

# Predicting Anxiety Diagnosis from Survey Data

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```
#I am looking to find the relationship between people who have been diagnosed with anxiety  
#vs their veteran status, height and weight.  
#Anxiety diagnosis(ADANXEV) is my response variable  
#Veteran status(VETERAN3), height(HEIGHT3), and weight(WEIGHT2) are my predictor variables
```

```
#Anxiety diagnosis(ADANXEV) and Veteran status(VETERAN3) won't have outliers since they  
#are yes/no  
#I will choose to not remove any outliers because part of my project will be seeing if  
#extreme heights and weights contribute to an anxiety diagnosis
```

```
#Clean data
```

```
#Anxiety diagnosis(ADANXEV)
```

```
#Only keep people who answered Yes(1) or No(2)
```

```
data <- subset(data, ADANXEV == 1 | ADANXEV == 2)
```

```
#Veteran status(VETERAN3)
```

```
#Only keep people who answered Yes(1) or No(2)
```

```
data <- subset(data, VETERAN3 == 1 | VETERAN3 == 2)
```

```
#Height(HEIGHT3)
```

```
#Only keep people with a reported height (200-711) and (9000-9998)
```

```
data <- subset(data, (HEIGHT3 >=200 & HEIGHT3 <=711) | (HEIGHT3 >= 9000 & HEIGHT3 <= 9998))
```

```
#Convert reported heights to inches
```

```
to_in <- function(number) {
```

```
  result <-
```

```
    #convert ft/in to in
```

```
    ifelse(number >= 200 & number <= 711,  
           (number %/% 100) * 12 + (number %% 100),
```

```
           #convert 9/m/cm to cm then to in
```

```
           ifelse(number >= 9000 & number <= 9998,  
                  round(0.393701*(number %/% 1000),0),  
                  number))
```

```
  return(result)
```

```
}
```

```
#Apply the to_in function to the HEIGHT3 column
```

```
data <- data %>%
```

```
  mutate(HEIGHT = to_in(HEIGHT3))
```

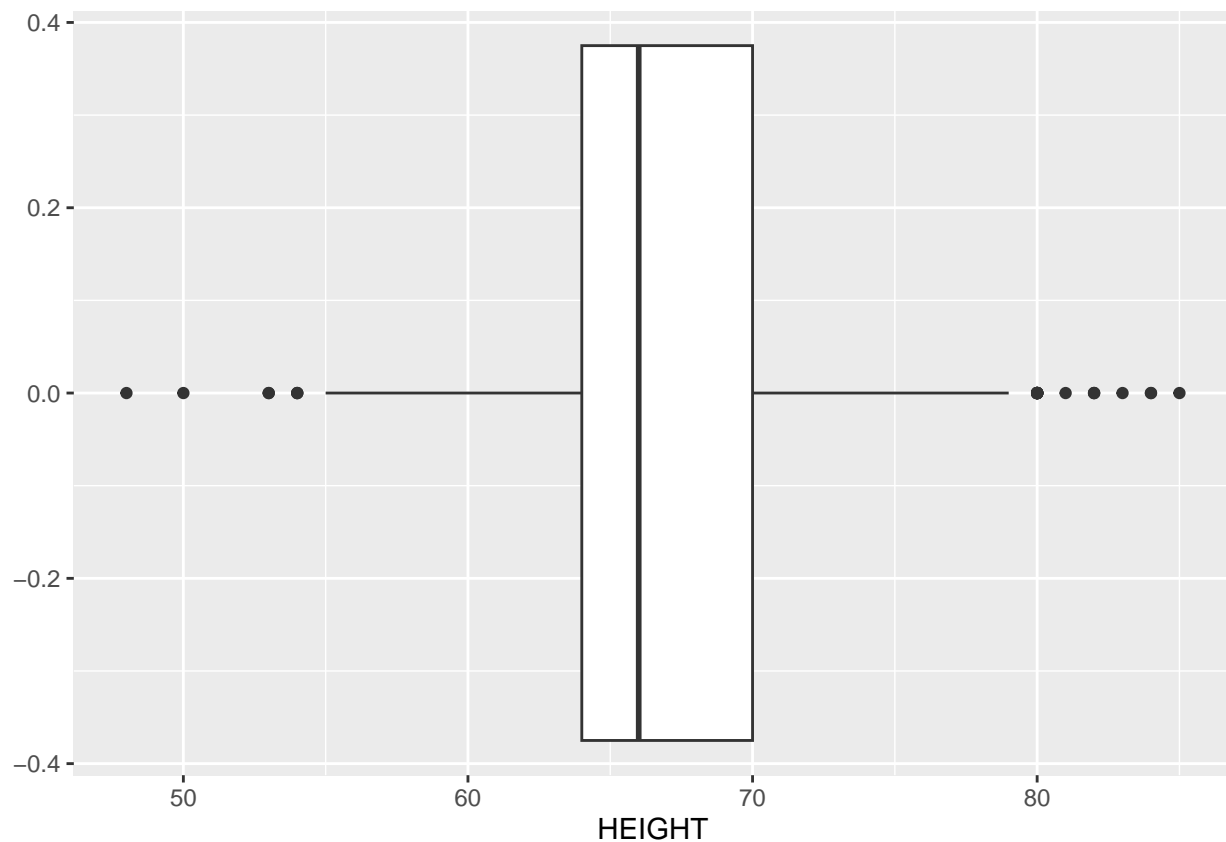
```
#Weight(WEIGHT3)
```

