

# Assignment2- Markdown

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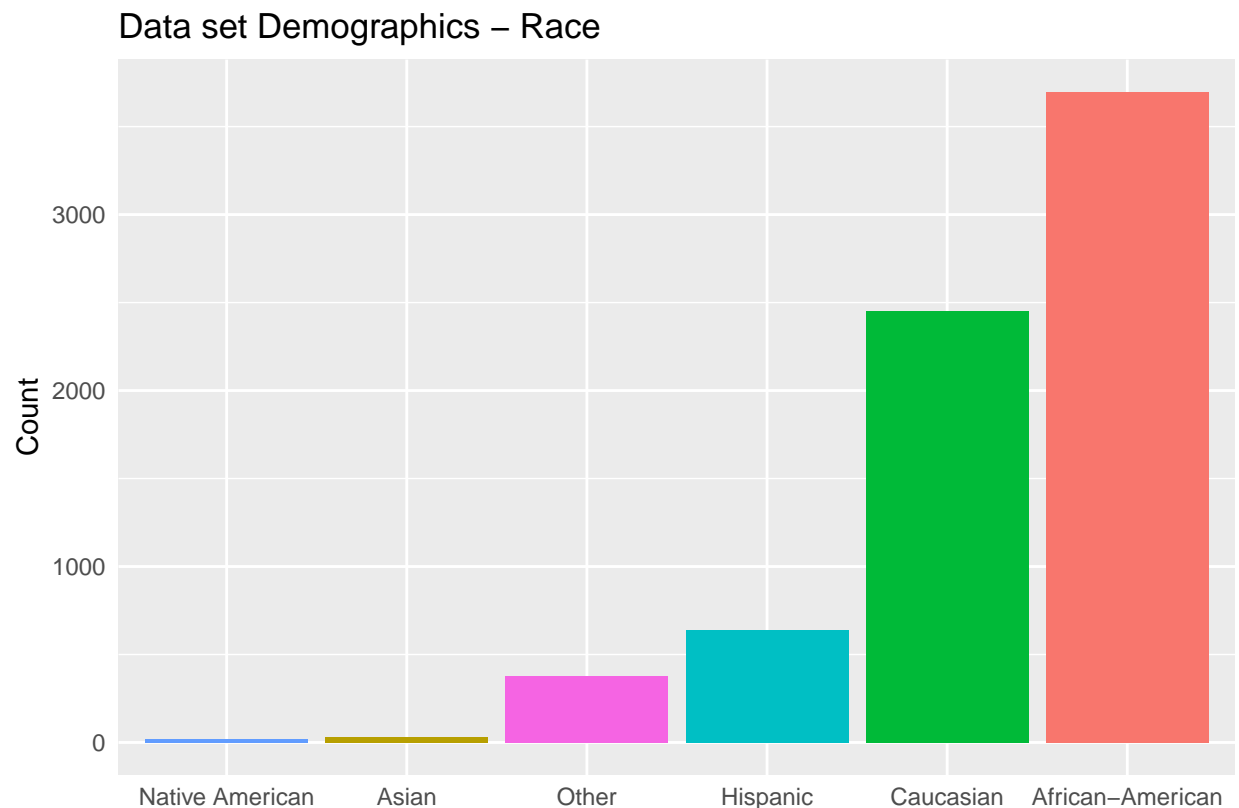
*February 18, 2019*

```
library(tidyverse)
library(data.table)
library(DT)
```

```
data <- fread(input = "D:/Columbia/Spring 2019/Data Science and Public Policy/Data Assignment 2/compas-")
```

