

## Final Project 2021, Jakob Pachucinski and Alon Kremerman

```
library(tidyverse)
library(mosaic)
library(Lahman)
library(mdsr)
library(tidymodels)

#For the project, we want to explore whether certain factors such as height,
weight, and birthplace
#effect a certain stat on hitters or pitchers
#For pitchers we will explore ERA and SO
#For the hitters, we evaluated OBP, AVG, SLG, SO, HR, SB
#We hypothesized that heavier players would have a bigger slugging percentage
#While lighter players would have more stolen bases
#Also, shorter players might have a higher OBP because they have a smaller
strike zone
#While taller players may have more SOs with a bigger strike zone
#For the pitching, we suggested that taller pitchers may have more strikeouts
#Since technically they release the ball closer to the plate
help("Lahman")

## starting httpd help server ... done

#Will use this data frame for the personal stats such as height, weight, etc.
head(People)

##   playerID birthYear birthMonth birthDay birthCountry birthState
birthCity
## 1 aardsda01      1981          12        27          USA          CO
Denver
## 2 aaronha01      1934           2         5          USA          AL
Mobile
## 3 aaronto01      1939           8         5          USA          AL
Mobile
## 4 aasedo01      1954           9         8          USA          CA
Orange
## 5 abadan01      1972           8        25          USA          FL Palm
Beach
## 6 abadfe01      1985          12        17          D.R. La Romana La
Romana
##   deathYear deathMonth deathDay deathCountry deathState deathCity
nameFirst
## 1          NA          NA        NA          <NA>          <NA>          <NA>
David
## 2        2021           1        22          USA          GA          Atlanta
Hank
## 3        1984           8        16          USA          GA          Atlanta
```

```

Tommie
## 4      NA      NA      NA      <NA>      <NA>      <NA>
Don
## 5      NA      NA      NA      <NA>      <NA>      <NA>
Andy
## 6      NA      NA      NA      <NA>      <NA>      <NA>
Fernando
##  nameLast      nameGiven weight height bats throws      debut
finalGame
## 1  Aardsma      David Allan   215    75    R      R 2004-04-06 2015-08-
23
## 2   Aaron      Henry Louis   180    72    R      R 1954-04-13 1976-10-
03
## 3   Aaron      Tommie Lee    190    75    R      R 1962-04-10 1971-09-
26
## 4    Aase  Donald William   190    75    R      R 1977-07-26 1990-10-
03
## 5    Abad   Fausto Andres   184    73    L      L 2001-09-10 2006-04-
13
## 6    Abad Fernando Antonio   235    74    L      L 2010-07-28 2019-09-
28
##  retroID  bbrefID  deathDate  birthDate
## 1 aardd001 aardsda01      <NA> 1981-12-27
## 2 aaroh101 aaronha01 2021-01-22 1934-02-05
## 3 aarot101 aaronto01 1984-08-16 1939-08-05
## 4 aased001 aasedo01      <NA> 1954-09-08
## 5 abada001 abadan01      <NA> 1972-08-25
## 6 abadf001 abadfe01      <NA> 1985-12-17

#Selected only the columns that we "might" use and dropped all the NA values
players1 <- People%>%
  select(playerID, birthYear, birthCountry, birthState, nameGiven, weight,
height, debut, bats)%>%
  drop_na()
head(players1)

##  playerID birthYear birthCountry birthState      nameGiven weight
height
## 1 aardsda01    1981      USA      CO    David Allan    215
75
## 2 aaronha01    1934      USA      AL    Henry Louis    180
72
## 3 aaronto01    1939      USA      AL      Tommie Lee    190
75
## 4 aasedo01    1954      USA      CA  Donald William    190
75
## 5 abadan01    1972      USA      FL   Fausto Andres    184
73
## 6 abadfe01    1985      D.R.   La Romana Fernando Antonio 235
74

```

```
##      debut bats
## 1 2004-04-06    R
## 2 1954-04-13    R
## 3 1962-04-10    R
## 4 1977-07-26    R
## 5 2001-09-10    L
## 6 2010-07-28    L
```

*#Headed the batting to see which stats we should explore*  
head(Batting)

```
##      playerID yearID stint teamID lgID  G  AB  R  H X2B X3B HR RBI SB CS BB
SO
## 1 abercda01   1871     1   TRO   NA   1   4  0  0  0  0  0  0  0  0  0
0
## 2 addybo01   1871     1   RC1   NA  25 118 30 32   6  0  0  13  8  1  4
0
## 3 allisar01   1871     1   CL1   NA  29 137 28 40   4  5  0  19  3  1  2
5
## 4 allisdo01   1871     1   WS3   NA  27 133 28 44  10  2  2  27  1  1  0
2
## 5 ansonca01   1871     1   RC1   NA  25 120 29 39  11  3  0  16  6  2  2
1
## 6 armstbo01   1871     1   FW1   NA  12  49  9 11   2  1  0   5  0  1  0
1
##      IBB HBP SH SF GIDP
## 1  NA  NA NA NA   0
## 2  NA  NA NA NA   0
## 3  NA  NA NA NA   1
## 4  NA  NA NA NA   0
## 5  NA  NA NA NA   0
## 6  NA  NA NA NA   0
```

*#Grouped by playerID so I could combine all the players years into one total career*

*#Then summed all the AB, H, HR, RBI, SB, and SO*

*#DBL and TRP were meant to be amount of doubles and triples*

*#Then after filtering dropped all the NAs*

```
batters1 <- Batting%>%
  group_by(playerID)%>%
  summarise(AB = sum(AB), H = sum(H), DBL = sum(X2B), TRP = sum(X3B), HR =
sum(HR), RBI = sum(RBI), SB = sum(SB), SO = sum(SO), BB = sum(BB))%>%
  drop_na()
head(batters1)
```

```
## # A tibble: 6 x 10
```

```
##   playerID      AB      H    DBL    TRP     HR    RBI    SB    SO    BB
##   <chr>      <int> <int> <int> <int> <int> <int> <int> <int> <int>
## 1 aardsda01      4      0      0      0      0      0      0      2      0
## 2 aaronha01  12364  3771   624    98   755  2297   240  1383  1402
## 3 aaronto01   944   216    42     6    13    94     9   145    86
```

```
## 4 aasedo01      5      0      0      0      0      0      0      3      0
## 5 abadan01     21      2      0      0      0      0      0      5      4
## 6 abadfe01      9      1      0      0      0      0      0      5      0
```

*#Explored the pitching metrics that the Lahman package gave us*  
head(Pitching)

```
##      playerID yearID stint teamID lgID  W  L  G  GS  CG  SHO  SV  IPouts   H  ER
HR BB
## 1 bechtge01   1871      1   PH1   NA   1  2  3  3  2   0  0    78  43  23
0 11
## 2 brainas01   1871      1   WS3   NA  12 15 30 30 30   0  0   792 361 132
4 37
## 3 fergubo01   1871      1   NY2   NA   0  0  1  0  0   0  0     3   8   3
0  0
## 4 fishech01   1871      1   RC1   NA   4 16 24 24 22   1  0   639 295 103
3 31
## 5 fleetfr01   1871      1   NY2   NA   0  1  1  1  1   0  0    27  20  10
0  3
## 6 flowedi01   1871      1   TRO   NA   0  0  1  0  0   0  0     3   1   0
0  0
##      SO BAOpp   ERA  IBB  WP  HBP  BK   BFP  GF   R  SH  SF  GIDP
## 1  1      NA  7.96  NA   7   NA   0   146   0  42  NA  NA   NA
## 2 13      NA  4.50  NA   7   NA   0  1291   0 292  NA  NA   NA
## 3  0      NA 27.00  NA   2   NA   0    14   0   9  NA  NA   NA
## 4 15      NA  4.35  NA  20   NA   0  1080   1 257  NA  NA   NA
## 5  0      NA 10.00  NA   0   NA   0    57   0  21  NA  NA   NA
## 6  0      NA  0.00  NA   0   NA   0     3   1   0  NA  NA   NA
```

*#Grouped by playerID like we did for the batters to get the career stats*

*#Summed up the total stats that we wanted*

*#Dropped the NA values*

```
pitchers1 <- Pitching%>%
  group_by(playerID)%>%
  summarise(seasons = n(), ER = sum(ER), SO = sum(SO), OUTS = sum(IPouts))%>%
  drop_na()
head(pitchers1)
```

```
## # A tibble: 6 x 5
##   playerID seasons   ER   SO  OUTS
##   <chr>      <int> <int> <int> <int>
## 1 aardsda01      9  160  340  1011
## 2 aasedo01     13  468  641  3328
## 3 abadfe01     10  135  280   992
## 4 abbeybe01      6  285  161  1704
## 5 abbeych01      1    1    0     6
## 6 abbotda01      1    9    1    39
```

*#Joined the height/weight data with the batting statistics*

```
batters2 <- batters1%>%
```

```

  inner_join(players1, by = "playerID")
head(batters2)

## # A tibble: 6 x 18
##   playerID    AB      H   DBL   TRP   HR   RBI   SB   SO   BB
##   <chr>      <int> <int> <int> <int> <int> <int> <int> <int> <int>
## 1 aardsda01     4     0     0     0     0     0     0     2     0
## 2 aaronha01 12364  3771   624    98   755  2297   240  1383  1402
## 3 aaronto01   944   216    42     6    13    94     9   145    86
## 4 aasedo01     5     0     0     0     0     0     0     3     0
## 5 abadan01    21     2     0     0     0     0     0     5     4
## 6 abadfe01     9     1     0     0     0     0     0     5     0
## # ... with 7 more variables: birthCountry <chr>, birthState <chr>,
## #   nameGiven <chr>, weight <int>, height <int>, debut <chr>, bats <fct>

