A Final RMarkdown Document in HTML Format

Student R. Me
10 September, 2019

Homework #1	

The Question

A study was undertaken to explore the foraging behavior of adult ospreys during the breeding season. The first of several questions asked was whether prey choice (capture) was independent of osprey sex during the 3-month breeding season. Individual male and female ospreys were watched and the species of fish captured was recorded.

Objectives were to:

- Determine if prey selection was independent of osprey sex
- Build a plot of the data.
- Examine standardized residuals.

The Analysis Using R

First import the data, then examine the data:

```
# import data from external .csv file
osprey <- read.csv("data/ospreypreybysex.csv", header = T)
osprey # examine raw data</pre>
```

```
##
     ospreysex fishspp count
## 1
          Male Sunfish
## 2
        Female Sunfish
                           72
## 3
          Male
                  Bass
                           14
## 4
        Female
                   Bass
                           21
## 5
          Male
                   Shad
                          189
## 6
        Female
                   Shad
                          138
```

Next create a cross-tabulated table of frequencies by fish species and osprey sex.

```
# crosstabs to mimic data table
osprey.xtab <- xtabs(count ~ fishspp + ospreysex, data = osprey)
osprey.xtab # examine cross-tabulated data</pre>
```

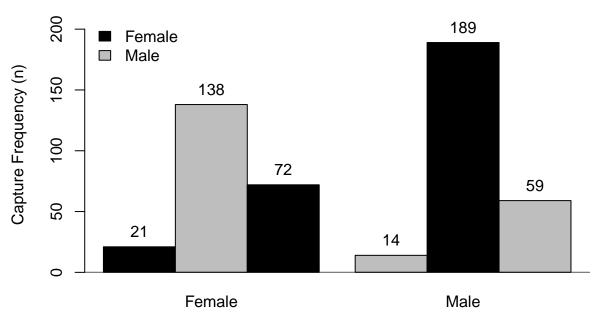
```
## ospreysex
## fishspp Female Male
## Bass 21 14
## Shad 138 189
## Sunfish 72 59
```

Construct a barplot of the capture frequencies.

```
# embed a barplot
freq.r <- range(0, signif(max(osprey$count), 1) + 10) # use raw data for ylim= range
bplot1 <- barplot(osprey.xtab, ylim = freq.r, xlab = "Fish Species",
    ylab = "Capture Frequency (n)", space = c(0, 0.5), col = c("black", "grey"), beside = T)
abline(h = 0) # add line to bottom of plot

# add legend
legend("topleft", c("Female", "Male"), fill = c("black", "grey"), bty = "n", cex = 1)
title(main = "Fish Species Capture Frequency by Osprey Sex")
text(x = bplot1, y = c(osprey.xtab[1:6]), labels = c(osprey.xtab[1:6]), pos = 3) # adds osprey.xtab</pre>
```

Fish Species Capture Frequency by Osprey Sex



Fish Species

Implement the χ^2 test, examine output, and see what objects are available after the analysis.

```
# Pearson chisq
osprey.chi2 <- chisq.test(osprey.xtab) # chisq test
osprey.chi2 # output from chisq test</pre>
```

```
Pearson's Chi-squared test
##
## data: osprey.xtab
## X-squared = 8.7294, df = 2, p-value = 0.01272
 ls(osprey.chi2) # available objects from chisq test
## [1] "data.name" "expected" "method"
                                             "observed"
                                                         "p.value"
                                                                      "parameter"
## [7] "residuals" "statistic" "stdres"
Last, examine some diagnostics for interpretation, especially the standardized residuals. The standardized
residuals provide an indication where deviations for expected capture frequencies deviate.
# examine diet preferences by sex
  chisq.test(osprey.xtab)$observed
                                    # observed captures
##
            ospreysex
             Female Male
## fishspp
##
     Bass
                 21
                       14
##
     Shad
                138
                    189
##
     Sunfish
                 72
                      59
 chisq.test(osprey.xtab)$expected # expected captures
##
            ospreysex
                Female
## fishspp
                             Male
##
     Bass
              16.39959 18.60041
             153.21907 173.78093
##
     Shad
     Sunfish 61.38134 69.61866
##
 100 * osprey.xtab/apply(osprey.xtab, 1, sum) # percent captures
##
            ospreysex
## fishspp
               Female
                           Male
             60.00000 40.00000
##
     Bass
##
     Shad
             42.20183 57.79817
##
     Sunfish 54.96183 45.03817
 chisq.test(osprey.xtab)$stdres # standardized residuals
##
            ospreysex
## fishspp
                Female
              1.616751 -1.616751
##
     Bass
             -2.906524 2.906524
##
     Shad
##
     Sunfish 2.169669 -2.169669
```