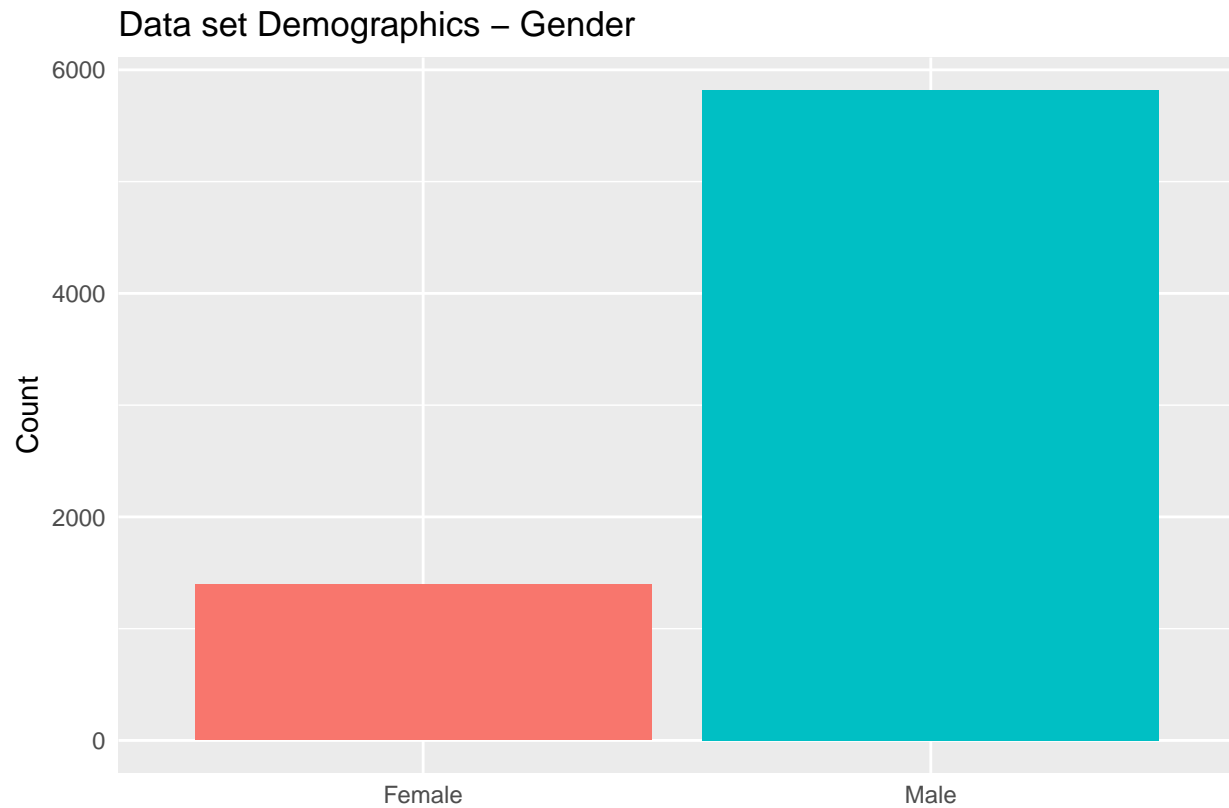
## Including Plots

```
ggplot(data[, .(count = .N), race], aes(x = reorder(race, count), y=count, fill= race)) +
  geom_bar(stat = "identity") +
  theme(axis.ticks = element_blank(), plot.title = element_text(size = rel(1.2)),
        legend.position="none") +
  labs(x = "", y = "Count",
        title = "Data set Demographics - Race")
```



