

# Suicide Rates Data

## Load CSV and use summary function

```
sData <- read.csv("~/Downloads/master.csv")
```

```
summary(sData)
```

```
##          country          year          sex          age
## Austria   : 382   Min.   :1985   female:13910   15-24 years:4642
## Iceland   : 382   1st Qu.:1995   male  :13910   25-34 years:4642
## Mauritius  : 382   Median :2002                   35-54 years:4642
## Netherlands: 382   Mean    :2001                   5-14 years :4610
## Argentina  : 372   3rd Qu.:2008                   55-74 years:4642
## Belgium    : 372   Max.    :2016                   75+ years  :4642
## (Other)    :25548
## suicides_no population suicides.100k.pop
## Min.   : 0.0   Min.   : 278   Min.   : 0.00
## 1st Qu.: 3.0   1st Qu.: 97498 1st Qu.: 0.92
## Median : 25.0  Median : 430150 Median : 5.99
## Mean   : 242.6 Mean   : 1844794 Mean   : 12.82
## 3rd Qu.: 131.0 3rd Qu.: 1486143 3rd Qu.: 16.62
## Max.   :22338.0 Max.   :43805214 Max.   :224.97
##
##          country.year      HDI.for.year      gdp_for_year....
## Albania1987: 12   Min.   :0.483   1,002,219,052,968: 12
## Albania1988: 12   1st Qu.:0.713   1,011,797,457,139: 12
## Albania1989: 12   Median :0.779   1,016,418,229 : 12
## Albania1992: 12   Mean    :0.777   1,018,847,043,277: 12
## Albania1993: 12   3rd Qu.:0.855   1,022,191,296 : 12
## Albania1994: 12   Max.    :0.944   1,023,196,003,075: 12
## (Other)    :27748   NA's   :19456   (Other)          :27748
## gdp_per_capita.... generation
## Min.   : 251   Boomers      :4990
## 1st Qu.: 3447   G.I. Generation:2744
## Median : 9372   Generation X   :6408
## Mean   : 16866   Generation Z   :1470
## 3rd Qu.: 24874   Millenials    :5844
## Max.   :126352   Silent        :6364
##
```

## Gender Data

```
#Filtering data for density because outliers are rare and skew density graph
```

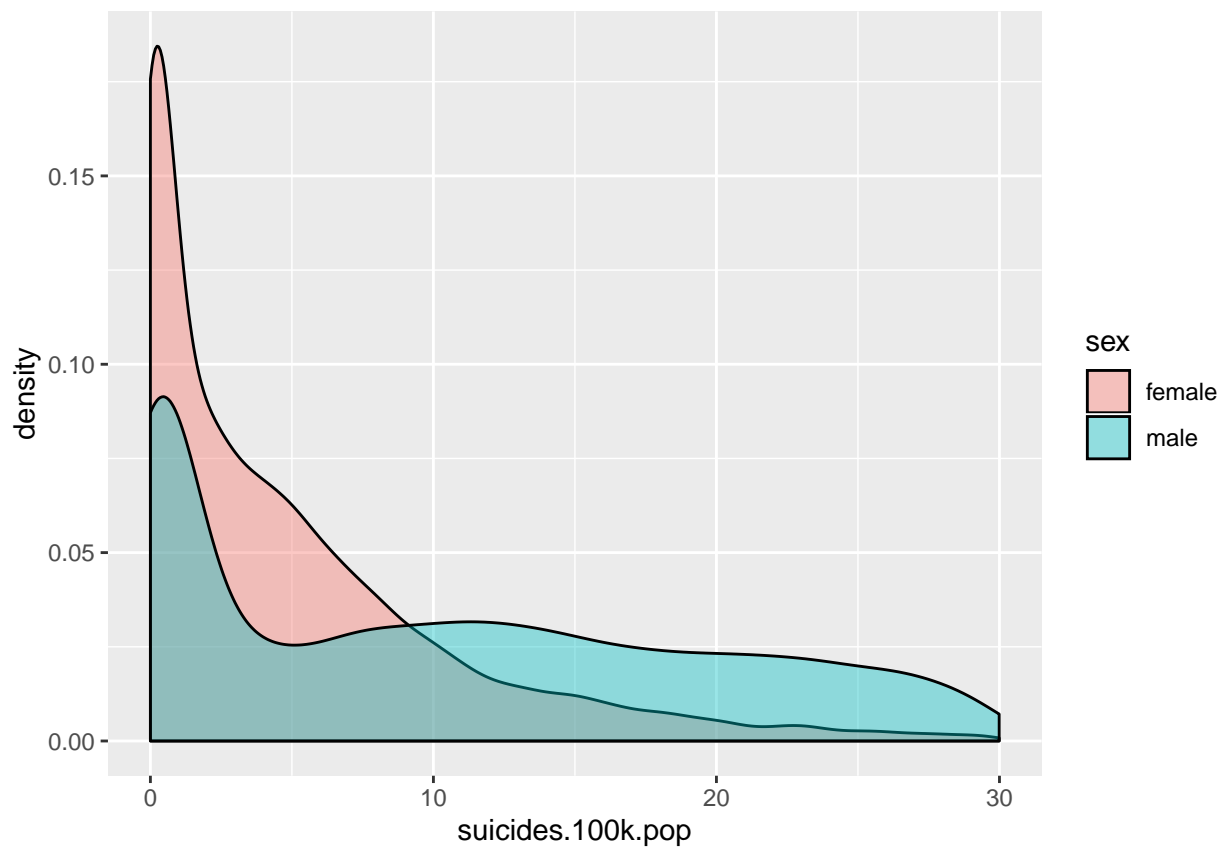
```
densityData <- sData %>% filter(suicides.100k.pop < 30)
```

```
library(ggplot2)
```