```
#Only keep people with a reported weight (50-0999) and (9000-9998)
```

```
data <- subset(data, (WEIGHT2 >=50 & WEIGHT2 <=999) | (WEIGHT2 >= 9000 & WEIGHT2 <= 9998))

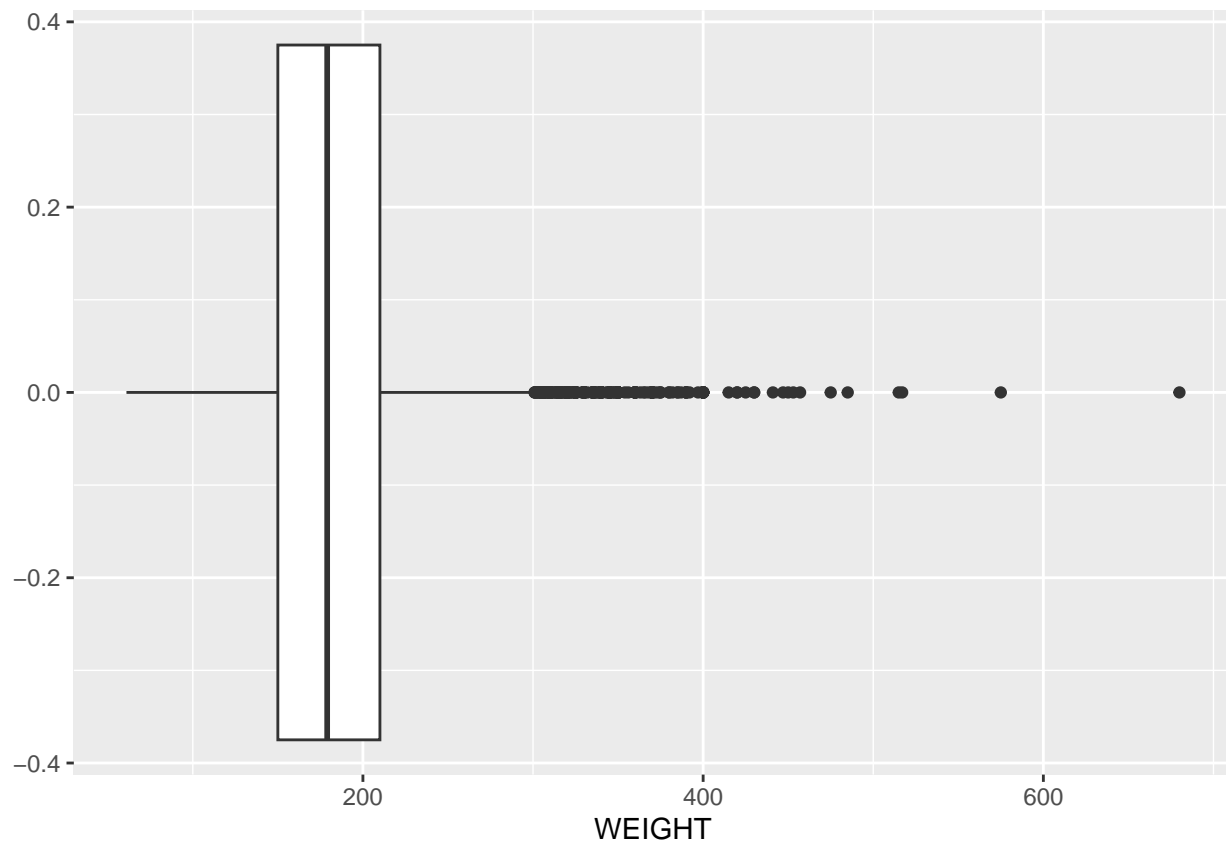
#Convert reported weights from kilograms to pounds
to_lbs <- function(number) {
  result <-
    #convert kgs to lbs
    ifelse(number >= 9000 & number <= 9998,
            round(2.20462262185*(number %% 1000),0),
            number)
  return(result)
}
#Apply the to_lbs function to the WEIGHT2 column
data <- data %>%
  mutate(WEIGHT = to_lbs(WEIGHT2))

