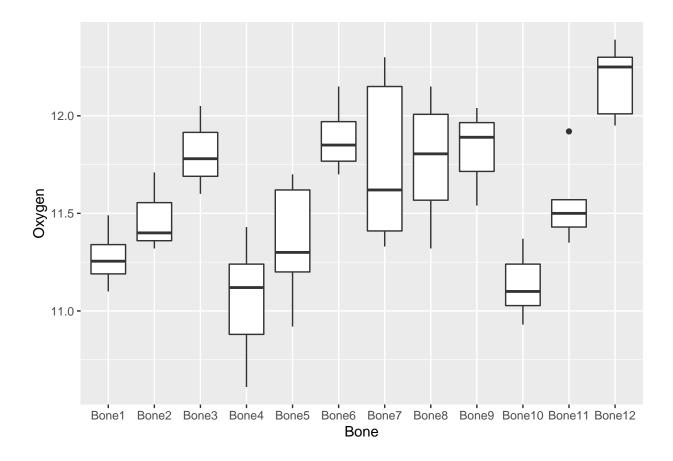
(a) Make side by side box plots of the data

Bone = col_character()

##

)

```
ggplot(data=trex, aes(x=Bone, y=Oxygen)) +
geom_boxplot()
```



(b) Bones 2 and 3 were both gastralia (roughly similar to human ribs), and bones 4 and 5 were both dorsal vertebra (part of the backbone). Is there evidence that the mean oxygen isotopic composition is different for the gastralia than for the dorsal vertebra? Specify a linear combination of means for the 12 bones that could be used to address this question, and conduct a relevant hypothesis test and find a confidence interval. Interpret all of your results in context.

Let μ_i denote the mean oxygen isotopic composition in bone i, for i=2,3,4,5.

$$H_0: \frac{1}{2}(\mu_2 + \mu_3) - \frac{1}{2}(\mu_4 - \mu_5) = 0$$
 versus $H_0: \frac{1}{2}(\mu_2 + \mu_3) - \frac{1}{2}(\mu_4 - \mu_5) \neq 0$

```
lm_trex <- lm(Oxygen ~ Bone, data=trex)
contrast <- c(0,0.5,0.5,-0.5,-0.5,rep(0,7))
fit.contrast(lm_trex, "Bone", contrast, conf.int=0.95)</pre>
```

We estimate the mean oxygen isotopic composition is about 0.441 units higher for the gastralia than for the dorsal vertebra. We are 95% confident that the true mean difference in oxygen isotopic composition is

between 0.16 and 0.73. (In about 95% of samples, a confidence interval calculated in this way would contain the difference in mean oxygen isotopic concentrations for these bones.)

The p-value for this hypothesis test is 0.003, indicating that there is strong evidence against the null that there is no difference. The data indicate that the mean oxygen isotopic concentration is different in these bone groups, suggesting that there were differences in body temperatures in those regions of the dinosaur's body and lending support to the theory that the T.rex was warm blooded.