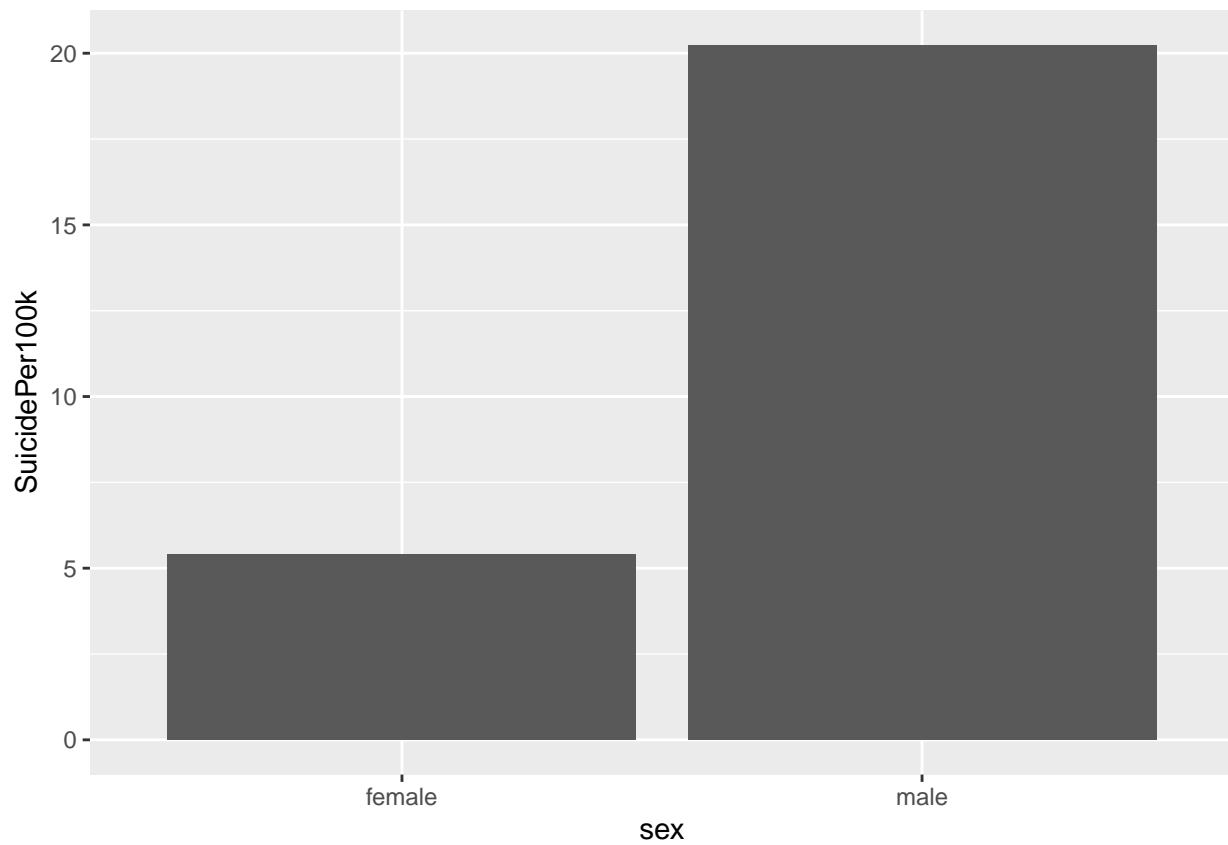
```
# Density based on Gender, Males tend to commit suicide far more often, shifted data to not include ext
```

```
suicideBySex<- sData %>% select(sex, suicides.100k.pop) %>% group_by(sex) %>% summarise(SuicidePer100k=
```

```
ggplot(densityData, aes(x=suicides.100k.pop,fill=sex))+
  geom_density(alpha=0.4)
```



```
ggplot(suicideBySex, aes(x=sex, y=SuicidePer100k)) +  
  geom_bar(stat="identity")
```