ggplot(data) +
  geom_boxplot(mapping = aes(HEIGHT))
```



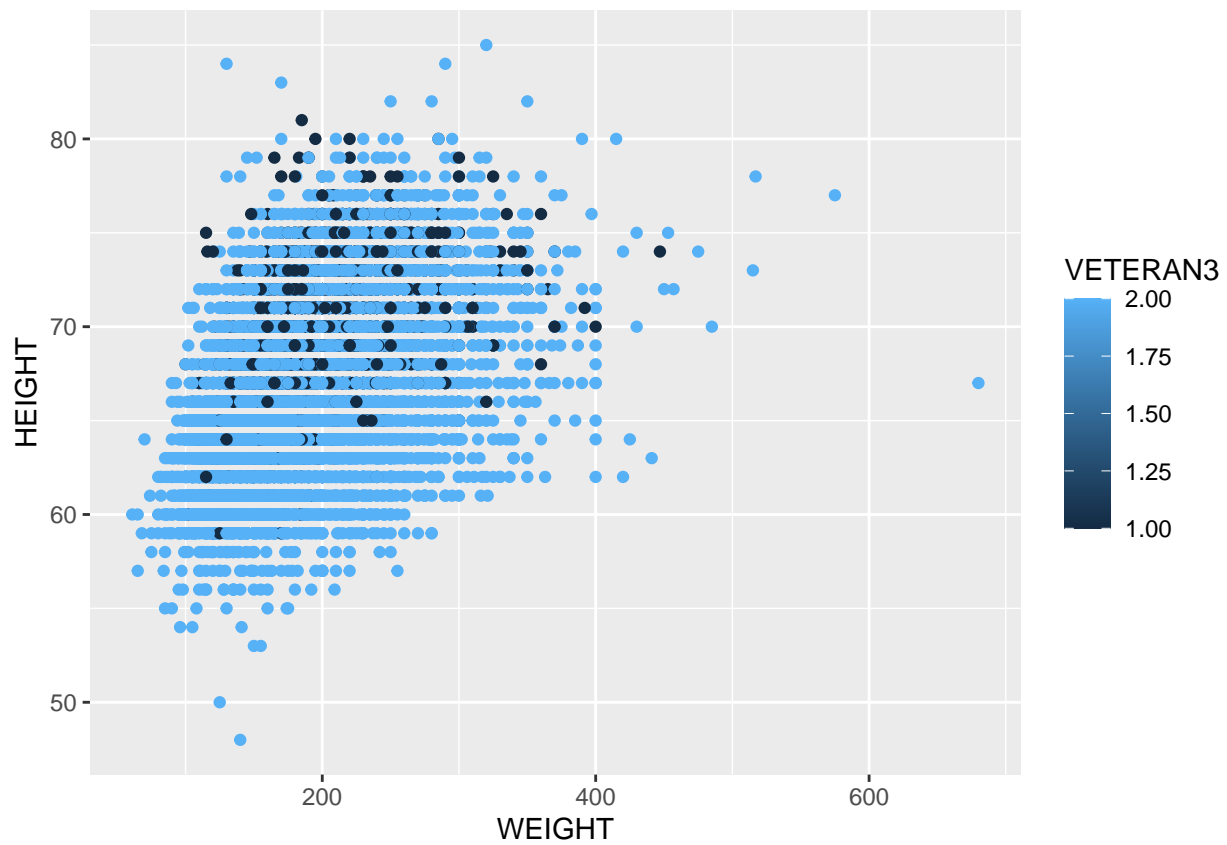
*#There are 4 low outliers and 6 higher outliers in terms of height, but overall it seems #decently normally distributed*

```
ggplot(data) +
  geom_boxplot(mapping = aes(WEIGHT))
```



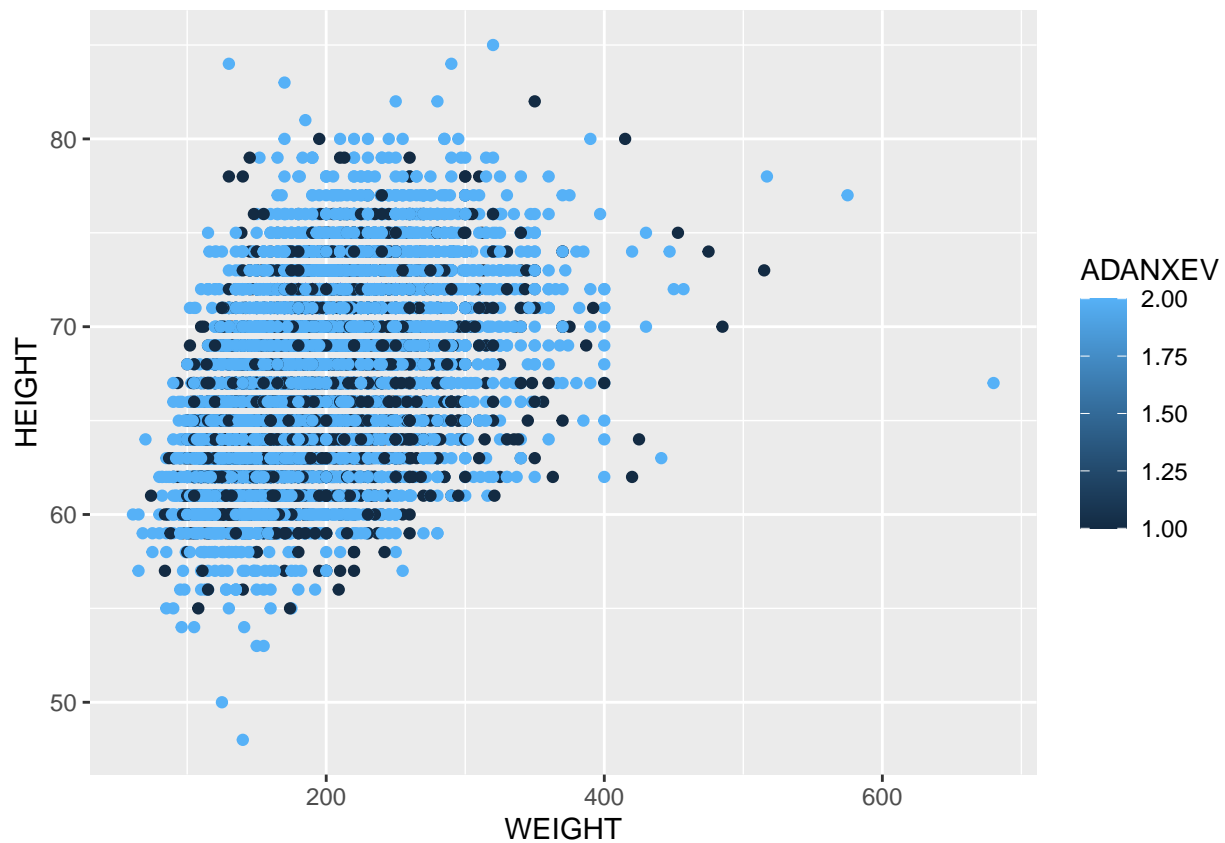
*#The weight distribution is right-skewed*