Interpretation

The Analysis Using R

Male and female ospreys differ in their diets ($\chi^2 = 8.729$, p = 0.013, df = 2). Shad makes up more than half the diet of both male and female birds, but for the males it is over 72%, whereas for the females it is about 60%. Proportionally, bass and sunfish make up slightly smaller portions of the average male's diet than in the female's diet.

Overall, both males and females appear to be selecting bass at approximately their relative availability. Males preferentially capture shad at higher proportions than expected, while under-utilizing sunfish. Females preferentially capture sunfish and ignore shad.

The RMD code used to build the HTML output is in the box below

```
title: "A Final **R**Markdown Document in HTML Format"
author: "Student R. Me"
date: "10 September, 2019"
output:
   html_document:
       keep_md: true
   pdf_document: default
   word_document: default
<!-- set root directory here -->
<!-- your directory will be specific to you -->
```{r global_options, include=FALSE}
knitr::opts_knit$set(root.dir = "~/words/github/useRfiles/rmd-files")
knitr::opts_chunk$set(warning=FALSE)
Homework \#1
The Question
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Objectives were to:
* Determine if prey selection was independent of osprey sex
* Build a plot of the data.
* Examine standardized residuals.
```

```
First import the data, then examine the data:
```{r}
# import data from external .csv file
 osprey <- read.csv("data/ospreypreybysex.csv", header = T)</pre>
 osprey # examine raw data
Next create a cross-tabulated table of frequencies by fish species and osprey sex.
```{r}
crosstabs to mimic data table
 osprey.xtab <- xtabs(count ~ fishspp + ospreysex, data = osprey)
 osprey.xtab # examine cross-tabulated data
Construct a barplot of the capture frequencies.
```{r}
# embed a barplot
  freq.r <- range(0, signif(max(osprey$count), 1) + 10) # use raw data for ylim= range
  bplot1 <- barplot(osprey.xtab, ylim = freq.r, xlab = "Fish Species",</pre>
   ylab = "Capture Frequency (n)", space = c(0, 0.5), col = c("black", "grey"), beside = T)
  abline(h = 0) # add line to bottom of plot
# add legend
  legend("topleft", c("Female", "Male"), fill = c("black", "grey"), bty = "n", cex = 1)
  title(main = "Fish Species Capture Frequency by Osprey Sex")
 text(x = bplot1, y = c(osprey.xtab[1:6]), labels = c(osprey.xtab[1:6]), pos = 3) # adds osprey.xtab
Implement the $\chi^2$ test, examine output, and see what objects are available after the analysis.
```{r}
Pearson chisq
 osprey.chi2 <- chisq.test(osprey.xtab) # chisq test</pre>
 osprey.chi2 # output from chisq test
 ls(osprey.chi2) # available objects from chisq test
Last, examine some diagnostics for interpretation, especially the standardized residuals. The standard
```{r}
# examine diet preferences by sex
  chisq.test(osprey.xtab)$observed # observed captures
  chisq.test(osprey.xtab)$expected # expected captures
  100 * osprey.xtab/apply(osprey.xtab, 1, sum) # percent captures
 chisq.test(osprey.xtab)$stdres # standardized residuals
```

Interpretation

Male and female ospreys differ in their diets (\$\chi^2\$ = `r round(osprey.chi2\$statistic, 3)`, *p* = `r Overall, both males and females appear to be selecting bass at approximately their relative availability.