```

*#Did the same with the pitching statistics*

```

pitchers2 <- pitchers1%>%
  inner_join(players1, by = "playerID")
head(pitchers2)

## # A tibble: 6 x 13
##   playerID seasons      ER      SO  OUTS birthYear birthCountry birthState
##   <chr>      <int> <int> <int> <int>      <int> <chr>      <chr>
## 1 aardsda~      9   160   340  1011      1981 USA      CO
## 2 aasedo01     13   468   641  3328      1954 USA      CA
## 3 abadfe01     10   135   280   992      1985 D.R.    La Romana
## 4 abbeybe~      6   285   161  1704      1869 USA      VT
## 5 abbeych~      1     1     0     6      1866 USA      NE
## 6 abbotda~      1     9     1    39      1862 USA      OH
## # ... with 4 more variables: weight <int>, height <int>, debut <chr>,
## #   bats <fct>

```

*#Filter out to get the players with a certain amount of at-bats*

*#Decided to use 2000 at-bats as the cutoff*

*#That way there wouldn't be players with small sample sizes*

```

#Also filtered out players that were born before 1940
#Then, calculated XBH in order to get slugging percentage
#Also calculated average and on base percentage
#Did not include RBI because we did not think it would be affected by
anything
batters3 <- batters2%>%
  filter(AB > 2000, birthYear > 1940)%>%
  mutate(XBH = DBL + TRP + HR, AVG = H/AB, OBP = (H+BB)/AB)%>%
  mutate(SLG = ((H-XBH)+(DBL*2)+(TRP*3)+(HR*4))/AB, SOpAB=SO/AB, HRpH = HR/H,
SBpH = SB/H)%>%
  select(AB, AVG, OBP, SLG, HRpH, SBpH, SOpAB, birthYear, birthCountry,
birthState, weight, height, bats)
head(batters3)

```

```

## # A tibble: 6 x 13
##      AB    AVG    OBP    SLG    HRpH    SBpH SOpAB birthYear birthCountry
birthState
##    <int> <dbl> <dbl> <dbl>    <dbl>    <dbl> <dbl>    <int> <chr>
<chr>
## 1  2044 0.256 0.321 0.423 0.119  0.0421 0.279    1969 USA      OH
## 2  8480 0.291 0.465 0.475 0.117  0.162   0.217    1974 Venezuela
Aragua
## 3  3787 0.294 0.364 0.520 0.178  0.00898 0.220    1987 Cuba
Cienfuegos
## 4  2125 0.241 0.332 0.367 0.0898 0.0605 0.197    1988 USA      NC
## 5  2385 0.259 0.327 0.467 0.191  0.00647 0.266    1988 USA      PA
## 6  3912 0.255 0.343 0.412 0.130  0.167   0.235    1942 USA      AL
## # ... with 3 more variables: weight <int>, height <int>, bats <fct>

```

```

#Gave the pitchers a lower threshold for OUTS because they don't play the
whole game
#While most of the time, starting fielders do
#Calculated ERA by multiplying the number of outs in a game by the total
earned runs and dividing by total outs
#Used the same method to calculate a pitchers strikeouts per game
#Then selected the columns we wanted to use
pitchers3 <- pitchers2%>%
  filter(OUTS > 1800, birthYear > 1940)%>%
  mutate(ERA = (27*ER)/OUTS, SOpG = (27*SO)/OUTS)%>%
  select(OUTS, ERA, SOpG, birthYear, birthCountry, birthState, weight,
height)
head(pitchers3)

```

```

## # A tibble: 6 x 8
##      OUTS    ERA    SOpG birthYear birthCountry birthState weight height
##    <int> <dbl> <dbl>    <int> <chr>        <chr>    <int> <int>
## 1  3328  3.80  5.20    1954 USA         CA        190    75
## 2  3858  4.39  3.39    1951 USA         AR        200    78
## 3  5022  4.25  4.77    1967 USA         MI        200    75
## 4  2162  4.92  6.19    1967 USA         CA        185    75

```

```
## 5 2713 3.97 4.80 1958 USA TX 210 74
## 6 2608 4.17 7.15 1973 USA AL 180 75
```

*#Set the seed for the sample and got samples of all the batters height/weight and hitting stats*

*#Then went through and compared to see if there was any good fitting line in the linear regression models*

*#Color coordinated by making weight with the orange dashed line and height with the blue*

*#Also color coordinated the hitting stats*

```
set.seed(51321)
```

```
n <- 150
```

```
samp_batter_height <- sample(batters3$height,n)
```

```
samp_batter_weight <- sample(batters3$weight,n)
```

```
samp_batter_avg <- sample(batters3$AVG,n)
```

```
samp_batter_obp <- sample(batters3$OBP,n)
```

```
samp_batter_slg <- sample(batters3$SLG,n)
```

```
samp_batter_hr <- sample(batters3$HRpH,n)
```

```
samp_batter_sb <- sample(batters3$SBpH,n)
```

```
samp_batter_so <- sample(batters3$SOpAB,n)
```

```
fav_stats(batters3$AVG)
```

```
##      min      Q1    median      Q3      max      mean      sd
n
## 0.1941676 0.2516878 0.2643683 0.2772197 0.3381783 0.2648844 0.01906285
1248
## missing
##      0
```

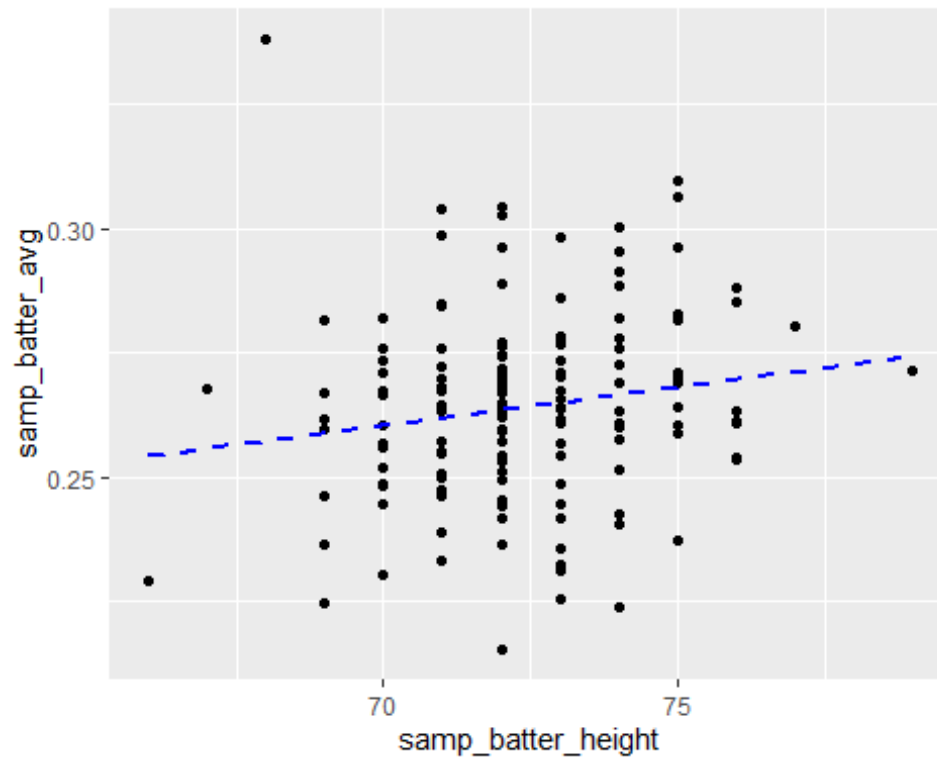
```
avg_height <- lm(samp_batter_avg ~ samp_batter_height)
```

```
msummary(avg_height)
```

```
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.1519203  0.0556311   2.731  0.00708 **
## samp_batter_height 0.0015481  0.0007684   2.015  0.04573 *
##
## Residual standard error: 0.01903 on 148 degrees of freedom
## Multiple R-squared:  0.0267, Adjusted R-squared:  0.02012
## F-statistic: 4.06 on 1 and 148 DF, p-value: 0.04573
```

```
gf_point(samp_batter_avg ~ samp_batter_height)%>%
```

```
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```

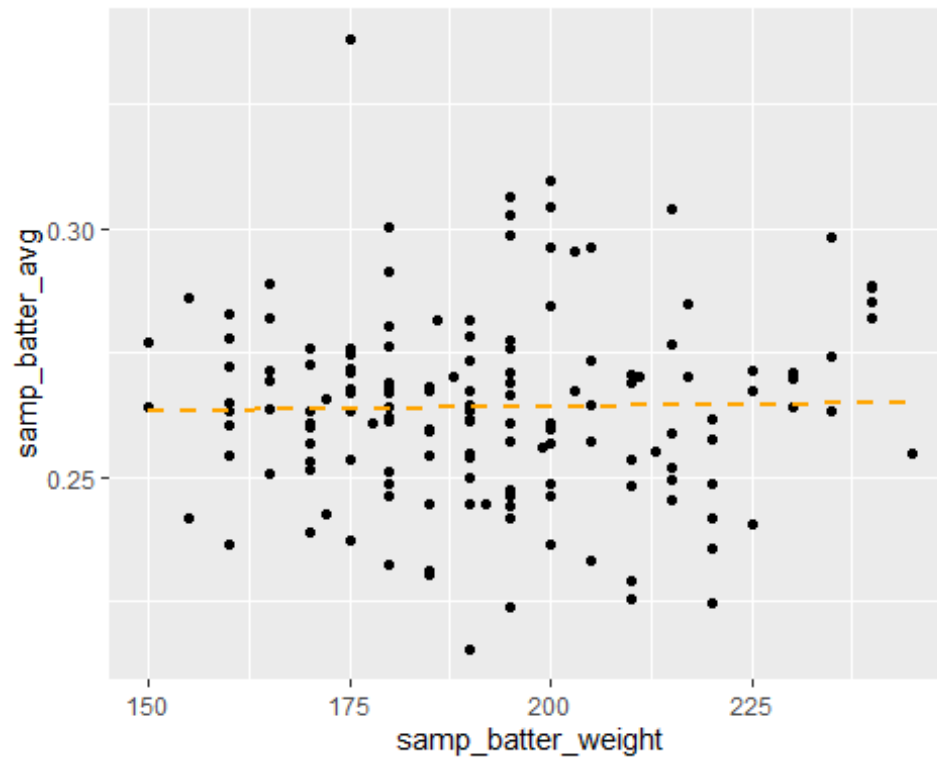