```
ggplot(data) +  
  geom_point(aes(x = WEIGHT, y = HEIGHT, color = VETERAN3))
```



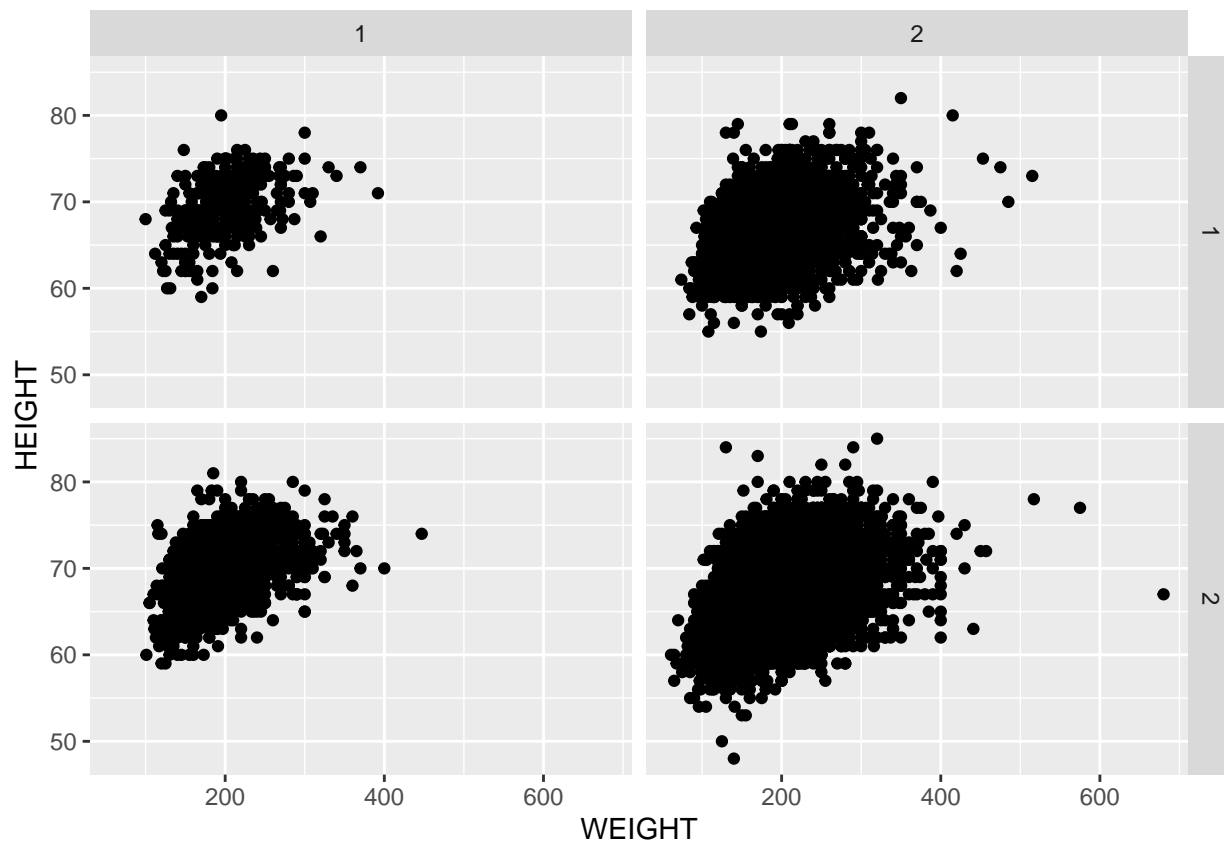
*#This plot shows people's height/weight measurement color coded by their Veteran status  
#Black is a veteran, blue is not  
#Veterans' height/weight measurements are mostly in the middle of the plot*

```
ggplot(data) +  
  geom_point(aes(x = WEIGHT, y = HEIGHT, color = ADANXEV))
```



*#This plot shows people's height/weight measurement color coded by their anxiety diagnosis  
 #Black has anxiety, blue does not  
 #Initially, it does seem like there are more people with anxiety in the bottom right area  
 #This shows roughly that shorter people and heavier people tend to have anxiety more often  
 #than taller and/or light people*

