SOC 325: Quantified-Self

Basics of Statistics

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Road Map

Section Introduction

Data

Descriptive Statistics

Statistical Inference

Linear Regression

Visualizations

Summary



Section Introduction



R Lab

- ► No R Lab this week
- ► Everyone gets full credit
- ightharpoonup Extra week with Lab 1



Overview

- ▶ In this lecture we are going to review basic statistics with R
 - ► Central Tendencies (e.g. mean, median, mode)
 - ► Volatility (e.g. variance, sd)
 - ► Statistical test (e.g. permutation and t-test)
 - ► Linear Regression



Data



Data

Half Marathon Times at Cherry Blossom Run





Data: Overview

- ▶ Results for Seattle Cherry Blossom Run 2022
 - ▶ I used Chrome to download an archive of these results
 - ► To directly scrape is a bit complicated because you have to get the javascript to precompile
- ▶ We will use the following packages to read and manipulate the data

```
library(tidyverse)
library(textreadr)
library(rvest)
library(lubridate)
library(AMR)
library(here)
```



Data: Import and clean

```
## Read in the archived results
cb<-rvest::read_html(here("data/","half", "Seattle Cherry Blossom Run Results.html"))
## Clean and pull the table
cb_tab<-cb/b/% minimal_html()%>%
html_node("table") %>%
html_table(header=i)

## Clean up name and add time in minutes
cb_tab<-cb_tab/%/mutate(
ChipTime_min=hns(ChipTime),
ChipTime_min=hns(ChipTime),
html chipTime_min=hour(ChipTime_min)*60 + minute(ChipTime_min)*second(ChipTime_min)/100,
Name=substr(str_remove_all(str_remove_all(Name,"\\n"),"\\t"),2,1000000L)

## Add age groups in 5s
cb_tab$age groups<-AMR::age groups(cb_tab$Age,"fives")
```



Data: Import and clean

```
glimpse(cb_tab)
```

```
## Rows: 1.198
## Columns: 13
                     <int> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 1~
## $ Place
                     <int> 19025, 17022, 17561, 17591, 17659, 17614, 17133, 171~
## $ Bib
                     <chr> "KyleCochrun", "JuliaBudniak", "HaroldDanks", "Jared~
## $ Name
## $ Gender
                     ## $ Citv
                     <chr> "Seattle", "Lakewood", "Chattanooga", "Washington", ~
## $ State
                     <chr> "WA", "WA", "TN", "DC", "WA", "WA", "WA", "WA", "WA", "WA"~
## $ ChipTime
                     <chr> "1:19:46.38", "1:21:18.23", "1:21:39.60", "1:21:45.9~
                     <int> 29, 41, 25, 41, 12, 24, 36, 26, 27, 26, 29, 32, 33, ~
## $ Age
## $ DivisionPlace
                     <int> 1, 1, 2, 1, 1, 1, 1, 3, 5, 4, 6, 1, 2, 3, 8, 7, 2, 2~
                     <chr> "M 25-29", "F 40-44", "M 25-29", "M 40-44", "M 10-14~
## $ Division
## $ `Group/Team Name` <chr> "", "", "", "", "", "Seattle Frontrunners", "Sea~
## $ ChipTime min
                     <dbl> 79.4638, 81.1823, 81.3960, 81.4590, 82.3811, 83.3603~
## $ age groups
                     <ord> 25-29, 40-44, 25-29, 40-44, 10-14, 20-24, 35-39, 25-~
```





```
cb_tab%>%group_by(age_groups)%>%summarise(n=n())
## # A tibble: 13 x 2
     age_groups
##
     <ord>
                 <int>
   1 10-14
   2 15-19
                   19
  3 20-24
                  127
  4 25-29
                  275
## 5 30-34
                  214
  6 35-39
                  166
## 7 40-44
                  143
## 8 45-49
                   89
  9 50-54
                   80
## 10 55-59
                    37
## 11 60-64
                    28
## 12 65-69
                    12
## 13 70-74
```