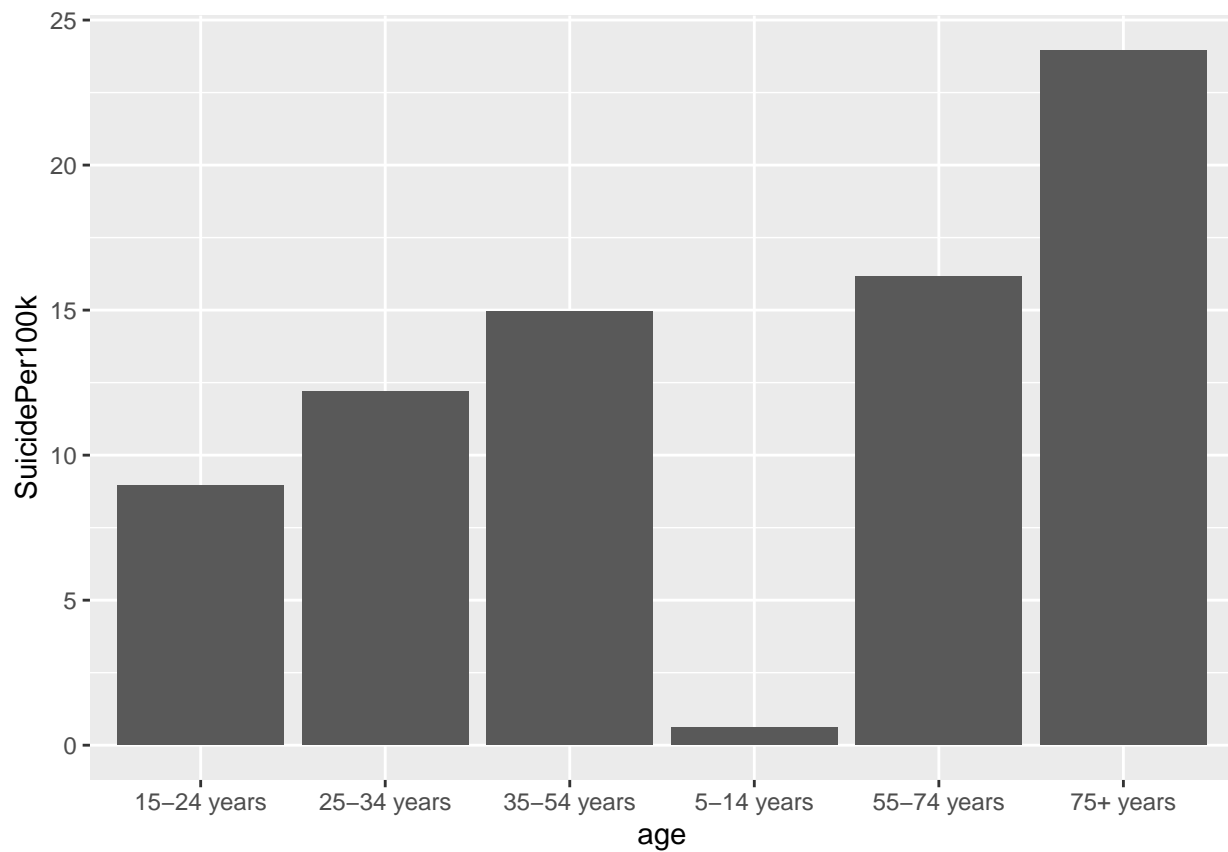


## Age Data

*#Density based on age groups, rates steadily climb as people get older*

```
suicideByAge <- sData %>% select(age, suicides.100k.pop) %>% group_by(age) %>% summarise(SuicidePer100k =
  suicides.100k.pop / population)

generationAverages<-ggplot(data=suicideByAge, aes(x=age, y=SuicidePer100k)) +
  geom_bar(stat="identity")
generationAverages
```