```
ggplot(data = data) +  
  geom_point(mapping = aes(x = WEIGHT, y = HEIGHT)) +  
  facet_grid(ADANXEV ~ VETERAN3)
```

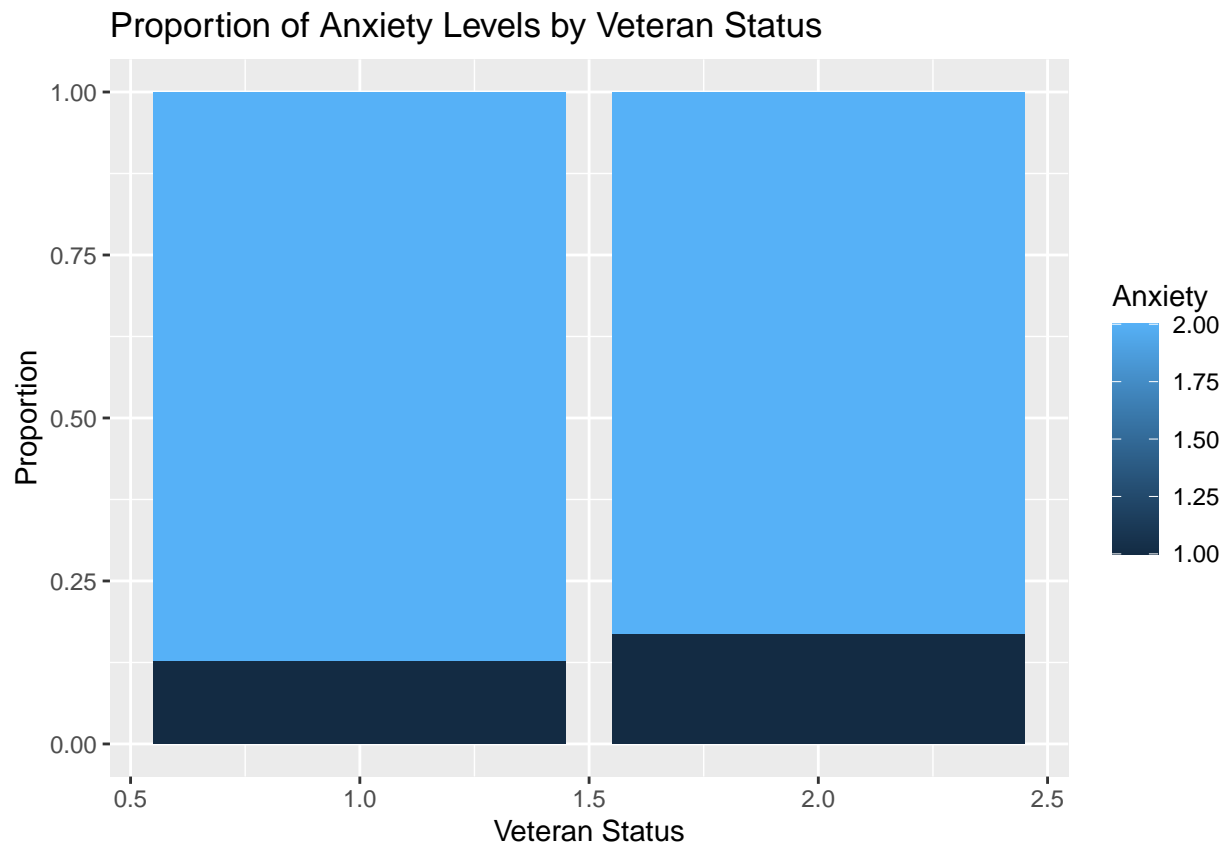


*#1 on the right = anxiety, 2 on the right = no anxiety diagnosis  
 #1 on top = veteran, 2 on top = not a veteran  
 #This confirms that veterans look much more alike each other in terms of height/weight  
 #than non-veterans  
 #It also shows that anxiety diagnoses are slightly more spread out from the big cluster  
 #than non-anxiety diagnoses*

```
#Calculate the proportion of anxiety diagnoses within each veteran status
summary_data <- data %>%
  group_by(VETERAN3, ADANXEV) %>%
  summarise(proportion = n()) %>%
  ungroup() %>%
  mutate(proportion = proportion / sum(proportion))
```

```
## `summarise()` has grouped output by 'VETERAN3'. You can override using the
## `.groups` argument.
```