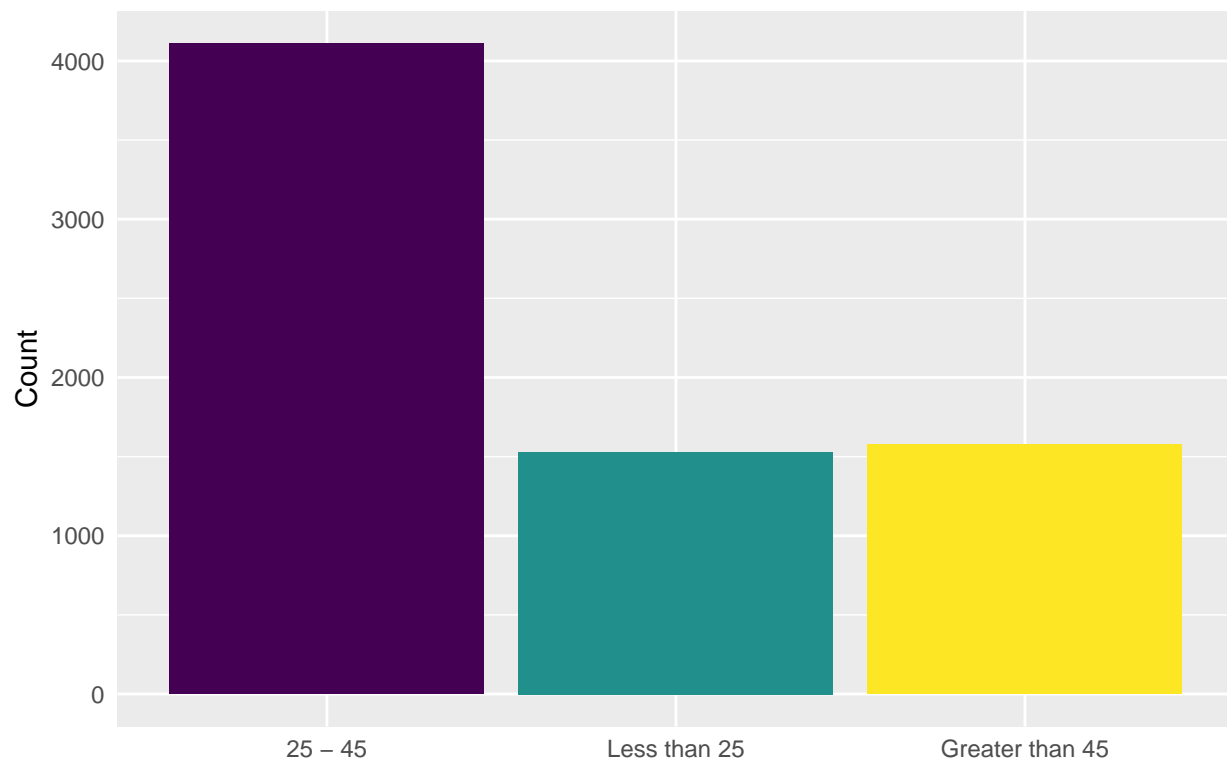
```
ggplot(data[, .(count = .N), sex], aes(x = reorder(sex, count), y=count, fill= sex)) +
  geom_bar(stat = "identity") +
  theme(axis.ticks = element_blank(), plot.title = element_text(size = rel(1.2)),
        legend.position="none") +
```

```
labs( x = "",y = "Count",
      title = "Data set Demographics - Gender")
```



```
data$age_cat <- ordered(data$age_cat, levels = c( "25 - 45", "Less than 25", "Greater than 45"))
ggplot(data[, .(count = .N), age_cat], aes(x = age_cat, y=count, fill= age_cat)) +
  geom_bar(stat = "identity") +
  theme(axis.ticks = element_blank(),plot.title = element_text(size = rel(1.2)),
        legend.position="none") +
  labs( x = "",y = "Count",
        title = "Data set Demographics - Age")
```

Data set Demographics – Age



```
data[score_text == "LOW"] [.N]
```

```
## Empty data.table (0 rows) of 53 cols: id,name,first,last,compas_screening_date,sex...
```

```
type = c('Correlation','Correlation', 'Coefficient', 'Coefficient')
```

```
gender = c('Male','Female', 'Male','Female')
```

```
value = c(-0.058, 0.058, 0.033, -0.033)
```

```
gender_score <- data.frame( "Analysis Type" = type, "Gender" = gender, "Value" = value)
```

```
ggplot(gender_score, aes(x = Analysis.Type, y = Value, fill = Gender)) +
```

```
  geom_bar(stat = "identity", position = "dodge") +
```

```
  theme(axis.ticks = element_blank(),plot.title = element_text(size = rel(1.2))) +
```

