t-test-hw-Key

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- Q1. (2 points) Describe null and alternative hypotheses for:
 - The difference in means of seal count in Wilhelmenia vs. Marguarite bay
 - the difference in means of fish count in Wilhelmenia vs. Marguarite bay
- Q2. (2 points) Generate helpful visualizations and descriptive statistics for the above data
- Q3. (4 points) Perform t-tests to evaluate your hypotheses, and interpret the results. Reject the null hypothesis if p<0.05.

Answers

Q1.

 H_0 : There is no difference in the mean seal count in Wilhelmia and Marguarite bay. There is no difference in the mean fish count in Wilhelmia and Marguarite bay.

 H_a :There is a difference in the mean seal count in Wilhelmia and Marguarite bay. There is a difference in the mean fish count in Wilhelmia and Marguarite bay.

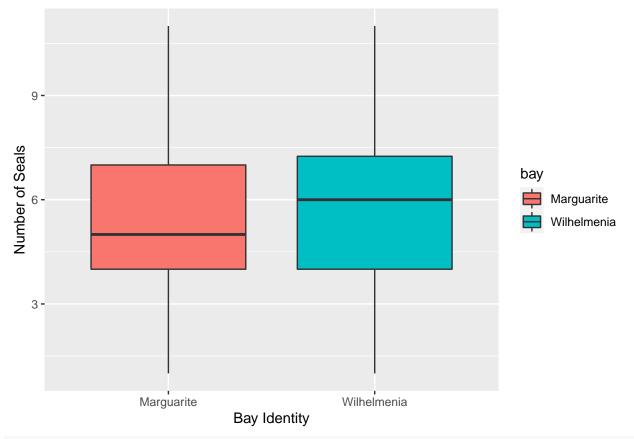
Q2.

```
library("tidyverse")
```

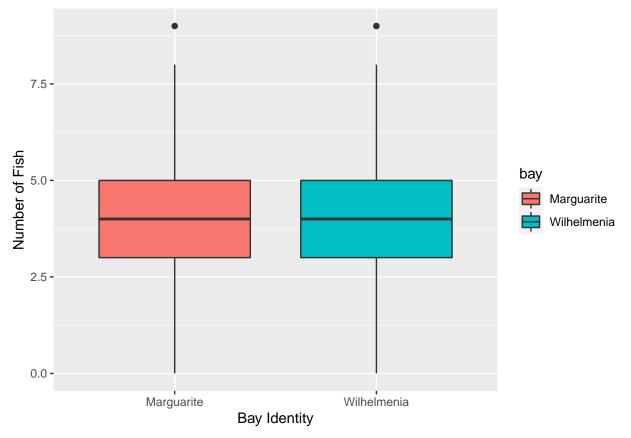
Rows: 640 Columns: 5

```
## -- Attaching packages ----- tidyverse 1.3.2 --
## v ggplot2 3.3.6
                                0.3.4
                      v purrr
## v tibble 3.1.8
                      v dplyr
                                1.0.10
## v tidyr
            1.2.1
                      v stringr 1.4.1
## v readr
            2.1.2
                      v forcats 0.5.2
## -- Conflicts -----
                              ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
seals <- read_csv("arctic-seals.csv")</pre>
## Rows: 640 Columns: 5
## -- Column specification -
## Delimiter: ","
## chr (2): time, bay
## dbl
       (2): area, num_seals
## date (1): date
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
fish <- read_csv("arctic-fish.csv")</pre>
```

```
## -- Column specification -----
## Delimiter: ","
## chr (2): time, bay
## dbl (2): net, num_fish
## date (1): date
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
# descriptive stats
summarySeals <- seals %>% group_by(bay) %>% summarize(mean(num_seals), sd(num_seals))
summaryFish <- fish %>% group_by(bay) %% summarize(mean(num_fish), sd(num_fish))
# data visualization: best to do a boxplot or bar chart
sealsPlot <- seals %>%
  ggplot(aes(bay, num_seals, fill= bay)) +
 geom_boxplot() +
 xlab("Bay Identity") +
 ylab("Number of Seals")
fishPlot <- fish %>% ggplot(aes(bay, num_fish, fill = bay)) +
  geom_boxplot() +
 xlab("Bay Identity") +
 ylab("Number of Fish")
summarySeals
## # A tibble: 2 x 3
##
    bay
                `mean(num_seals)` `sd(num_seals)`
##
     <chr>
                             <dbl>
                                             <dbl>
## 1 Marguarite
                             5.25
                                              2.10
## 2 Wilhelmenia
                              5.95
                                              2.10
summaryFish
## # A tibble: 2 x 3
                `mean(num_fish)` `sd(num_fish)`
##
    bay
##
     <chr>
                           <dbl>
                                          <dbl>
## 1 Marguarite
                            3.91
                                           1.76
## 2 Wilhelmenia
                            4.16
                                           1.96
sealsPlot
```



fishPlot



```
Q3.
```

data: num_fish by bay

```
# t test for seals
sealsT <- t.test(data = seals, num_seals ~ bay)</pre>
sealsT
##
##
    Welch Two Sample t-test
## data: num_seals by bay
## t = -4.2182, df = 638, p-value = 2.82e-05
## alternative hypothesis: true difference in means between group Marguarite and group Wilhelmenia is n
## 95 percent confidence interval:
## -1.0258729 -0.3741271
## sample estimates:
##
   mean in group Marguarite mean in group Wilhelmenia
                        5.25
# Interpretation: The p-value is 0.0000272, which is less than our alpha level of 0.05, so we can rejec
# We conclude that the mean number of seals is different between Marguarite and Wilhelmia Bay
# t test for fish
fishT <- t.test(data = fish, num_fish ~ bay)</pre>
fishT
##
   Welch Two Sample t-test
##
```

Interpretation: The p-value was 0.08, which is greater than our alpha level of 0.05, so we cannot rej # We conclude that the mean number of fish is not different between Marguarite and Wilhelmia Bay.