Counts by age



... with 22 more rows

```
## Counts by age
cb_tab%>%group_by(age_groups,Gender)%>%summarise(n=n())
## # A tibble: 32 x 3
## # Groups:
               age_groups [13]
      age_groups Gender
      <ord>
##
                 <chr> <int>
   1 10-14
                            3
   2 10-14
   3 15-19
                 "F"
   4 15-19
                           10
   5 20-24
                            1
   6 20-24
                           63
  7 20-24
                           63
   8 25-29
                            3
   9 25-29
                          122
## 10 25-29
                          148
```



... with 22 more rows

```
## Counts by age
cb_tab%>%group_by(age_groups,Gender)%>%summarise(n=n())
## # A tibble: 32 x 3
## # Groups:
               age_groups [13]
      age_groups Gender
      <ord>
##
                 <chr> <int>
   1 10-14
                            3
   2 10-14
   3 15-19
                 "F"
   4 15-19
                           10
   5 20-24
                            1
   6 20-24
                           63
  7 20-24
                           63
   8 25-29
                            3
   9 25-29
                          122
## 10 25-29
                          148
```



Update data: Limit to just "M" and "F" cases

 $\verb|cb_tab_mf<-cb_tab|| > & filter(Gender in & ("F", "M")) \\$



Age groups and Gender

... with 16 more rows

```
cb_tab_mf%>%group_by(age_groups,Gender)%>%summarise(n=n())
## # A tibble: 26 x 3
## # Groups:
               age_groups [13]
     age_groups Gender
     <ord>
##
                <chr> <int>
   1 10-14
   2 10-14
  3 15-19
  4 15-19
                          10
  5 20-24
                          63
  6 20-24
                          63
## 7 25-29
                         122
   8 25-29
                         148
   9 30-34
                         106
## 10 30-34
                         106
```



Total AvgTime

med <int> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> < ## 1 1188 138. 0.0280 131. 96.3 203.

```
stats_overall<-cb_tab_mf%>%summarise(
 Total=n().
 AvgTime=mean(ChipTime_min),
 se=sd(ChipTime_min)/Total,
 med=median(ChipTime_min),
 q5=quantile(ChipTime_min,.05),
 q95=quantile(ChipTime_min,.95)
## Print
stats_overall
## # A tibble: 1 x 6
```



```
stats_gender<-cb_tab_mf%>%group_by(Gender)%>%summarise(
   Total=n(),
   AvgTime=mean(ChipTime_min),
   se=sd(ChipTime, min)/Total,
   med=median(ChipTime min),
   q5=quantile(ChipTime_min,.05),
   q95=quantile(ChipTime_min,.95)
)

## Print
stats_gender
```

```
## # A tibble: 2 x 7

## Gender Total AvgTime se med q5 q95

## <chr> <int> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> = 137. 109. 216.

## 2 M 551 128. 0.0537 120. 89.8 184.
```





We will use the infer package

library(infer)



ightharpoonup Mean: $\bar{x} = \frac{1}{n} \sum_{i=1}^{n} x_i$

calculate the observed statistic
observed statistic <- cb_tab mf %>%

► We will use a simple permutation/simulation test

```
specify(response = ChipTime_min) %>%
calculate(stat = "mean")

observed_statistic

## Response: ChipTime_min (numeric)
## # A tibble: 1 x 1
## stat
## <dbl>
## 138.
```

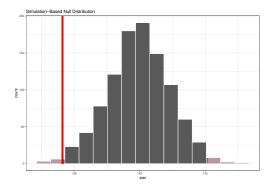


- ▶ Mean: $\bar{x} = \frac{1}{n} \sum_{1}^{n} x_i$
- ► The average Female Time for a half Marathon for 20-24 year olds is 140 Min
 - ➤ Our null hypthesis is that average time for a runner in the CBR is 140 Min
- We will generate a plausable null distribution from this assumption

```
# generate the null distribution
null_dist_1_sample <- cb_tab_mf %>%
specify(response = ChipTime_min) %>%
hypothesize(null = "point", mu = 140) %>%
generate(reps = 1000, type = "bootstrap") %>%
calculate(stat = "mean")
```



```
ightharpoonup Mean: \bar{x} = \frac{1}{n} \sum_{1}^{n} x_i
```





```
## # A tibble: 1 x 1
## p_value
## <dbl>
## 1 0.014
```