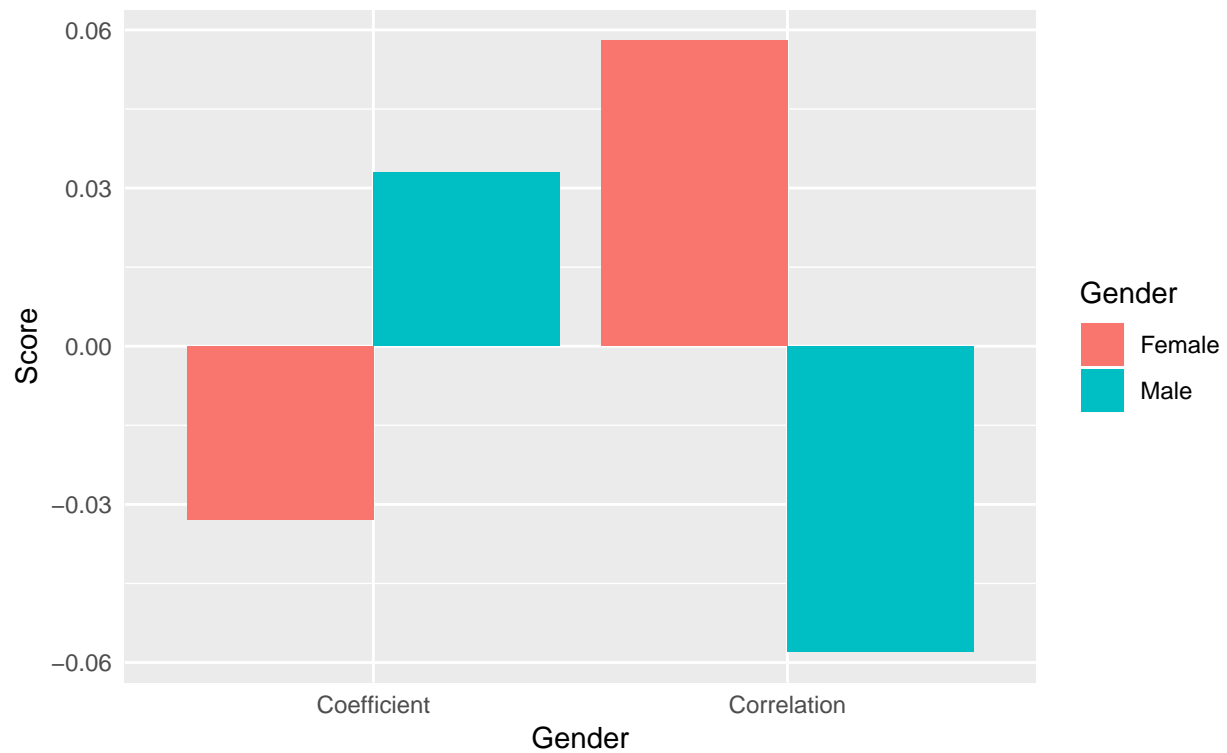
```
  labs(x = "Gender", y = "Score",
```

```
  title = "Correlation vs Regression Coefficient ",
```

```
  subtitle = "Correlation is not Causation")
```