```
# Create a percent stacked bar plot
ggplot(summary_data, aes(x = VETERAN3, y = proportion, fill = ADANXEV)) +
  geom_bar(position = "fill", stat = "identity") +
  labs(x = "Veteran Status", y = "Proportion", fill = "Anxiety") +
  ggtitle("Proportion of Anxiety Levels by Veteran Status")
```

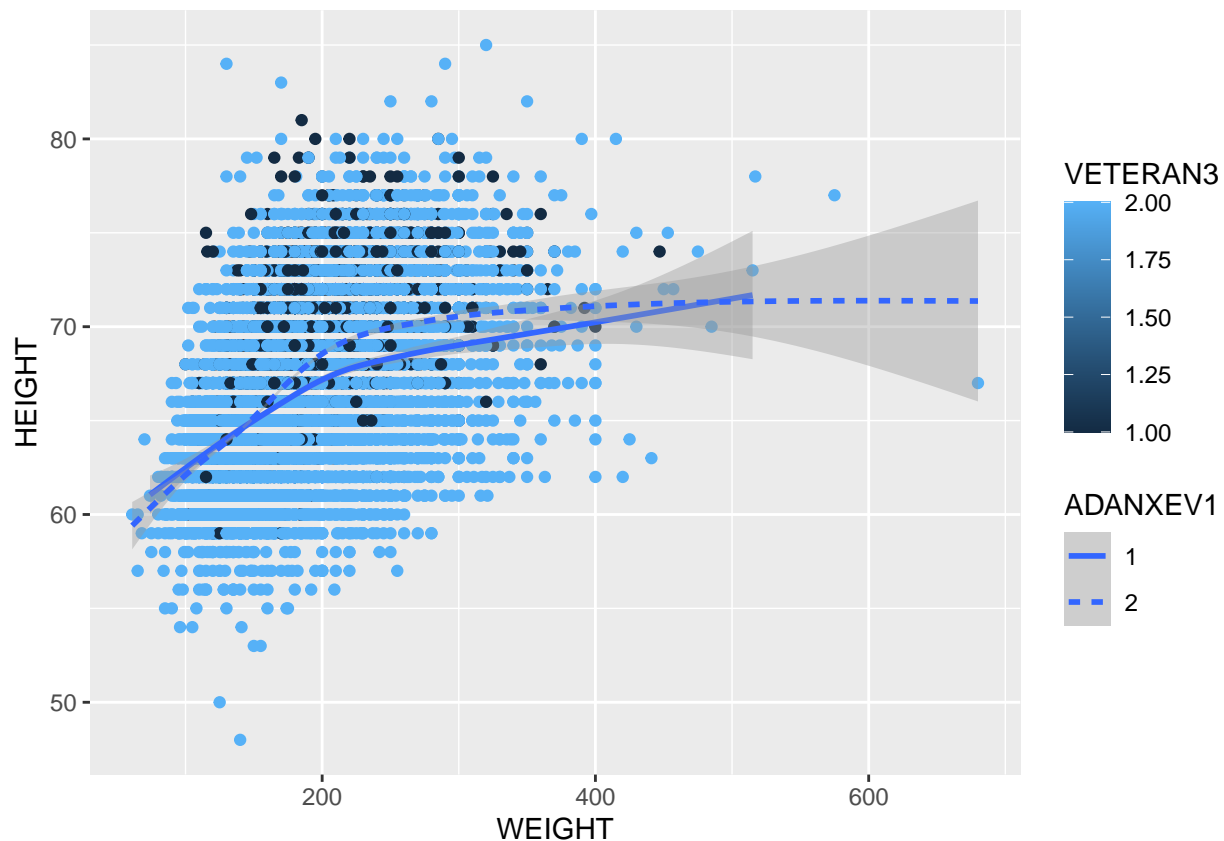


```
#Veteran = 1 means veteran; Veteran = 2 means not a veteran
#Anxiety = 1 or black means diagnosed anxiety
#Anxiety = 2 or blue means no anxiety diagnosis
#This plot shows that veterans (left side) have a slightly lower anxiety proportion than
#the non-veteran group (right side)
#We do not know if this difference is significant though
```

```
data$ADANXE1 <- factor(data$ADANXE1)
ggplot(data = data, mapping = aes(x = WEIGHT, y = HEIGHT, color = VETERAN3)) +
  geom_point() +
  geom_smooth(mapping = aes(linetype = ADANXE1))
```

```
## `geom_smooth()` using method = 'gam' and formula = 'y ~ s(x, bs = "cs")'
```

```
## Warning: The following aesthetics were dropped during statistical transformation: colour
## i This can happen when ggplot fails to infer the correct grouping structure in
## the data.
## i Did you forget to specify a `group` aesthetic or to convert a numerical
## variable into a factor?
```



```
#ADANXE1 = 1 means anxiety; ADANXE1 = 2 means no anxiety diagnosis
#The anxiety trendline is lower which might signal that shorter people get diagnosed with
#anxiety more
#Because veterans look to be taller on average than a non-veteran, this also might signal
#that veterans get diagnosed with anxiety less than a non-veteran
```

```
#Anxiety diagnosis(ADANXE1)
#Calculate descriptive statistics of 'ADANXE1'
summary(data$ADANXE1)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      1.000  2.000   2.000  1.836  2.000   2.000
```

```
#1st quartile is 2 indicating a majority of people are not diagnosed with anxiety
```

```
#Count of people with (1) and without (2) anxiety diagnosis
table(data$ADANXE1)
```

```
##
##      1      2
## 3145 15982
```

```
#With: 3,145 people
#Without: 15,982 people
```

```
#Veteran status(VETERAN3)
#Calculate descriptive statistics of 'VETERAN3'
summary(data$VETERAN3)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
```



```
##    1.000    2.000    2.000    1.865    2.000    2.000
```

*#1st quartile is 2 indicating an overwhelming majority of people are not veterans*

*#Count of veterans (1) and non-veterans (2)*

```
table(data$VETERAN3)
```

```
##
```

```
##      1      2
```

```
## 2585 16542
```

*#Veterans: 2,585 people*

*#Non-veterans: 16,542 people*

*#Height (HEIGHT)*

*#Calculate descriptive statistics of 'HEIGHT'*