```
avg_weight <- lm(samp_batter_avg ~ samp_batter_weight)
msummary(avg_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    2.606e-01  1.423e-02  18.310   <2e-16 ***
## samp_batter_weight 1.755e-05  7.354e-05   0.239    0.812
##
## Residual standard error: 0.01928 on 148 degrees of freedom
## Multiple R-squared:  0.0003849, Adjusted R-squared:  -0.006369
## F-statistic: 0.05698 on 1 and 148 DF,  p-value: 0.8117

gf_point(samp_batter_avg ~ samp_batter_weight)%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```

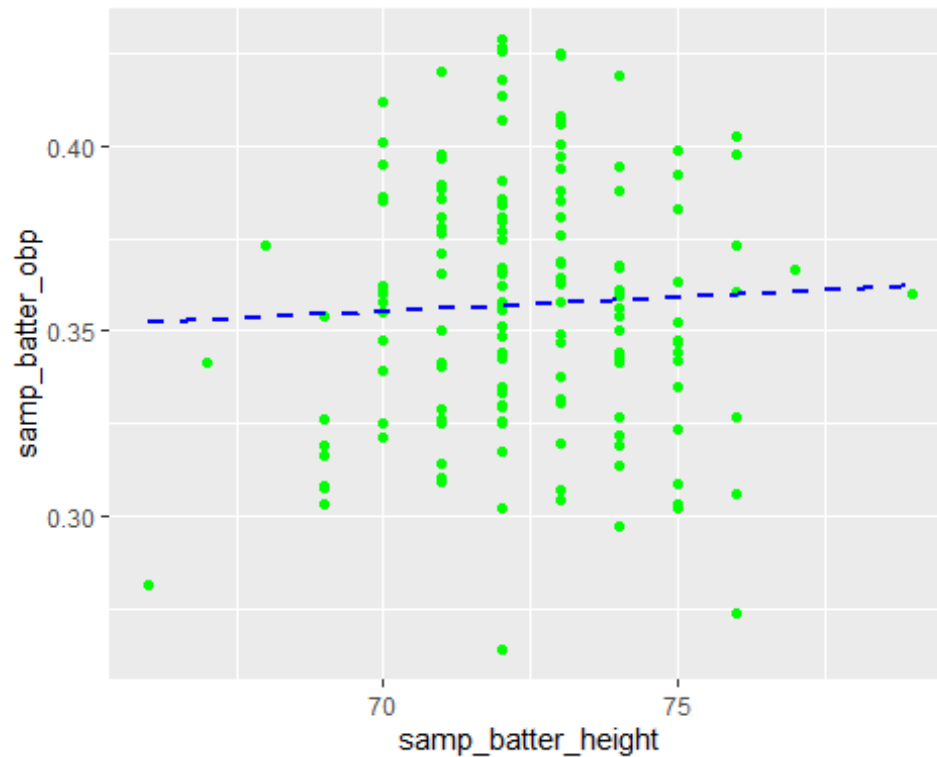




```
obp_height <- lm(samp_batter_obp ~ samp_batter_height)
msummary(obp_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.302008   0.100898   2.993  0.00324 **
## samp_batter_height 0.000761   0.001394   0.546  0.58585
##
## Residual standard error: 0.03451 on 148 degrees of freedom
## Multiple R-squared:  0.002011,    Adjusted R-squared:  -0.004732
## F-statistic: 0.2982 on 1 and 148 DF,  p-value: 0.5858

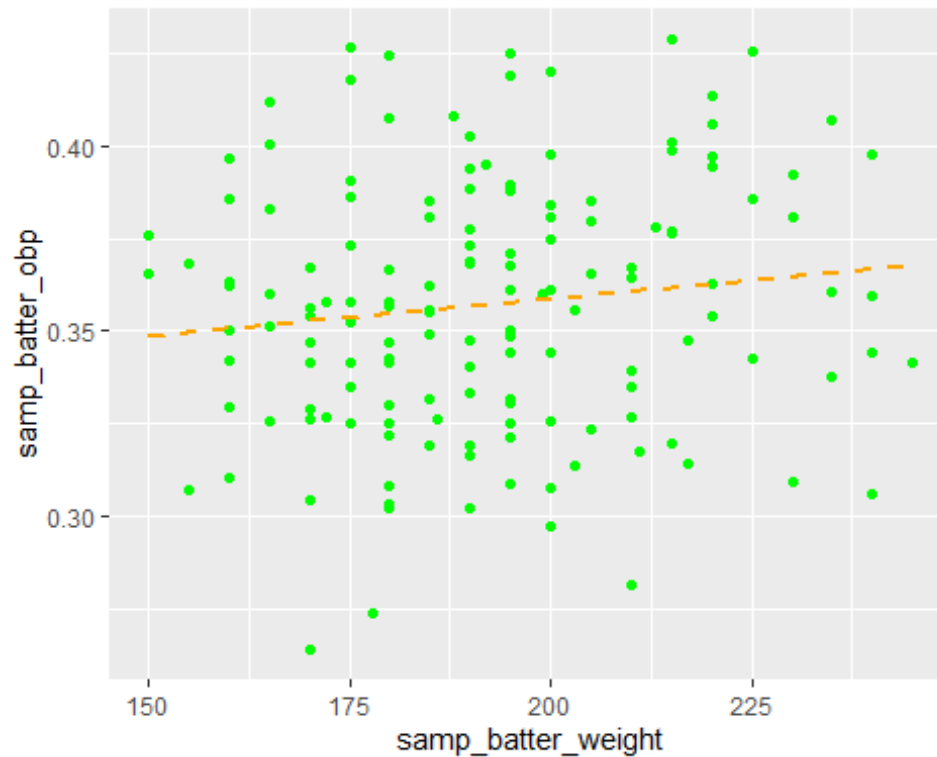
gf_point(samp_batter_obp ~ samp_batter_height, color = "green")%>%
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```



```
obp_weight <- lm(samp_batter_obp ~ samp_batter_weight)
msummary(obp_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.3186746  0.0252970  12.597  <2e-16 ***
## samp_batter_weight 0.0001997  0.0001307   1.528   0.129
##
## Residual standard error: 0.03427 on 148 degrees of freedom
## Multiple R-squared:  0.01552,    Adjusted R-squared:  0.008872
## F-statistic: 2.334 on 1 and 148 DF,  p-value: 0.1287