- $\blacktriangleright \text{ Mean: } \bar{x} = \frac{1}{n} \sum_{1}^{n} x_{i}$
- ▶ Instead of the permutation/simulation test we can do a t-test
 - ► Approximate normal test
- $\blacktriangleright \ t = \frac{\bar{x}}{se(\bar{x})}$
 - $ightharpoonup se(\bar{x}) = sd(x)/\sqrt{n}$
 - $sd(x) = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (x_i \bar{x})^2}$ (mle)
 - $sd(x) = \sqrt{\frac{1}{n-1} \sum_{i=1}^{n} (x_i \bar{x})^2}$ (unbiased)

```
##t.test
cb_tab_mf%>%t_test(response =ChipTime_min , mu = 140)
```

```
## # A tibble: 1 x 7

## statistic t_df p_value alternative estimate lower_ci upper_ci
## <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> 138. 136. 140.
```



- ▶ P-value of the t-statistic
 - ▶ Provides simple decision heuristic for t-test
 - ▶ p-value < 0.05 statistically significant (there is a difference!)
 - ▶ p-value > 0.05 statistically not significant (there is **not** a difference!)

```
# calculate the observed statistic
observed_statistic <- cb_tab_mf %>%
specify(response = ChipTime_min) %>%
hypothesize(null = "point", mu = 40) %>%
calculate(stat = "t") %>%
dplyn::pull()

# calculate 2-tail t-test
pt(observed_statistic, df = nrow(gss) - 1, lower.tail = FALSE)*2
```

t ## 0



- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a permutation terst here too!

```
# calculate the observed statistic
observed_statistic <- cb_tab_mf %>%
    specify( ChipTime_min - Gender) %>%
    calculate(stat = "diff in means", order = c("F", "M"))
observed_statistic
```

```
## Response: ChipTime_min (numeric)
## Explanatory: Gender (factor)
## # A tibble: 1 x 1
## stat
## <dbl>
## 1 18.5
```

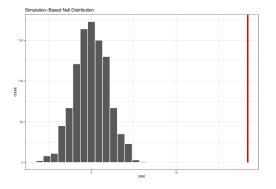


- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a permutation terst here too!

```
# generate the null distribution with randomization
null_dist_2_sample <- cb_tab_mf %>%
specify(ChipTime_min ~ Gender) %>%
hypothesize(null = "independence") %>%
generate(reps = 1000, type = "permute") %>%
calculate(stat = "diff in means", order = c("F", "M"))
```



- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a permutation terst here too!





- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a permutation terst here too!

```
## # A tibble: 1 x 1
## p_value
## <dbl>
## 1 0
```



- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a t-test here too

```
t = \bar{x}_1 - \bar{x}_2/se(\bar{x}_1 - \bar{x}_2)
```

```
## # A tibble: 1 x 7
## statistic t_df p_value alternative estimate lower_ci upper_ci
## <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> 21.1
## 1 10.0 1186 8.62e-23 two.sided 18.5 14.9 22.1
```



Difference of Means Test

- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a t-test here too
- $t = \bar{x}_1 \bar{x}_2/se(\bar{x}_1 \bar{x}_2)$

```
# calculate the observed statistic
observed_statistic <- cb_tab_mf %>%
specify(ChipTime_min - Gender,) %>%
hypothesize(mull = "point", mu = 0) %>%
calculate(stat = "t", order = c("F", "M")) %>%
dplyr::pull()
```

 $observed_statistic$

```
## t
## 10.03263
```



- $ightharpoonup \bar{x}_1 \bar{x}_2$
- ▶ We can do a t-test here too
- $t = \bar{x}_1 \bar{x}_2/se(\bar{x}_1 \bar{x}_2)$
- ▶ We can again compute the p-value
 - ▶ p-value < 0.05 statistically significant (there is a difference!)
 - ▶ p-value > 0.05 statistically not significant (there is **not** a difference!)

```
# calculate the observed statistic
observed_statistic <- cb_tab_mf %%
specify(ChipTime_min - Gender,) %%
hypothesize(null = "point", mu = 0) %%
calculate(stat = "t", order = c("F", "M")) %>%
dplyr::pull()
observed_statistic
```

```
## t
## 10.03263
pt(observed_statistic, df = nrow(gss) - 2, lower.tail = FALSE)*2
## t
## 1.070199e-21
```