```
summary(data$HEIGHT)
```

```
##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.
```

```
##  48.00   64.00   66.00   66.87   70.00   85.00
```

*#Shortest person is 48in (4')*

*#Median person is 66in (5'6")*

*#Tallest person is 85in (7'1")*

```
table(data$HEIGHT)
```

```
##
```

```
##  48  50  53  54  55  56  57  58  59  60  61  62  63  64  65  66
```

```
##    1    1    2    3    7   18   30   40  174  596  639 1459 1474 1767 1587 1776
```

```
##  67  68  69  70  71  72  73  74  75  76  77  78  79  80  81  82
```

```
## 1515 1351 1290 1296 1108 1233  654  500  277  180   74   33   22   12    1    3
```

```
##   83   84   85
```

```
##    1    2    1
```

*#There are relatively few people shorter than 60in (5') or taller than 74in (6'2") which is true of American society so this height dataset feels valid*

*#Weight (WEIGHT)*

*#Calculate descriptive statistics of 'WEIGHT'*

```
summary(data$WEIGHT)
```

```
##      Min. 1st Qu.  Median      Mean 3rd Qu.      Max.
```

```
##   61.0   150.0   179.0   183.3   210.0   680.0
```

*#Lightest person is 61lbs*

*#Median person is 179lbs*

*#Heaviest person is 680lbs*

*#Count of veterans (1) and non-veterans (2)*

```
table(data$WEIGHT)
```

```
##
```

```
##  61  65  68  70  74  75  80  82  84  85  86  87  88  89  90  91
```

```
##    1    2    1    1    1    2    3    3    2    5    4    2    1    1   14    3
```

```
##  92  93  94  95  96  97  98  99 100 101 102 103 104 105 106 107
```

```
##    4    2    4   14    9    5   11    3   50    5   18   11   20   70   23   19
```

```
## 108 109 110 111 112 113 114 115 116 117 118 119 120 121 122 123
```

```
##   31   16  127   10   52   17   27  151   26   34   54   20  264   15   42   35
```

```
## 124 125 126 127 128 129 130 131 132 133 134 135 136 137 138 139
## 46 263 38 39 78 31 463 18 71 32 41 401 51 40 96 43
## 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155
## 570 23 88 69 46 439 46 51 103 47 823 34 82 48 58 382
## 156 157 158 159 160 161 162 163 164 165 166 167 168 169 170 171
## 56 41 91 34 769 24 87 44 48 488 28 55 110 42 808 32
## 172 173 174 175 176 177 178 179 180 181 182 183 184 185 186 187
## 116 47 73 500 52 42 72 31 919 31 74 63 44 505 51 64
## 188 189 190 191 192 193 194 195 196 197 198 199 200 201 202 203
## 56 50 641 29 63 35 38 306 39 47 82 29 1050 26 46 36
## 204 205 206 207 208 209 210 211 212 213 214 215 216 217 218 219
## 36 198 30 29 39 23 416 16 59 25 45 251 32 23 49 19
## 220 221 222 223 224 225 226 227 228 229 230 231 232 233 234 235
## 485 9 27 17 24 252 17 16 23 13 391 5 25 14 11 136
## 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251
## 19 8 25 8 374 10 19 11 12 121 8 14 20 9 367 7
## 252 253 254 255 256 257 258 259 260 261 262 263 264 265 266 267
## 10 9 8 50 11 8 16 6 175 8 8 8 5 67 4 5
## 268 269 270 271 272 273 274 275 276 277 278 279 280 282 283 284
## 13 7 139 3 8 5 6 58 8 3 9 3 126 3 3 2
## 285 286 287 288 289 290 291 292 293 294 295 296 297 298 299 300
## 38 4 7 2 9 80 4 3 1 2 21 2 3 2 2 142
## 301 302 303 304 305 306 307 308 309 310 311 312 313 314 315 316
## 3 2 2 2 8 7 3 2 1 27 1 1 2 2 15 4
## 317 319 320 321 322 323 324 325 326 329 330 332 335 336 337 338
## 3 3 30 1 2 1 1 21 1 2 16 1 6 3 1 1
## 339 340 343 344 345 346 347 348 350 354 356 360 363 365 366 368
## 1 22 1 1 9 1 1 2 27 1 1 13 1 1 1 1
## 370 372 374 375 380 382 385 387 390 392 397 400 415 420 425 430
## 11 1 1 2 2 1 2 1 4 1 1 12 1 2 1 2
## 441 447 450 453 457 475 485 515 517 575 680
## 1 1 1 1 1 1 1 1 1 1 1 1
```

```
#Most people reported weights that are divisible by 5
#There are relatively few people that weigh less than 100lbs or more than 300lbs which
#is generally true of America
```

```
#ADANXEV has to be 0 and 1 for the logistic regression to run
data$ANXIETY <- ifelse(data$ADANXEV == 2, 1, 0)
```

```
#Logistic regression of how height and weight affect anxiety diagnosis frequency
model1 <- glm(ANXIETY ~ HEIGHT + WEIGHT, data = data, family = binomial)
# Display the summary of the logistic regression model
summary(model1)
```