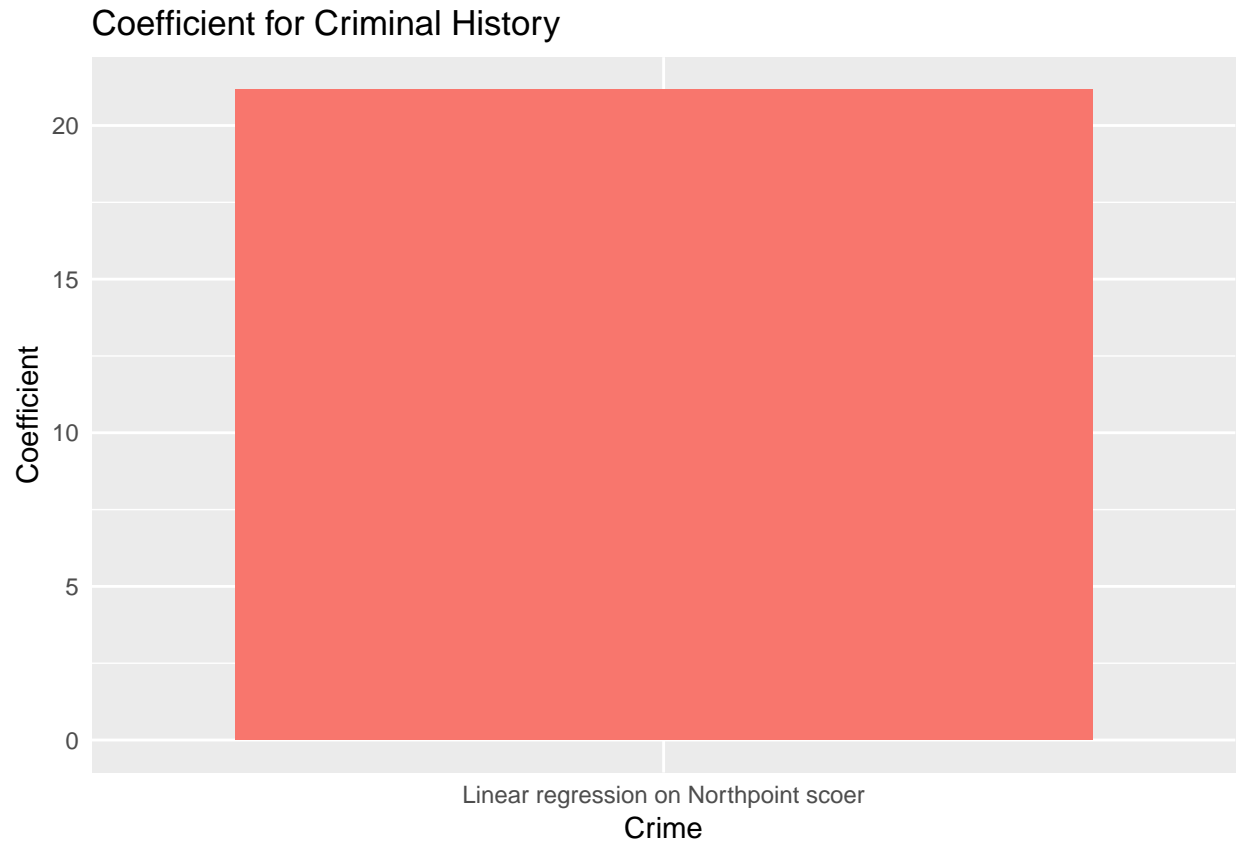
## Correlation vs Regression Coefficient

Correlation is not Causation



```
Race = c('Linear regression on Northpoint scoer')
Score = c(5.86129229, 2.22553838, 3.4098577, 9.66977697)
Race_coeff <- data.frame( "Race" = Race, "Score" = Score)

ggplot(Race_coeff, aes(x = reorder(Race, desc(abs(Score))), y=Score, fill= Race)) +
  geom_bar(stat = "identity") +
  theme(axis.ticks = element_blank(), plot.title = element_text(size = rel(1.2)),
        legend.position="none") +
  labs( x = "Crime", y = "Coefficient",
        title = "Coefficient for Criminal History")
```



This is for Medium High - 1 and Low = 0

Overall

```
print("Overall")

## [1] "Overall"

TP = data[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
FN = data[is_recid == 1 & score_text == "Low", .N ]
FP = data[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
TN = data[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##           Type Positive Negative
## 1 Positive      2140      1177
## 2 Negative      1331      2566

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.65"

print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.35"
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.34"
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))

## [1] "False Positive Rate = 0.31"
```

## African American

```
print("Race = African American")

## [1] "Race = African American"
aad = data[race == "African-American"]

TP = aad[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
FN = aad[is_recid == 1 & score_text == "Low", .N ]
FP = aad[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
TN = aad[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##           Type Positive Negative
## 1 Positive      1445       729
## 2 Negative       591       931

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.64"
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))

## [1] "False Discovery Rate = 0.34"
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.39"
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))

## [1] "False Positive Rate = 0.44"
```

## Caucasian

```
print("Race = Caucasian")

## [1] "Race = Caucasian"
cad = data[race == "Caucasian"]
TP = cad[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
FN = cad[is_recid == 1 & score_text == "Low", .N ]
FP = cad[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
TN = cad[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
```

```
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
## 1 Positive      523      331
## 2 Negative      502     1098
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.66"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.39"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.31"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.23"
```

## AGE CATEGORY

### Less than 25

```
print("age_cat = Less than 25")
```

```
## [1] "age_cat = Less than 25"
```

```
low_age = data[age_cat == "Less than 25"]
TP = low_age[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
FN = low_age[is_recid == 1 & score_text == "Low", .N ]
FP = low_age[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
TN = low_age[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
## 1 Positive      669      330
## 2 Negative      244      286
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.62"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.33"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.46"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.54"
```