Difference of Means Test

▶ Old (over 35) versus Young (under 35)

```
## [1] "Young" "Old"
```



Difference of Means Test

▶ Old (over 35) versus Young (under 35)

```
observed_statistic <- cb_tab_mf%>%filter(Gender=="M") %>%
    specify(ChipTime_min ~ youngOld) %>%
    calculate(stat = "diff in means", order = c("Old", "Young"))

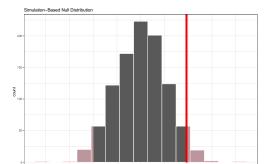
observed_statistic
```

```
## Response: ChipTime_min (numeric)
## Explanatory: youngOld (factor)
## # A tibble: 1 x 1
## stat
## <dbl>
## 1 4.82
```



Difference of Means Test

▶ Old (over 35) versus Young (under 35)





Statistical Inference

<dbl>

1

Difference of Means Test

▶ Old (over 35) versus Young (under 35)



Statistical Inference

Difference of Means Test

▶ Old (over 35) versus Young (under 35)



Statistical Inference

Difference of Means Test

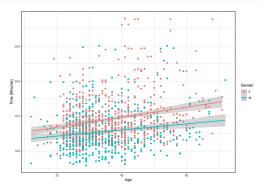
▶ Old (over 35) versus Young (under 35)

```
## t-test
cb_tab_mf%>%filter(Gender=="M")%>%t_test(
  formula = ChipTime_min - youngOld,
    order = c("Dld", "Young"),
    alternative = "two-sided")
```





```
cb_tab_mf%>%ggplot(aes(y=ChipTime_min,x=Age,color=Gender))+
geom_point()+
#stat_summary(fun.data= mean_cl_normal) +
geom_smooth(method='lm')+
theme_bw()+
xlab("Age")+
ylab("Time (Minutes)")
```





► Simple Linear Regression

$$Y_i = \beta_0 + \beta_1 X_i + \epsilon_i$$

* Y in this case is run time in minutes (e.g. 140 minutes) * X is Age (e.g. 25) * Epsilon is deviation around the score (or measurment error) $+ \epsilon \sim N(0,\sigma)$ (Normal Distribution) * R^2 is a measure of fit that ranges from 0 [low] to 1 [high]

```
summary(lm(ChipTime_min~Age,data=cb_tab_mf))
```

```
##
## Call:
## lm(formula = ChipTime_min ~ Age, data = cb_tab_mf)
##
## Residuals:
     Min
            1Q Median 3Q
## -60 03 -21 61 -6 13 14 30 148 21
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 113.36433 3.18620 35.580 < 2e-16 ***
              0.67918
                          0.08515 7.976 3.53e-15 ***
## Age
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 32.4 on 1186 degrees of freedom
## Multiple R-squared: 0.05091, Adjusted R-squared: 0.05011
```



$$Y_i = \beta_0 + \beta_1 X_{1i} + \dots + \beta_{ki} X_{ki} + \epsilon_{ki}$$

```
models<-list(
lm(ChipTime_min-1,data=cb_tab_mf),
lm(ChipTime_min-Age,data=cb_tab_mf),
lm(ChipTime_min-Age+Gender,data=cb_tab_mf),
)

r2<-models%>%purrr::map(function(x){summary(x)$r.squared})%>%unlist()
ar2<-models%>%purrr::map(function(x){summary(x)$adj.r.squared})%>%unlist()
data.frame(model=1:3,r2,ar2)
```

```
## model r2 ar2
## 1 1 0.00000000 0.0000000
## 2 2 2 0.05091324 0.0501130
## 3 3 0.11509494 0.1136014
```