```
##
## Call:
## glm(formula = ANXIETY ~ HEIGHT + WEIGHT, family = binomial, data = data)
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -2.1936732  0.3335161  -6.577 4.79e-11 ***
## HEIGHT      0.0677929  0.0054842  12.361 < 2e-16 ***
## WEIGHT      -0.0037819  0.0004527  -8.354 < 2e-16 ***
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 17097  on 19126  degrees of freedom
## Residual deviance: 16935  on 19124  degrees of freedom
## AIC: 16941
##
## Number of Fisher Scoring iterations: 4
#Logistic regression of how height and weight and veteran status affect anxiety diagnosis frequency
model2 <- glm(ANXIETY ~ HEIGHT + WEIGHT + VETERAN3, data = data, family = binomial)
# Display the summary of the logistic regression model
summary(model2)

##
## Call:
## glm(formula = ANXIETY ~ HEIGHT + WEIGHT + VETERAN3, family = binomial,
##      data = data)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.6030687  0.3983857  -4.024 5.72e-05 ***
## HEIGHT      0.0637003  0.0056765  11.222 < 2e-16 ***
## WEIGHT     -0.0037462  0.0004523  -8.283 < 2e-16 ***
## VETERAN3    -0.1728791  0.0650980  -2.656  0.00792 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 17097  on 19126  degrees of freedom
## Residual deviance: 16928  on 19123  degrees of freedom
## AIC: 16936
##
## Number of Fisher Scoring iterations: 4
#Logistic regression of how height and veteran status affect anxiety diagnosis frequency
model3 <- glm(ANXIETY ~ HEIGHT + VETERAN3, data = data, family = binomial)
# Display the summary of the logistic regression model
summary(model3)

##
## Call:
## glm(formula = ANXIETY ~ HEIGHT + VETERAN3, family = binomial,
##      data = data)
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.85614    0.38804  -2.206  0.02736 *
## HEIGHT      0.04248    0.00505   8.411 < 2e-16 ***
## VETERAN3    -0.18505    0.06501  -2.846  0.00442 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 17097 on 19126 degrees of freedom
## Residual deviance: 16995 on 19124 degrees of freedom
## AIC: 17001
##
## Number of Fisher Scoring iterations: 4
#Logistic regression of how weight and veteran status affect anxiety diagnosis frequency
model14 <- glm(ANXIETY ~ WEIGHT + VETERAN3, data = data, family = binomial)
# Display the summary of the logistic regression model
summary(model14)

##
## Call:
## glm(formula = ANXIETY ~ WEIGHT + VETERAN3, family = binomial,
## data = data)
##
## Coefficients:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) 2.5364809 0.1482581 17.109 < 2e-16 ***
## WEIGHT -0.0013259 0.0004074 -3.255 0.00114 **
## VETERAN3 -0.3550936 0.0627469 -5.659 1.52e-08 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 17097 on 19126 degrees of freedom
## Residual deviance: 17056 on 19124 degrees of freedom
## AIC: 17062
##
## Number of Fisher Scoring iterations: 4
model1$aic

## [1] 16941.43
#16941.43

model2$aic

## [1] 16936.19
#16936.19

model3$aic

## [1] 17000.64
#17000.64

model4$aic

## [1] 17062.32
#17062.32
```

```

#Model 2 has the lowest AIC value so it is likely to be the best fit model

#The HEIGHT coefficient signals that for every 1" increase in HEIGHT, the log-odds of
#having an anxiety diagnosis increases by 0.0637003.
#HEIGHT is statistically significant with a P-value below 0.05 and the relationship is
#unlikely to be from random chance alone
#This does not line up with my prediction from the previous visualizations where I assumed
#that shorter people would be diagnosed with anxiety more
#This could be because there are more shorter people, so it seems like there are more
#anxiety diagnoses initially at the bottom, but the proportion of taller people having
#anxiety could be higher. But that is somewhat hard to discern off a visualization with
#thousands of data points

#The WEIGHT coefficient signals that for every 1lb increase in WEIGHT, the log-odds of
#having an anxiety diagnosis decreases by 0.0037462.
#WEIGHT is statistically significant with a P-value below 0.05 and the relationship is
#unlikely to be from random chance alone
#This does line up with my prediction that heavier people are more likely to have an
#anxiety diagnosis.

#The VETERAN3 coefficient signals that for every 1 unit increase in VETERAN3 (from yes
#to no), the log-odds of having an anxiety diagnosis decreases by 0.1728791.
#VETERAN3 is statistically significant with a P-value below 0.05 and the relationship is
#unlikely to be from random chance alone
#This does not line up with one of the visualizations showing that anxiety diagnosis rate
#increases from a veteran to a non-veteran

model2.chi <- model2$null.deviance - model2$deviance

model2.df <- model2$df.null - model2$df.residual
cat("p-value = ", 1-pchisq(model2.chi, model2.df))

## p-value = 0
#p-value = 0
#A p-value < 0.05 means the observed Chi-square difference is unlikely to have occurred by
#random chance alone

cat("Chi-square difference = ", model2.chi)

## Chi-square difference = 169.0151
#Chi-square difference = 169.0151
#169.0151 is the difference in deviances between the null model and Model 2, which
#includes all of the predictor variables (HEIGHT, WEIGHT, and VETERAN3)
#This is a relatively high Chi-square difference meaning that adding more predictor
#variables will improve my model fit in a logistic regression model

```