## 25 - 45

```
print("age_cat = 25 - 45")

## [1] "age_cat = 25 - 45"
med_age = data[age_cat == "25 - 45"]
TP = med_age[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
FN = med_age[is_recid == 1 & score_text == "Low", .N]
FP = med_age[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
TN = med_age[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
P_value = c(TP, FN)
N_value = c(FP, TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##           Type Positive Negative
## 1 Positive      1244       680
## 2 Negative       776      1409

print(sprintf("Accuracy = %.2f", round((TP+TN)/(TP+TN+FP+FN), 2) ))

## [1] "Accuracy = 0.65"

print(sprintf("False Discovery Rate = %.2f", round((FP)/(TP+FP), 2) ))

## [1] "False Discovery Rate = 0.35"

print(sprintf("False Omission Rate = %.2f", round((FN)/(TN+FN), 2) ))

## [1] "False Omission Rate = 0.36"

print(sprintf("False Positive Rate = %.2f", round((FP)/(TN+FP), 2) ))

## [1] "False Positive Rate = 0.33"
```

## Greater than 45

```
print("age_cat = Greater than 45")

## [1] "age_cat = Greater than 45"
high_age = data[age_cat == "Greater than 45"]
TP = high_age[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
FN = high_age[is_recid == 1 & score_text == "Low", .N]
FP = high_age[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
TN = high_age[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
P_value = c(TP, FN)
N_value = c(FP, TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##           Type Positive Negative
## 1 Positive        227       167
## 2 Negative        311       871

print(sprintf("Accuracy = %.2f", round((TP+TN)/(TP+TN+FP+FN), 2) ))

## [1] "Accuracy = 0.70"
```



```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.42"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.26"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.16"
```

## Gender

### MALE

```
print("sex = Male")
```

```
## [1] "sex = Male"
```

```
md = data[sex == "Male"]
```

```
TP = md[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
```

```
FN = md[is_recid == 1 & score_text == "Low", .N]
```

```
FP = md[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
```

```
TN = md[is_recid == 0 & score_text == "Low", .N]
```

```
names = c('Positive', 'Negative')
```

```
P_value = c(TP,FN)
```

```
N_value = c(FP,TN)
```

```
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
```

```
## 1 Positive      1821       905
```

```
## 2 Negative      1120      1973
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.65"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.33"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.36"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.31"
```

### Female

```
print("sex = Female")
```

```
## [1] "sex = Female"
```

```
fd = data[sex == "Female"]
```

```
TP = fd[is_recid == 1 & (score_text == "Medium" | score_text == "High"), .N]
```

```
FN = fd[is_recid == 1 & score_text == "Low", .N]
```

```
FP = fd[is_recid == 0 & (score_text == "Medium" | score_text == "High"), .N]
```

```

TN = fd[is_recid == 0 & score_text == "Low", .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##          Type Positive Negative
## 1 Positive      319      272
## 2 Negative      211      593

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.65"

print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))

## [1] "False Discovery Rate = 0.46"

print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.26"

print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))

## [1] "False Positive Rate = 0.31"

```

This is for High - 1 and Low, Medium = 0

Overall

```

print("Overall")

## [1] "Overall"

TP = data[is_recid == 1 & score_text == "High", .N]
FN = data[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
FP = data[is_recid == 0 & score_text == "High", .N]
TN = data[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##          Type Positive Negative
## 1 Positive     1041      362
## 2 Negative     2430     3381

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.61"

print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))

## [1] "False Discovery Rate = 0.26"

print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.42"

```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.10"
```

## African-American

```
print("Race = African American")
```

```
## [1] "Race = African American"
```

```
aad = data[race == "African-American"]
```

```
TP = aad[is_recid == 1 & score_text == "High", .N]
```

```
FN = aad[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
```

```
FP = aad[is_recid == 0 & score_text == "High", .N]
```

```
TN = aad[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
```

```
names = c('Positive', 'Negative')
```

```
P_value = c(TP,FN)
```

```
N_value = c(FP,TN)
```

```
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
```

```
## 1 Positive      771      254
```

```
## 2 Negative     1265     1406
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.59"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.25"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.47"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.15"
```

## Caucasian