$$Y_i = \beta_0 + \beta_1 X_{1i} + \dots + \beta_{ki} X_{ki} + \epsilon_{ki}$$

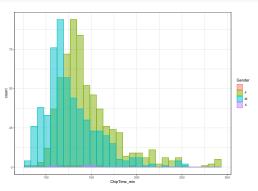
| ## | ## # A tibble: 6 x 6 | | | | | | |
|----|----------------------|-------------|-------------|-------------|-------------|-------------|-------------|
| ## | | models | term | estimate | std.error | statistic | p.value |
| ## | | <chr></chr> | <chr></chr> | <dbl></dbl> | <dbl></dbl> | <dbl></dbl> | <dbl></dbl> |
| ## | 1 | 1 | (Intercept) | 138. | 0.964 | 143. | 0 |
| ## | 2 | 2 | (Intercept) | 113. | 3.19 | 35.6 | 2.90e-189 |
| ## | 3 | 2 | Age | 0.679 | 0.0851 | 7.98 | 3.53e- 15 |
| ## | 4 | 3 | (Intercept) | 124. | 3.30 | 37.7 | 4.39e-205 |
| ## | 5 | 3 | Age | 0.593 | 0.0828 | 7.16 | 1.42e- 12 |
| ## | 6 | 3 | GenderM | -17.0 | 1.83 | -9.27 | 8.46e- 20 |





► Histogram

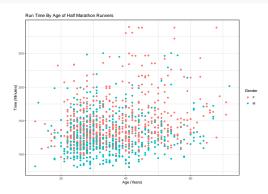
```
cb_tab%/%ggplot(aes(x=ChipTime_min, color=Gender,fill=Gender)) +
  geom_histogram(alpha=0.5, position="identity")+
  theme_bw()
```





► Scatter Plot

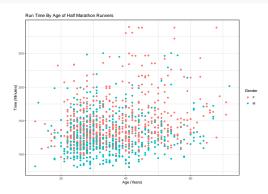
```
cb_tab_mf%>%ggplot(aes(y=ChipTime_min,x=Age,color=Gender))+
geom_point()+
theme_bw()+
ggtitle("Run Time By Age of Half Marathon Runners")+
xlab("Age (Years)")+
ylab("Time (Minutes)")
```





► Scatter Plot

```
cb_tab_mf%>%ggplot(aes(y=ChipTime_min,x=Age,color=Gender))+
geom_point()+
theme_bw()+
ggtitle("Run Time By Age of Half Marathon Runners")+
xlab("Age (Years)")+
ylab("Time (Minutes)")
```

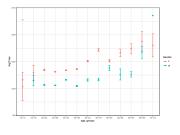




► Line plot with SE

```
stats<-cb_tab_mf%>%group_by(age_groups,Gender)%>%summarise(
   Total=n(),
   AvgTime=mean(ChipTime_min),
   se=sd(ChipTime_min)/Total,
   med=median(ChipTime_min),
   q5=quantile(ChipTime_min,.05),
   q95=quantile(ChipTime_min,.95)
)

stats%>%ggplot(aes(y=AvgTime,x=age_groups,group=Gender,color=Gender))+
   geom_point()+
   geom_errorbar(aes(ymin=AvgTime-2*se, ymax=AvgTime+2*se), width=.2)+
   ylim(95,205)+
   theme_bw()
```

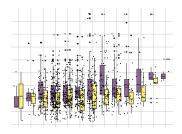




▶ Box Plot

```
library(viridis)
library(hrbrthemes)

cb_tab_mf%-%
    ggplot( aes(x=age_groups, y=ChipTime_min, fill=Gender)) +
    geom_boxplot() +
    scale_fill_viridis(discrete = TRUE, alpha=0.6) +
    geom_jitter(color="black", size=0.4, alpha=0.9) +
    hrbrthemes::theme_ipsum() +
    theme(
        legend.position="none",
        plot.title = element_text(size=11)
    ) +
    ggtitle("Boxplot of Times by Age Groups") +
    xlab("Age Groups")+
    ylab("Time (Minutes)")
```





Summary



Basic Statistics in R for QS

- ▶ Descriptive statistics (mean, median, sd, se)
- ▶ Inferential statistics (permutation test, t-test, p-value)
- ► Linear Regression (Linear relationship between variables)



End of Slides