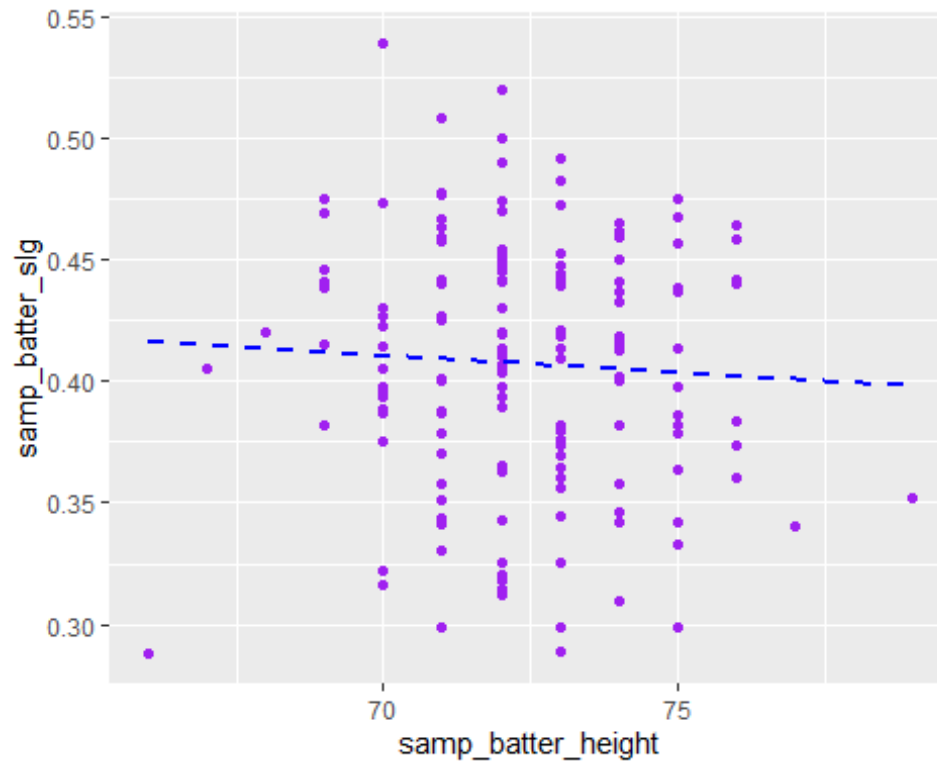
gf_point(samp_batter_obp ~ samp_batter_weight, color = "green")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```



```
slg_height <- lm(samp_batter_slg ~ samp_batter_height)
msummary(slg_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.508403   0.152629   3.331  0.00109 **
## samp_batter_height -0.001398   0.002108  -0.663  0.50821
##
## Residual standard error: 0.0522 on 148 degrees of freedom
## Multiple R-squared:  0.002963,    Adjusted R-squared:  -0.003773
## F-statistic: 0.4399 on 1 and 148 DF,  p-value: 0.5082

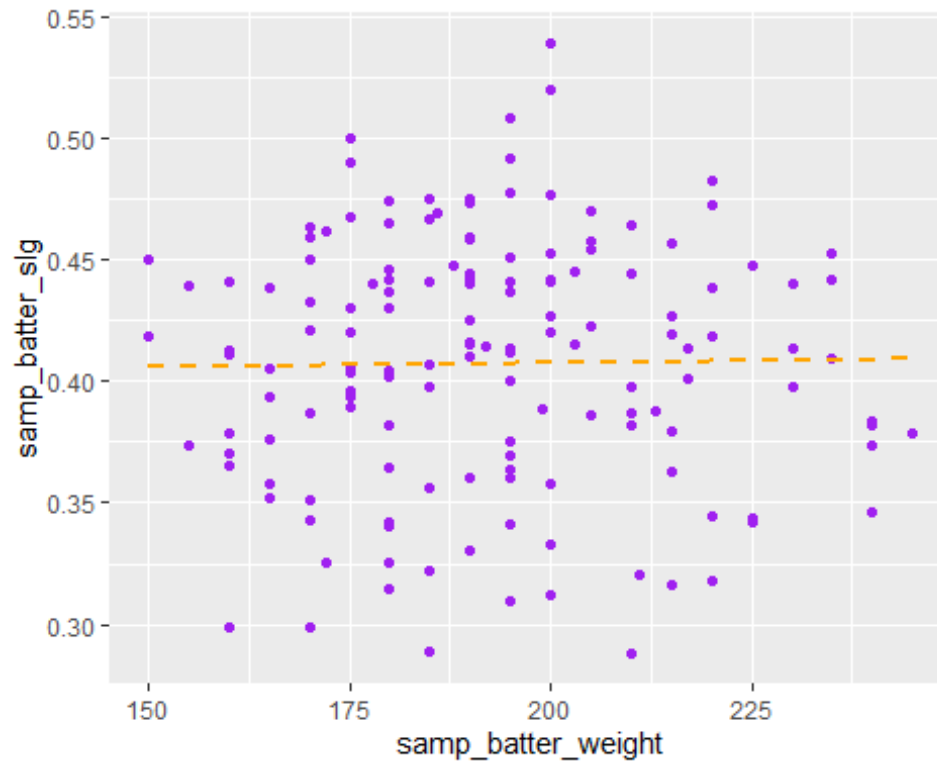
gf_point(samp_batter_slg ~ samp_batter_height, color = "purple")%>%
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```



```
slg_weight <- lm(samp_batter_slg ~ samp_batter_weight)
msummary(slg_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    4.004e-01  3.858e-02  10.379  <2e-16 ***
## samp_batter_weight 3.517e-05  1.994e-04   0.176    0.86
##
## Residual standard error: 0.05227 on 148 degrees of freedom
## Multiple R-squared:  0.0002102, Adjusted R-squared:  -0.006545
## F-statistic: 0.03112 on 1 and 148 DF,  p-value: 0.8602

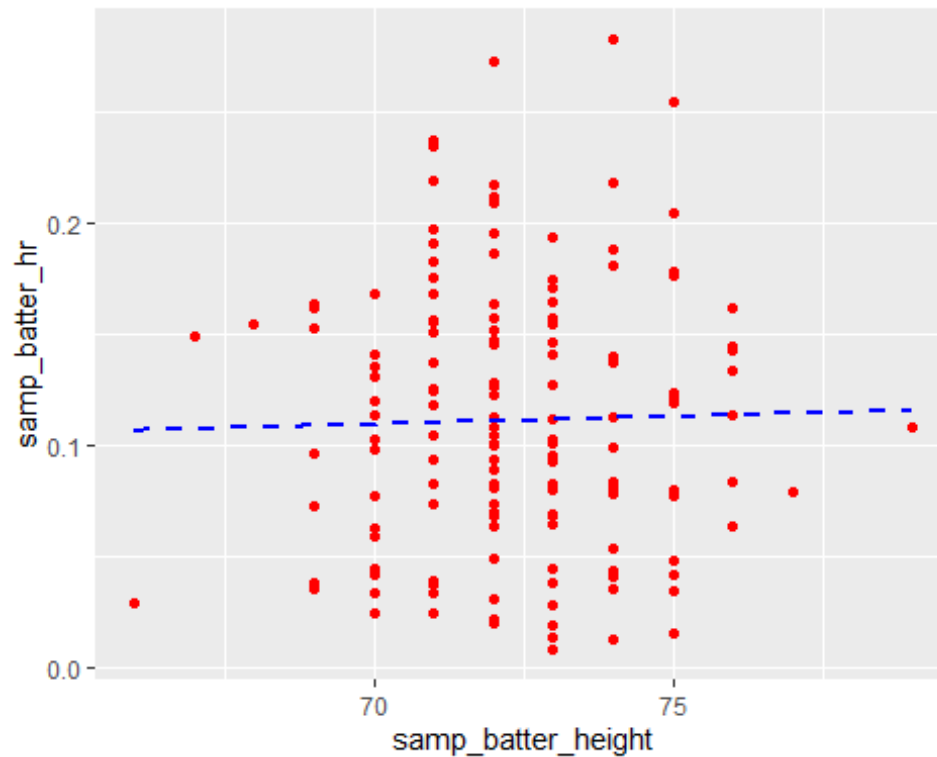
gf_point(samp_batter_slg ~ samp_batter_weight, color = "purple")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```



```
hr_height <- lm(samp_batter_hr ~ samp_batter_height)
msummary(hr_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.0607984  0.1793294   0.339   0.735
## samp_batter_height 0.0006967  0.0024769   0.281   0.779
##
## Residual standard error: 0.06133 on 148 degrees of freedom
## Multiple R-squared:  0.0005344, Adjusted R-squared:  -0.006219
## F-statistic: 0.07913 on 1 and 148 DF,  p-value: 0.7789

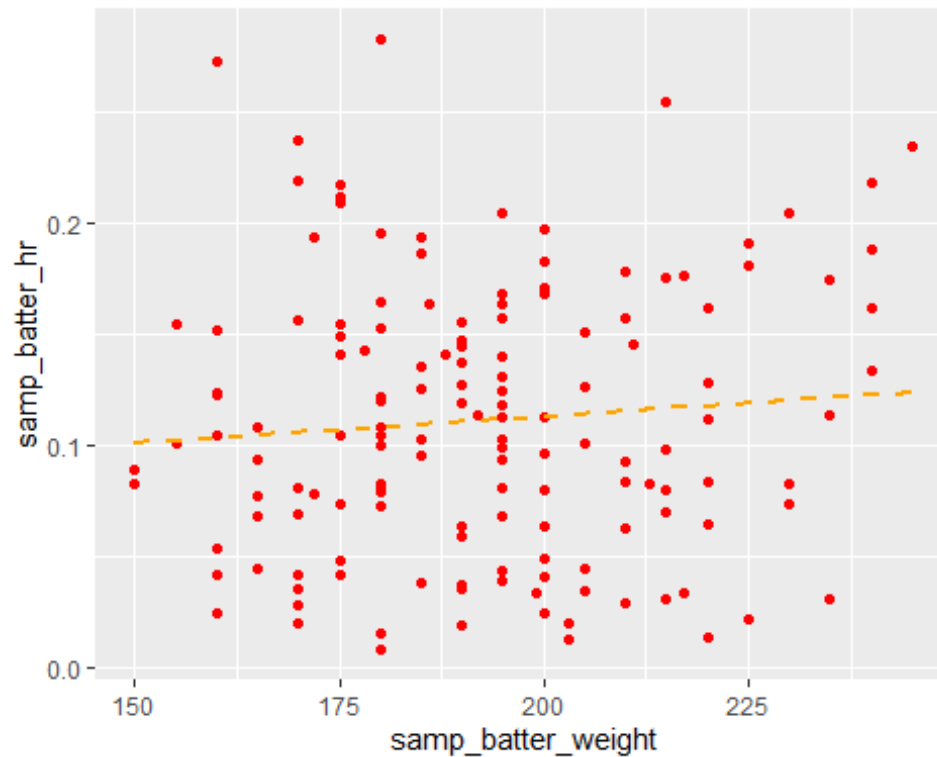
gf_point(samp_batter_hr ~ samp_batter_height, color = "red")%%
gf_lm(size = 1, color = "blue", linetype = "dashed")
```



```
hr_weight <- lm(samp_batter_hr ~ samp_batter_weight)
msummary(hr_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.0607984  0.1793294   0.339   0.735
## samp_batter_height 0.0006967  0.0024769   0.281   0.779
##
## Residual standard error: 0.06133 on 148 degrees of freedom
## Multiple R-squared:  0.0005344, Adjusted R-squared:  -0.006219
## F-statistic: 0.07913 on 1 and 148 DF,  p-value: 0.7789

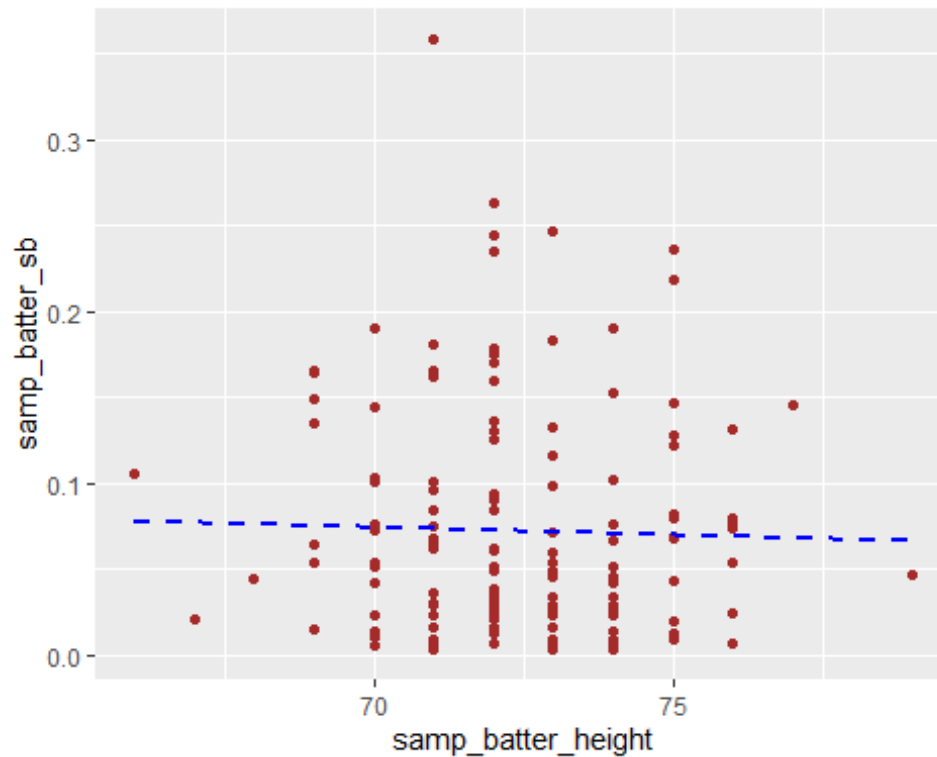
gf_point(samp_batter_hr ~ samp_batter_weight, color = "red")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```