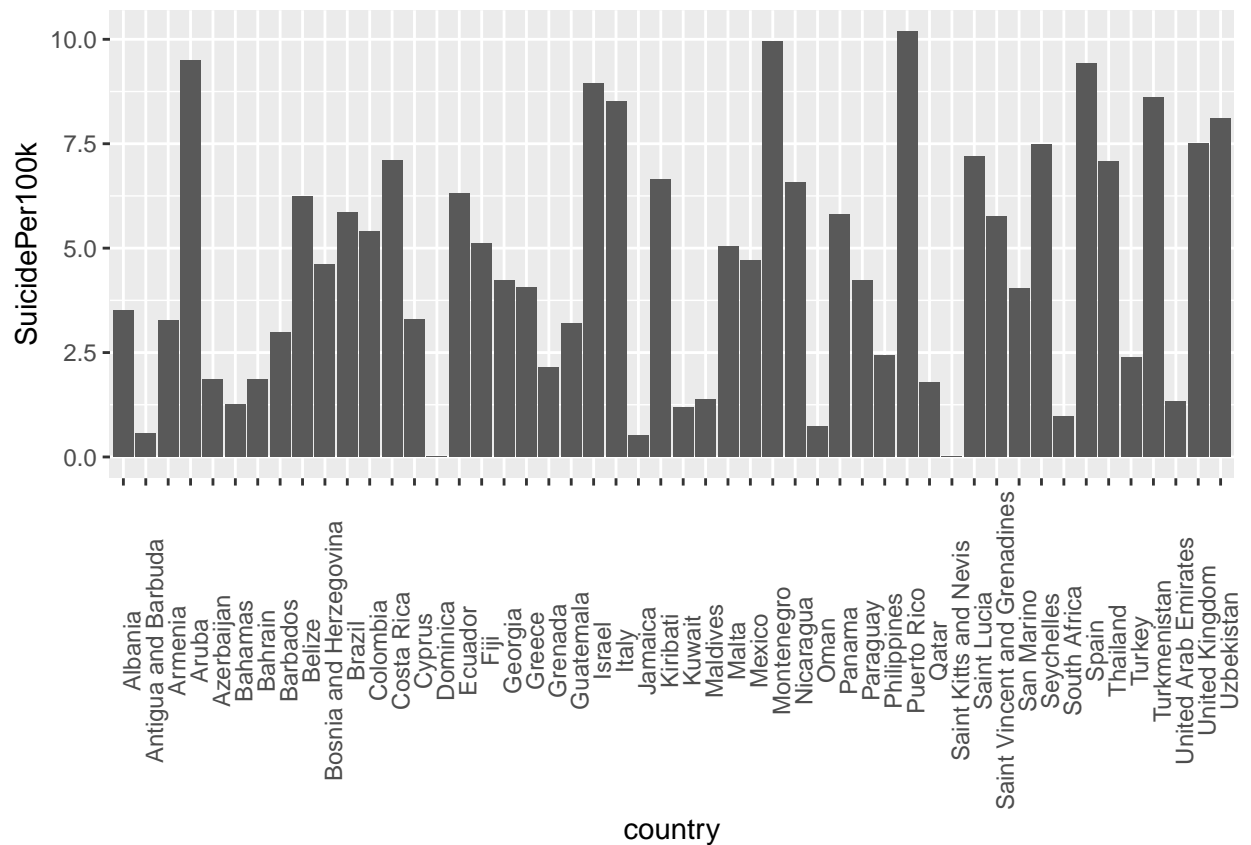


## Country Data

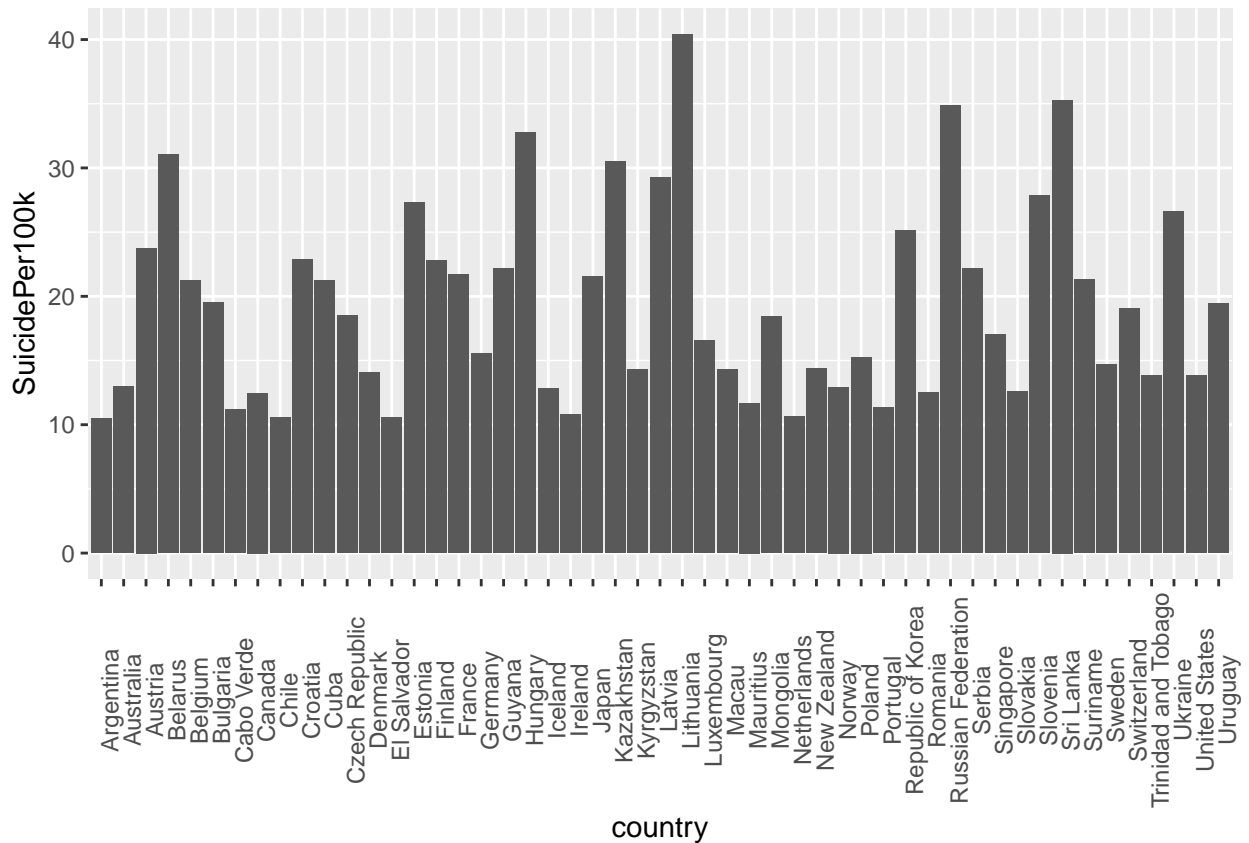
```
#Suicide by country, broken down into two sides, arranged by rate so countryone has lowe rate countries
suicideByCountry <- sData %>% select(country, suicides.100k.pop) %>% group_by(country) %>% summarise(SuicidePer100k = suicides.100k.pop)

suicideByCountry <- arrange(suicideByCountry, SuicidePer100k)
countryOne <- suicideByCountry[1:50,]
countryTwo <- suicideByCountry[51:101,]

countryOneAverage<-ggplot(data=countryOne, aes(x=country, y=SuicidePer100k)) +
  geom_bar(stat="identity") + theme(axis.text.x = element_text(angle = 90))
countryOneAverage
```