```
print("Race = Caucasian")
```

```
## [1] "Race = Caucasian"
```

```
cad = data[race == "Caucasian"]
```

```
TP = cad[is_recid == 1 & score_text == "High", .N]
```

```
FN = cad[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
```

```
FP = cad[is_recid == 0 & score_text == "High", .N]
```

```
TN = cad[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
```

```
names = c('Positive', 'Negative')
```

```
P_value = c(TP,FN)
```

```
N_value = c(FP,TN)
```

```
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
```

```
## 1 Positive      201      75
## 2 Negative      824     1354

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.63"

print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))

## [1] "False Discovery Rate = 0.27"

print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.38"

print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))

## [1] "False Positive Rate = 0.05"
```

## AGE

### Less than 25

```
print("age_cat = Less than 25")

## [1] "age_cat = Less than 25"

low_age = data[age_cat == "Less than 25"]
TP = low_age[is_recid == 1 & score_text == "High", .N]
FN = low_age[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
FP = low_age[is_recid == 0 & score_text == "High", .N]
TN = low_age[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##      Type Positive Negative
## 1 Positive      343      110
## 2 Negative      570      506

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.56"

print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))

## [1] "False Discovery Rate = 0.24"

print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.53"

print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))

## [1] "False Positive Rate = 0.18"
```

### 25 - 45

```
print("age_cat = 25 - 45")
```

```
## [1] "age_cat = 25 - 45"
mid_age = data[age_cat == "25 - 45"]
TP = mid_age[is_recid == 1 & score_text == "High", .N]
FN = mid_age[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
FP = mid_age[is_recid == 0 & score_text == "High", .N]
TN = mid_age[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
## 1 Positive      615      207
## 2 Negative     1405     1882
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.61"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.25"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.43"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.10"
```

## Greater than 45

```
print("age_cat = Greater than 45")
```

```
## [1] "age_cat = Greater than 45"
```

```
high_age = data[age_cat == "Greater than 45"]
TP = high_age[is_recid == 1 & score_text == "High", .N]
FN = high_age[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
FP = high_age[is_recid == 0 & score_text == "High", .N]
TN = high_age[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
## 1 Positive       83       45
## 2 Negative      455      993
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.68"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.35"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.31"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.04"
```

## Gender

### MALE

```
print("sex = Male")
```

```
## [1] "sex = Male"
```

```
md = data[sex == "Male"]
TP = md[is_recid == 1 & score_text == "High", .N]
FN = md[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
FP = md[is_recid == 0 & score_text == "High", .N]
TN = md[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)
```

```
##      Type Positive Negative
## 1 Positive      921      292
## 2 Negative     2020     2586
```

```
print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))
```

```
## [1] "Accuracy = 0.60"
```

```
print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))
```

```
## [1] "False Discovery Rate = 0.24"
```

```
print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))
```

```
## [1] "False Omission Rate = 0.44"
```

```
print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))
```

```
## [1] "False Positive Rate = 0.10"
```

### Female

```
print("sex = Female")
```

```
## [1] "sex = Female"
```

```
fd = data[sex == "Female"]
TP = fd[is_recid == 1 & score_text == "High", .N]
FN = fd[is_recid == 1 & (score_text == "Medium" | score_text == "Low"), .N ]
FP = fd[is_recid == 0 & score_text == "High", .N]
TN = fd[is_recid == 0 & (score_text == "Medium" | score_text == "Low"), .N]
names = c('Positive', 'Negative')
P_value = c(TP,FN)
```

```

N_value = c(FP,TN)
data.frame( "Type" = names, "Positive" = P_value, "Negative" = N_value)

##      Type Positive Negative
## 1 Positive     120       70
## 2 Negative     410      795

print(sprintf("Accuracy = %.2f",round((TP+TN)/(TP+TN+FP+FN),2) ))

## [1] "Accuracy = 0.66"

print(sprintf("False Discovery Rate = %.2f",round((FP)/(TP+FP),2) ))

## [1] "False Discovery Rate = 0.37"

print(sprintf("False Omission Rate = %.2f",round((FN)/(TN+FN),2) ))

## [1] "False Omission Rate = 0.34"

print(sprintf("False Positive Rate = %.2f",round((FP)/(TN+FP),2) ))

## [1] "False Positive Rate = 0.08"

```