```
sb_height <- lm(samp_batter_sb ~ samp_batter_height)
msummary(sb_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.1371409  0.1937783   0.708   0.480
## samp_batter_height -0.0009011  0.0026764  -0.337   0.737
##
## Residual standard error: 0.06627 on 148 degrees of freedom
## Multiple R-squared:  0.0007653, Adjusted R-squared:  -0.005986
## F-statistic: 0.1134 on 1 and 148 DF,  p-value: 0.7368

gf_point(samp_batter_sb ~ samp_batter_height, color = "brown")%>%
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```

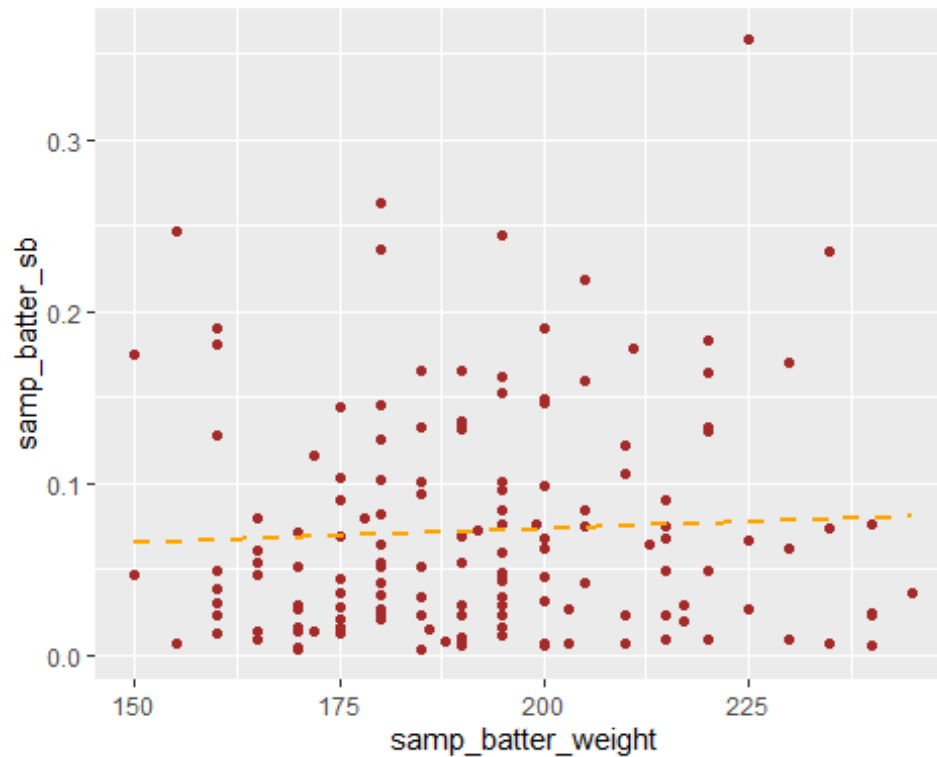


```
sb_weight <- lm(samp_batter_sb ~ samp_batter_weight)
msummary(sb_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.0408981  0.0488677   0.837   0.404
## samp_batter_weight 0.0001613  0.0002525   0.639   0.524
##
## Residual standard error: 0.0662 on 148 degrees of freedom
## Multiple R-squared:  0.00275,    Adjusted R-squared:  -0.003988
## F-statistic: 0.4081 on 1 and 148 DF,  p-value: 0.5239

gf_point(samp_batter_sb ~ samp_batter_weight, color = "brown")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```

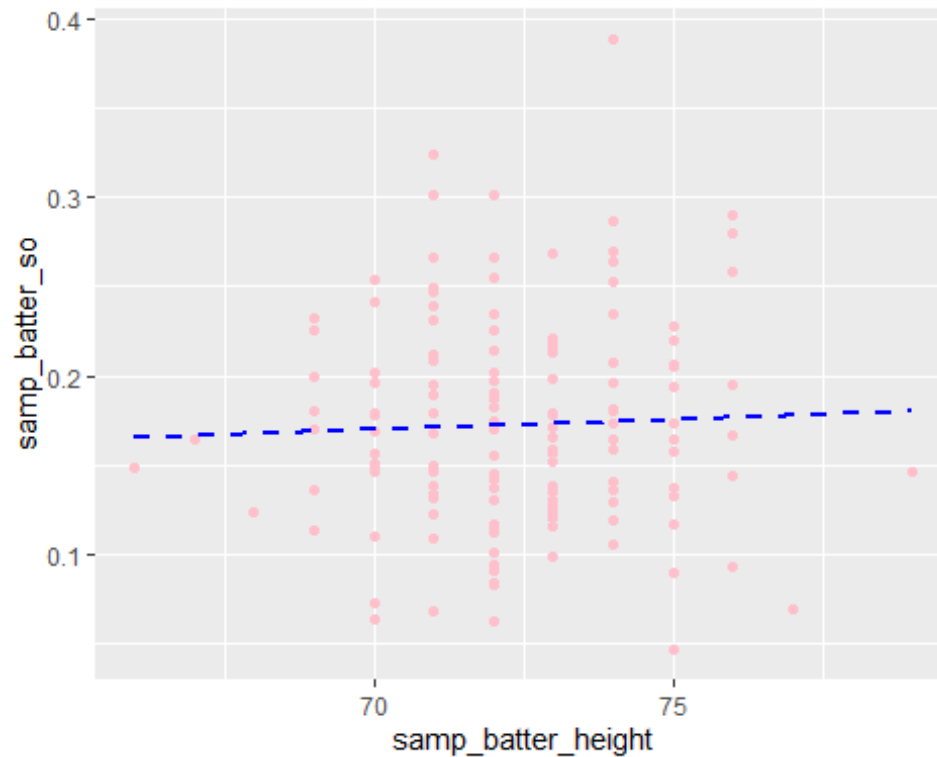




```
soH_height <- lm(samp_batter_so ~ samp_batter_height)
msummary(soH_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    0.091069   0.173172   0.526   0.600
## samp_batter_height 0.001123   0.002392   0.470   0.639
##
## Residual standard error: 0.05922 on 148 degrees of freedom
## Multiple R-squared:  0.001488,    Adjusted R-squared:  -0.005259
## F-statistic: 0.2206 on 1 and 148 DF,  p-value: 0.6393