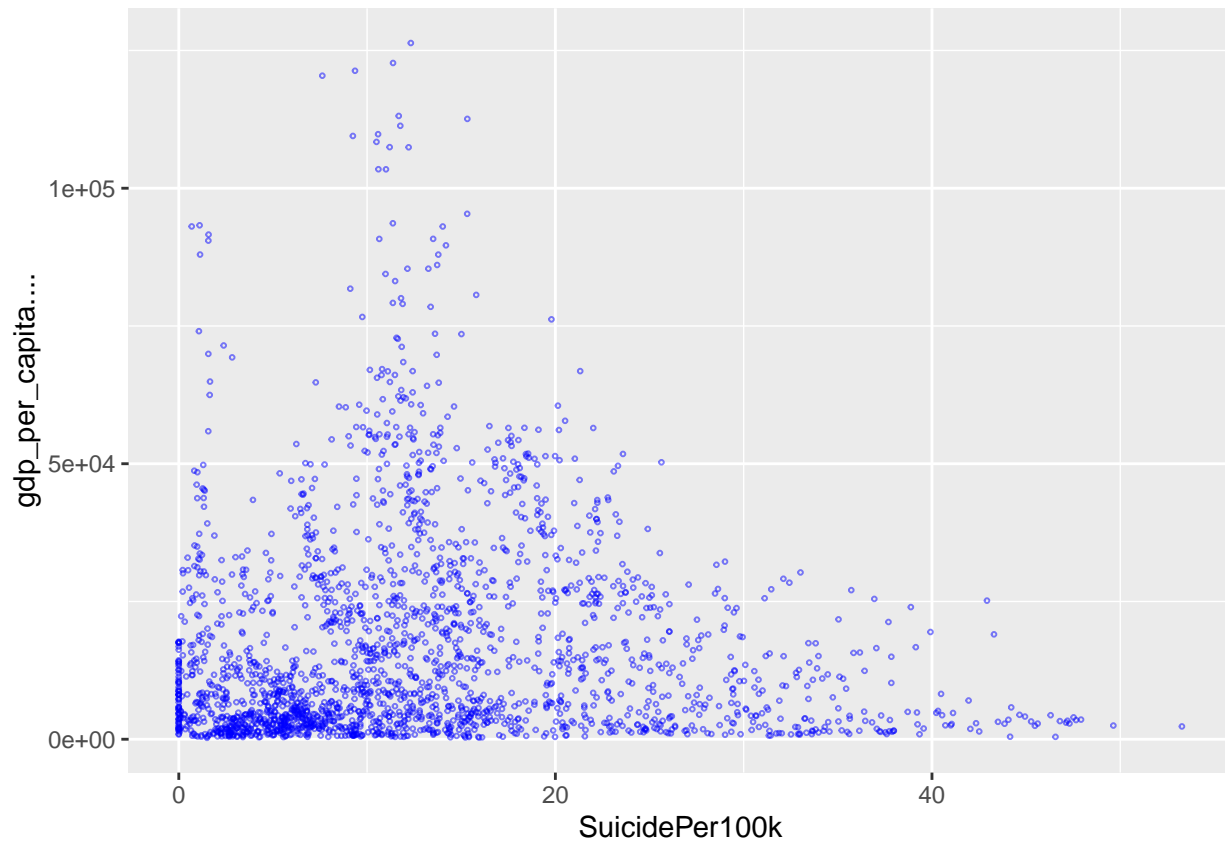
```
countryTwoAverage<-ggplot(data=countryTwo, aes(x=country, y=SuicidePer100k)) +
  geom_bar(stat="identity") + theme(axis.text.x = element_text(angle = 90))
countryTwoAverage
```