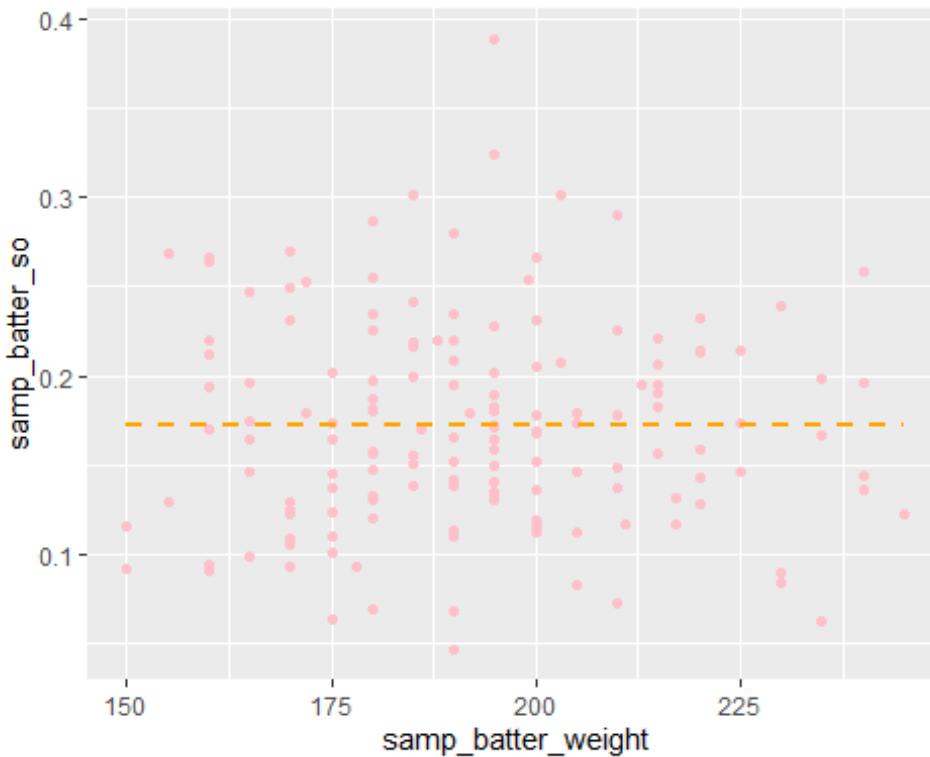
gf_point(samp_batter_so ~ samp_batter_height, color = "pink")%>%
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```



```
soH_weight <- lm(samp_batter_so ~ samp_batter_weight)
msummary(soH_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)    1.739e-01  4.375e-02   3.975  0.00011 ***
## samp_batter_weight -7.845e-06  2.260e-04  -0.035  0.97236
##
## Residual standard error: 0.05927 on 148 degrees of freedom
## Multiple R-squared:  8.138e-06, Adjusted R-squared:  -0.006749
## F-statistic: 0.001204 on 1 and 148 DF, p-value: 0.9724

gf_point(samp_batter_so ~ samp_batter_weight, color = "pink")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```



*#Now we do the same as above except with the pitchers data  
#We dont expect to get any significant results after seeing the hitting graphs*

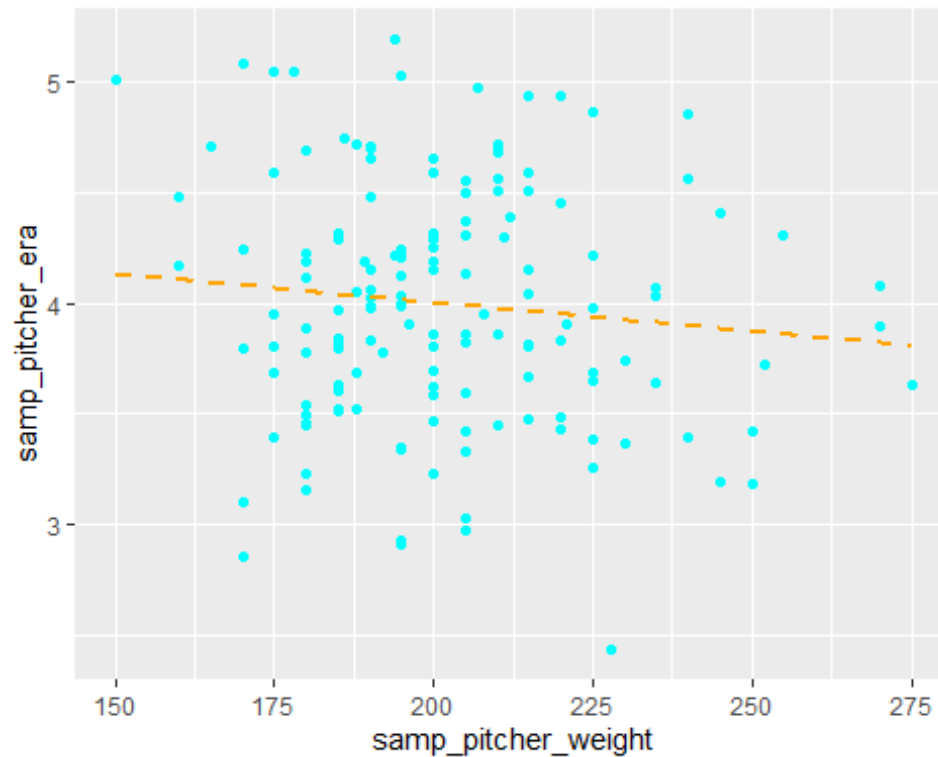
```
set.seed(51321)
n <- 150
samp_pitcher_height <- sample(pitchers3$height,n)
samp_pitcher_weight <- sample(pitchers3$weight,n)

samp_pitcher_era <- sample(pitchers3$ERA, n)
samp_pitcher_so <- sample(pitchers3$SOpG, n)

era_weight <- lm(samp_pitcher_era ~ samp_pitcher_weight)
msummary(era_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      4.519340   0.397582  11.367   <2e-16 ***
## samp_pitcher_weight -0.002588   0.001957  -1.322    0.188
##
## Residual standard error: 0.5409 on 148 degrees of freedom
## Multiple R-squared:  0.01167,    Adjusted R-squared:  0.004994
## F-statistic: 1.748 on 1 and 148 DF,  p-value: 0.1882

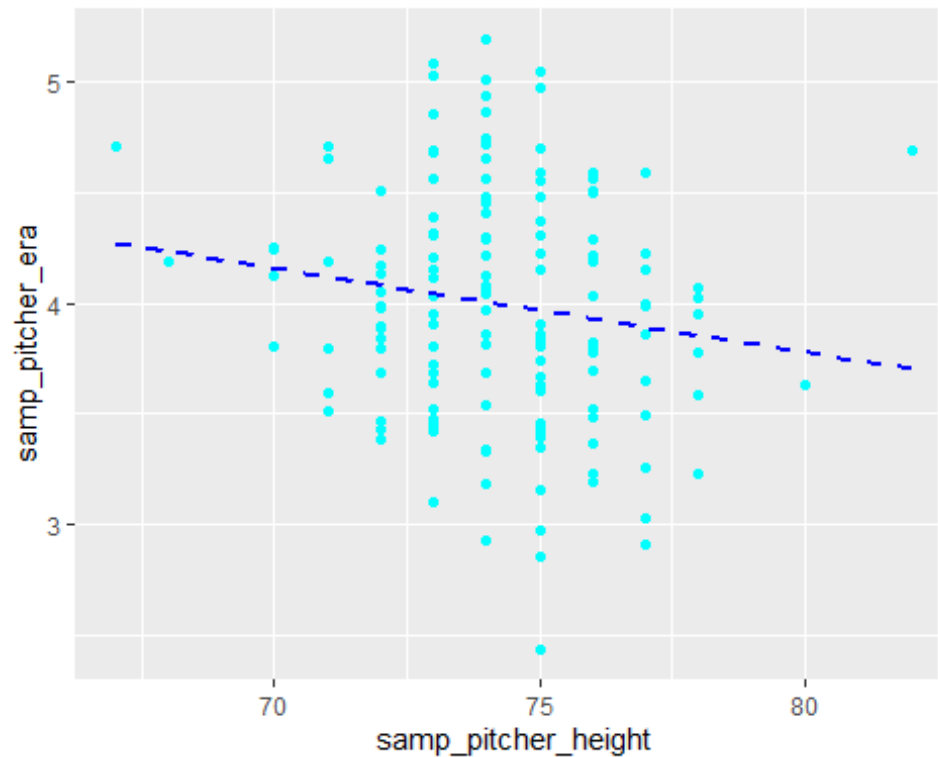
gf_point(samp_pitcher_era ~ samp_pitcher_weight, color = "cyan")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```



```
era_height <- lm(samp_pitcher_era ~ samp_pitcher_weight)
msummary(era_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      6.81856    1.53201   4.451 1.67e-05 ***
## samp_pitcher_weight -0.03802    0.02063  -1.843  0.0674 .
##
## Residual standard error: 0.5379 on 148 degrees of freedom
## Multiple R-squared:  0.02242,    Adjusted R-squared:  0.01582
## F-statistic: 3.395 on 1 and 148 DF,  p-value: 0.0674

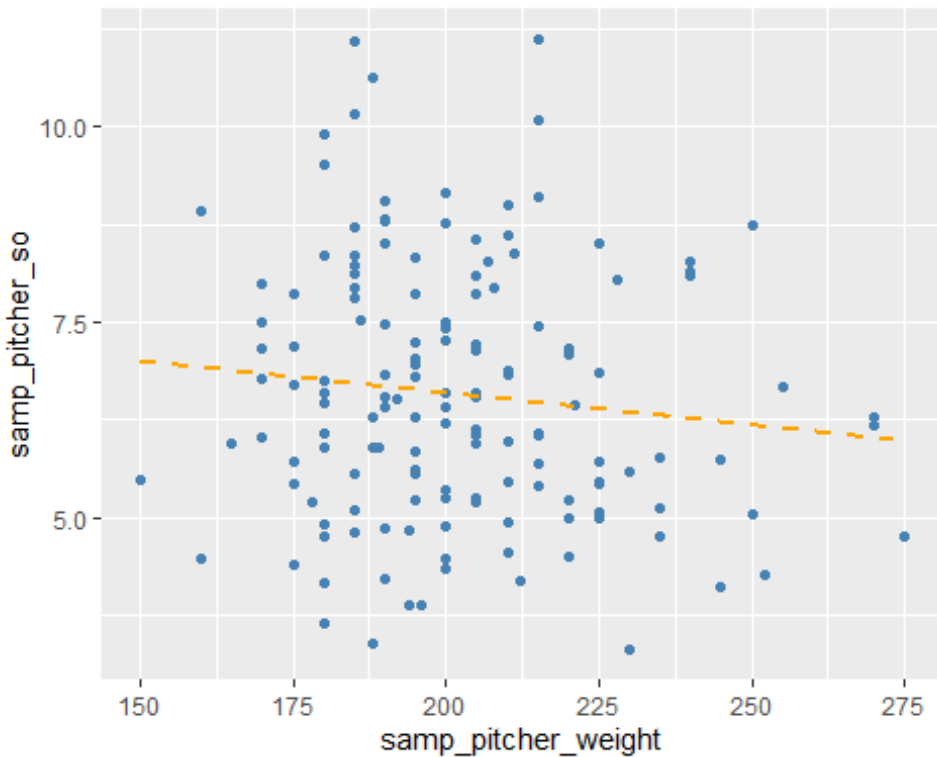
gf_point(samp_pitcher_era ~ samp_pitcher_weight, color = "cyan")%>%
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```



```
soP_weight <- lm(samp_pitcher_so ~ samp_pitcher_weight)
msummary(soP_weight)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      8.224363   1.202268   6.841 1.94e-10 ***
## samp_pitcher_weight -0.008174   0.005919  -1.381   0.169
##
## Residual standard error: 1.636 on 148 degrees of freedom
## Multiple R-squared:  0.01272,    Adjusted R-squared:  0.006053
## F-statistic: 1.907 on 1 and 148 DF,  p-value: 0.1693

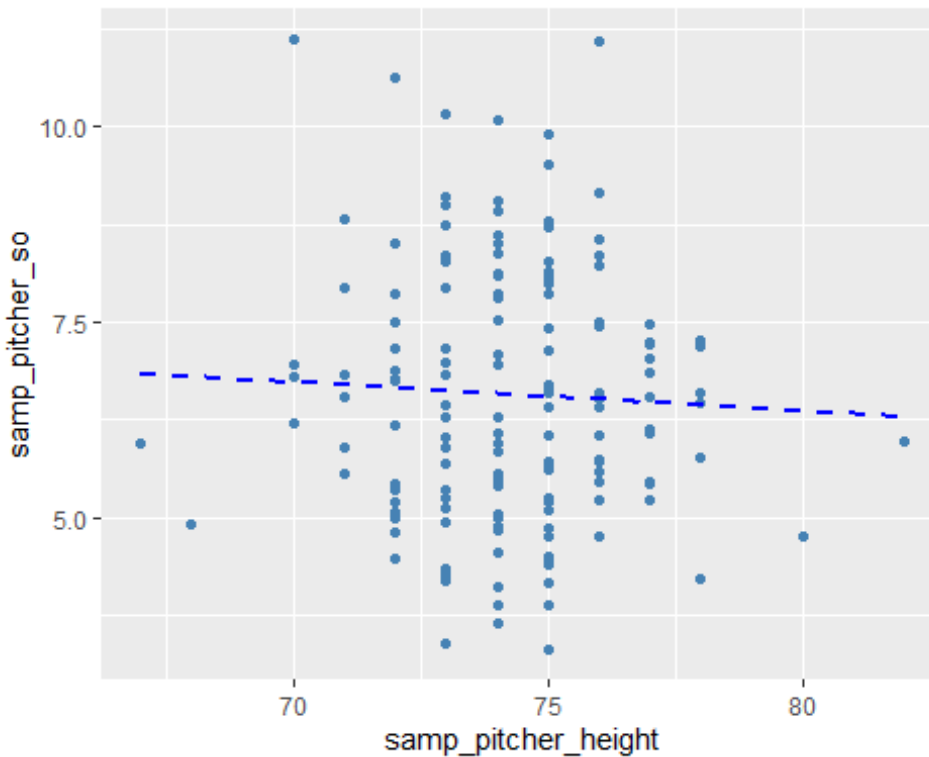
gf_point(samp_pitcher_so ~ samp_pitcher_weight, color = "steelblue")%>%
  gf_lm(size = 1, color = "orange", linetype = "dashed")
```



```
soP_height <- lm(samp_pitcher_so ~ samp_pitcher_height)
msummary(soP_height)

##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      9.28859    4.68273   1.984  0.0492 *
## samp_pitcher_height -0.03657    0.06307  -0.580  0.5629
##
## Residual standard error: 1.644 on 148 degrees of freedom
## Multiple R-squared:  0.002267,    Adjusted R-squared:  -0.004474
## F-statistic: 0.3363 on 1 and 148 DF,  p-value: 0.5629

gf_point(samp_pitcher_so ~ samp_pitcher_height, color = "steelblue")%>%
  gf_lm(size = 1, color = "blue", linetype = "dashed")
```



*#None of the linear regressions really showed a strong correlation*

*#Decided to stick with the ones that showed the most:*

*#Batter Avg vs Height*

*#Pitcher ERA vs Height*

*#Wanted to see if running an accuracy test and decision tree on Avg and Era would provide any interesting results*

*#Started with avg but had to make it a factor*

*#TRUE will be a good hitter ( $\geq .250$ )*

*#FALSE will be a bad hitter ( $< .250$ )*

```
batters4 <- batters3%>%
  mutate(good_hitter = ifelse(AVG >= 0.264,TRUE,FALSE))%>%
  select(good_hitter, birthCountry, birthState, weight, height, bats)%>%
  mutate(good_hitter = as.factor(good_hitter))
head(batters4)
```

## # A tibble: 6 x 6

	good_hitter	birthCountry	birthState	weight	height	bats
	<fct>	<chr>	<chr>	<int>	<int>	<fct>
## 1	FALSE	USA	OH	180	71	R
## 2	TRUE	Venezuela	Aragua	220	72	L
## 3	TRUE	Cuba	Cienfuegos	250	75	R
## 4	FALSE	USA	NC	205	73	L
## 5	FALSE	USA	PA	245	75	L
## 6	FALSE	USA	AL	195	71	R

```

#Split the hitters into training and test data sets with the proportion at 75%
nrow(batters4)

## [1] 1248

set.seed(51321)
split_batters <- batters4%>%
  initial_split(prop = 0.75)
train_bat <- split_batters%>%
  training()
test_bat <- split_batters%>%
  testing()
list(train_bat, test_bat)%>%
  map_int(nrow)

## [1] 936 312

#Built the null model as "hitter_null"
hitter_null <- logistic_reg(mode = "classification") %>%
  set_engine("glm") %>%
  fit(good_hitter ~ 1, data = train_bat)
#Created null_hit_pred to test the accuracy of the null prediction
null_hit_pred <- train_bat%>%
  bind_cols(predict(hitter_null, new_data = train_bat, type = "class"))%>%
  rename(good_null = .pred_class)
accuracy(null_hit_pred, good_hitter, good_null)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>    <chr>        <dbl>
## 1 accuracy binary      0.505

#The null model seems to be a little over 50% accuracy
#Now we want to see if we can improve it

#Built the first model with only height as a variable
bat_model1 <- logistic_reg(mode = "classification")%>%
  set_engine("glm")%>%
  fit(good_hitter ~ height, data = train_bat)

bat_pred1 <- train_bat%>%
  bind_cols(predict(bat_model1, new_data = train_bat, type = "class"))%>%
  rename(hit_model1 = .pred_class)
accuracy(bat_pred1, good_hitter, hit_model1)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>    <chr>        <dbl>
## 1 accuracy binary      0.505

#The estimate is the same as the null

```



```

#Decided to add weight onto the first model
bat_model2 <- logistic_reg(mode = "classification")%>%
  set_engine("glm")%>%
  fit(good_hitter ~ height + weight, data = train_bat)

bat_pred2 <- train_bat%>%
  bind_cols(predict(bat_model2, new_data = train_bat, type = "class"))%>%
  rename(hit_model2 = .pred_class)
accuracy(bat_pred2, good_hitter, hit_model2)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 accuracy binary      0.516

#The estimate increased a little bit
#Now we're going to try to add country and state of birth because in warmer
places they can play all year

#Since height and weight didn't really help too much, we added birthCountry
and state
bat_model3 <- logistic_reg(mode = "classification")%>%
  set_engine("glm")%>%
  fit(good_hitter ~ height + weight + birthCountry + birthState, data =
train_bat)

## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

bat_pred3 <- train_bat%>%
  bind_cols(predict(bat_model3, new_data = train_bat, type = "class"))%>%
  rename(hit_model3 = .pred_class)

## Warning in predict.lm(object, newdata, se.fit, scale = 1, type = if (type
== :
## prediction from a rank-deficient fit may be misleading

accuracy(bat_pred3, good_hitter, hit_model3)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 accuracy binary      0.573

#The prediction increased by about 7% now

bat_test <- logistic_reg(mode = "classification")%>%
  set_engine("glm")%>%
  fit(good_hitter ~ height + weight + birthCountry + birthState, data =
test_bat)

```