## GDP Data

```
#suicide by GDP, found some meaningful correlation rates are similar until 20 + that's when poorer coun
suicideByGDP <- sData %>% select(gdp_per_capita..., suicides.100k.pop) %>% group_by(gdp_per_capita...)

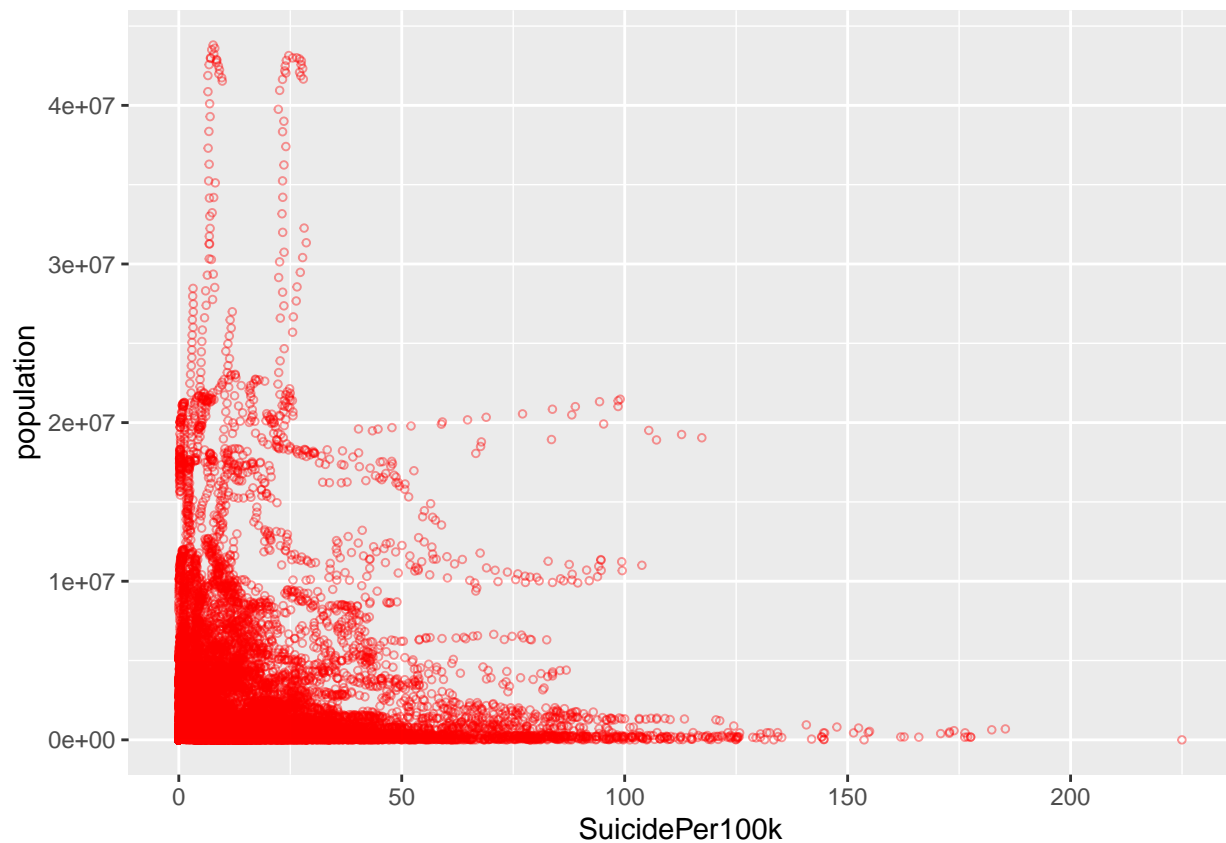
ggplot(suicideByGDP, aes(x=SuicidePer100k, y=gdp_per_capita...)) +
  geom_point(size=.5, shape=1, colour = "blue",alpha=0.5)
```