```

## Warning: glm.fit: algorithm did not converge
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

bat_test

## parsnip model object
##
## Fit time: 61ms
##
## Call: stats::glm(formula = good_hitter ~ height + weight + birthCountry +
##   birthState, family = stats::binomial, data = data)
##
## Coefficients:
##           (Intercept)                height
##           2.021e+14                -3.266e-02
##           weight                birthCountryCAN
##           -2.896e-03                4.301e+15
##           birthCountryCuba                birthCountryD.R.
##           -4.706e+15                1.688e+13
##           birthCountryJamaica                birthCountryMexico
##           -2.021e+14                -2.021e+14
##           birthCountryNicaragua                birthCountryPanama
##           4.301e+15                4.301e+15
##           birthCountryUSA                birthCountryVenezuela
##           -2.021e+14                -2.380e+15
##           birthStateAnzoategui                birthStateAR
##           2.178e+15                4.504e+15
##           birthStateAragua                birthStateAtlantico Sur
##           6.682e+15                NA
##           birthStateAZ                birthStateBC
##           -1.547e+00                1.429e+06
##           birthStateBolivar                birthStateCA
##           6.682e+15                -1.043e+00
##           birthStateCamaguey                birthStateCarabobo
##           9.007e+15                2.178e+15
##           birthStateChiriqui                birthStateColon
##           -9.007e+15                NA
##           birthStateCT                birthStateDE
##           -4.504e+15                -8.273e-02
##           birthStateDistrito Federal                birthStateDistrito Nacional
##           2.178e+15                -2.190e+14
##           birthStateDuarte                birthStateEl Seibo
##           -2.190e+14                -2.190e+14
##           birthStateEspaillat                birthStateFL
##           -2.190e+14                -5.398e-01
##           birthStateGA                birthStateGranma
##           -3.313e-01                4.504e+15
##           birthStateHI                birthStateIA
##           -2.618e+01                -1.445e+00

```

##	birthStateIL	birthStateIN
##	-1.336e+00	-7.614e-01
##	birthStateKingston	birthStateKS
##	NA	-1.873e+00
##	birthStateKY	birthStateLA
##	-1.429e+00	2.685e+01
##	birthStateLa Habana	birthStateLa Vega
##	4.504e+15	-2.190e+14
##	birthStateLara	birthStateMA
##	2.178e+15	-2.501e+07
##	birthStateMaracay	birthStateMI
##	2.178e+15	3.111e-01
##	birthStateMiranda	birthStateMN
##	2.178e+15	2.678e+01
##	birthStateMO	birthStateMonte Cristi
##	-7.454e-01	-2.190e+14
##	birthStateMS	birthStateNC
##	3.255e-01	-7.581e-01
##	birthStateND	birthStateNE
##	2.686e+01	-7.544e-01
##	birthStateNJ	birthStateNM
##	-1.401e+00	-2.637e+01
##	birthStateNY	birthStateOH
##	-8.263e-01	-5.701e-01
##	birthStateOK	birthStateON
##	-1.483e+00	1.103e+06
##	birthStateOR	birthStatePA
##	-4.658e-01	-4.225e-01
##	birthStatePeravia	birthStateRI
##	-2.190e+14	-4.504e+15
##	birthStateSamana	birthStateSan Cristobal
##	-2.190e+14	-2.190e+14
##	birthStateSan Pedro de Macoris	birthStateSantiago
##	-2.190e+14	-2.190e+14
##	birthStateSao Paulo	birthStateSC
##	-4.706e+15	2.567e-01
##	birthStateSD	birthStateSK
##	-8.689e-01	NA
##	birthStateSonora	birthStateTN
##	NA	-1.504e+00
##	birthStateTX	birthStateVA
##	-9.026e-01	3.060e-01
##	birthStateVilla Clara	birthStateWA
##	NA	-7.151e-02
##	birthStateWI	birthStateWV
##	-8.238e-02	-2.620e+01
##	birthStateWY	birthStateZulia
##	-8.138e-01	2.178e+15
##		

## Degrees of Freedom: 311 Total (i.e. Null); 234 Residual

```

## Null Deviance:      432.4
## Residual Deviance: 347.3      AIC: 503.3

bat_test_pred <- test_bat%>%
  bind_cols(predict(bat_test, new_data = test_bat, type = "class"))%>%
  rename(hit_test = .pred_class)

## Warning in predict.lm(object, newdata, se.fit, scale = 1, type = if (type
== :
## prediction from a rank-deficient fit may be misleading

accuracy(bat_test_pred, good_hitter, hit_test)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>      <dbl>
## 1 accuracy binary      0.676

#The testing estimate is even better
#Therefore this is a pretty decent model to use for predicting good_hitters
#However, 67.6% still isn't that good of a prediction, but for only using
non-game factors, its pretty good
#But since it increased so much, we may have overfit the data

fav_stats(pitchers3$ERA)

##      min      Q1   median      Q3      max      mean      sd      n
missing
## 2.208517 3.628254 3.990284 4.357752 6.029462 3.991612 0.5588547 1033
0

#Now want to build a model for pitchers era
#Will classify a good pitcher with an era <= 4

pitchers4 <- pitchers3%>%
  mutate(good_pitcher = ifelse(ERA <= 4.00, TRUE, FALSE))%>%
  select(good_pitcher, ERA, birthCountry, weight, height)%>%
  mutate(good_pitcher = as.factor(good_pitcher))
head(pitchers4)

## # A tibble: 6 x 5
##   good_pitcher  ERA birthCountry weight height
##   <fct>      <dbl> <chr>      <int>  <int>
## 1 TRUE        3.80 USA        190    75
## 2 FALSE       4.39 USA        200    78
## 3 FALSE       4.25 USA        200    75
## 4 FALSE       4.92 USA        185    75
## 5 TRUE        3.97 USA        210    74
## 6 FALSE       4.17 USA        180    75

```

```

#Set the same seed for the pitchers data and split the training and test by 75%
nrow(pitchers4)

## [1] 1033

set.seed(51321)
split_batters <- pitchers4%>%
  initial_split(prop = 0.75)
train_pitch <- split_batters%>%
  training()
test_pitch <- split_batters%>%
  testing()
list(train_pitch, test_pitch)%>%
  map_int(nrow)

## [1] 775 258

#Built the null model as "pitcher_null"
pitcher_null <- logistic_reg(mode = "classification") %>%
  set_engine("glm") %>%
  fit(good_pitcher ~ 1, data = train_pitch)
#Created null_pitch_pred to test the accuracy of the null prediction
null_pitch_pred <- train_pitch%>%
  bind_cols(predict(pitcher_null, new_data = train_pitch, type = "class"))%>%
  rename(good_pitch_null = .pred_class)
accuracy(null_pitch_pred, good_pitcher, good_pitch_null)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>      <dbl>
## 1 accuracy binary      0.510

#The null model seems to be a little over 50% accuracy
#Now we want to see if we can improve it

pitch_model1 <- logistic_reg(mode = "classification")%>%
  set_engine("glm")%>%
  fit(good_pitcher ~ height + weight + birthCountry, data = train_pitch)

pitch_pred1 <- train_pitch%>%
  bind_cols(predict(pitch_model1, new_data = train_pitch, type = "class"))%>%
  rename(pitch1 = .pred_class)
accuracy(pitch_pred1, good_pitcher, pitch1)

## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>      <dbl>
## 1 accuracy binary      0.581

#The estimate increased by around 7%, but the estimate still isn't very strong

```

*#It was definitely easier to predict the hitters average rather than the pitchers ERA*

*#Lets see if the test set performs better*

```
pitch_test <- logistic_reg(mode = "classification")%>%
  set_engine("glm")%>%
  fit(good_pitcher ~ height + weight + birthCountry, data = test_pitch)

pitch_test_pred <- test_pitch%>%
  bind_cols(predict(pitch_test, new_data = test_pitch, type = "class"))%>%
  rename(testing_pitch = .pred_class)
accuracy(pitch_test_pred, good_pitcher, testing_pitch)
```

```
## # A tibble: 1 x 3
##   .metric .estimator .estimate
##   <chr>   <chr>       <dbl>
## 1 accuracy binary      0.585
```

*#The testing set was a tiny bit more accurate but still not as good as the hitters data*

*#Created a decision tree to see what the different variables affected*

```
form <- as.formula("good_pitcher ~ height + weight + birthCountry")
decision_tree <- decision_tree(mode = "classification")%>%
  set_engine("rpart")%>%
  fit(form, data = train_pitch)
decision_tree

## parsnip model object
##
## Fit time: 11ms
## n= 775
##
## node), split, n, loss, yval, (yprob)
##      * denotes terminal node
##
##  1) root 775 380 TRUE (0.4903226 0.5096774)
##    2) weight>=201.5 335 142 FALSE (0.5761194 0.4238806) *
##    3) weight< 201.5 440 187 TRUE (0.4250000 0.5750000)
##      6) birthCountry=D.R.,Mexico,Nicaragua,Panama,South Korea 36 14 FALSE
##         (0.6111111 0.3888889) *
##      7) birthCountry=CAN,Germany,Japan,Netherlands,USA,Venezuela 404 165
##         TRUE (0.4084158 0.5915842)
##        14) weight>=186 205 96 TRUE (0.4682927 0.5317073)
##          28) weight< 194.5 69 30 FALSE (0.5652174 0.4347826) *
##          29) weight>=194.5 136 57 TRUE (0.4191176 0.5808824) *
##          15) weight< 186 199 69 TRUE (0.3467337 0.6532663) *
```

*#Conclusion:*

*#Looking at the graphs and models that we made, you can't predict if a hitter or pitcher will be good based off*

*#physical attributes*

*#What surprised us the most was that there weren't even significant data for heavier hitters having*

*#a higher slugging percentage*

*#Also, the fact that there was even a little correlation between a hitters height, and their average surprised me*

*#Only because the taller hitters had a higher average*

*#Especially those that were 75 inches tall*

*#This could be because they walk less, with having a larger strike zone.*

*#Looking at the graph for OBP, the higher values tend to go to shorter players*

*#Which would back up the last statement*

*#As for the pitchers, it seemed that physical attributes didn't seem to affect their stats as much*

*#That could be because pitching is such a game of mechanics rather than physical status*

*#Both short pitchers such as Marcus Stroman, and formerly Tim Lincecum, throw hard*

*#While hitters who are big, tend to always be power hitters (Aaron Judge, Giancarlo Stanton, David Ortiz...)*