## Population Data

```
#suicide by population, this was completely random no correlation
suicideByPop <- sData %>% select(population, suicides.100k.pop) %>% group_by(population) %>% summarise(

ggplot(suicideByPop, aes(x=SuicidePer100k, y=population)) +
  geom_point(size=1, shape=1, color = "red",alpha=0.4)
```

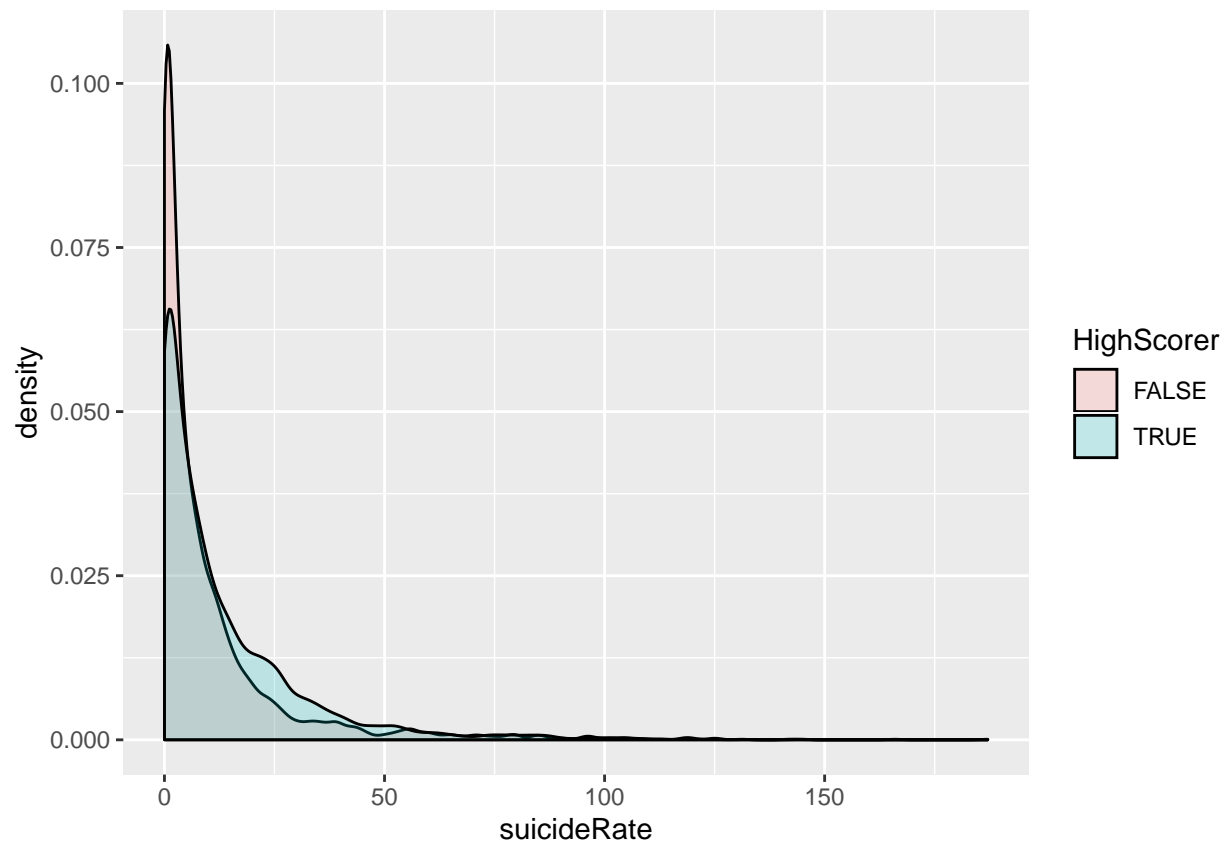


## HDI Data

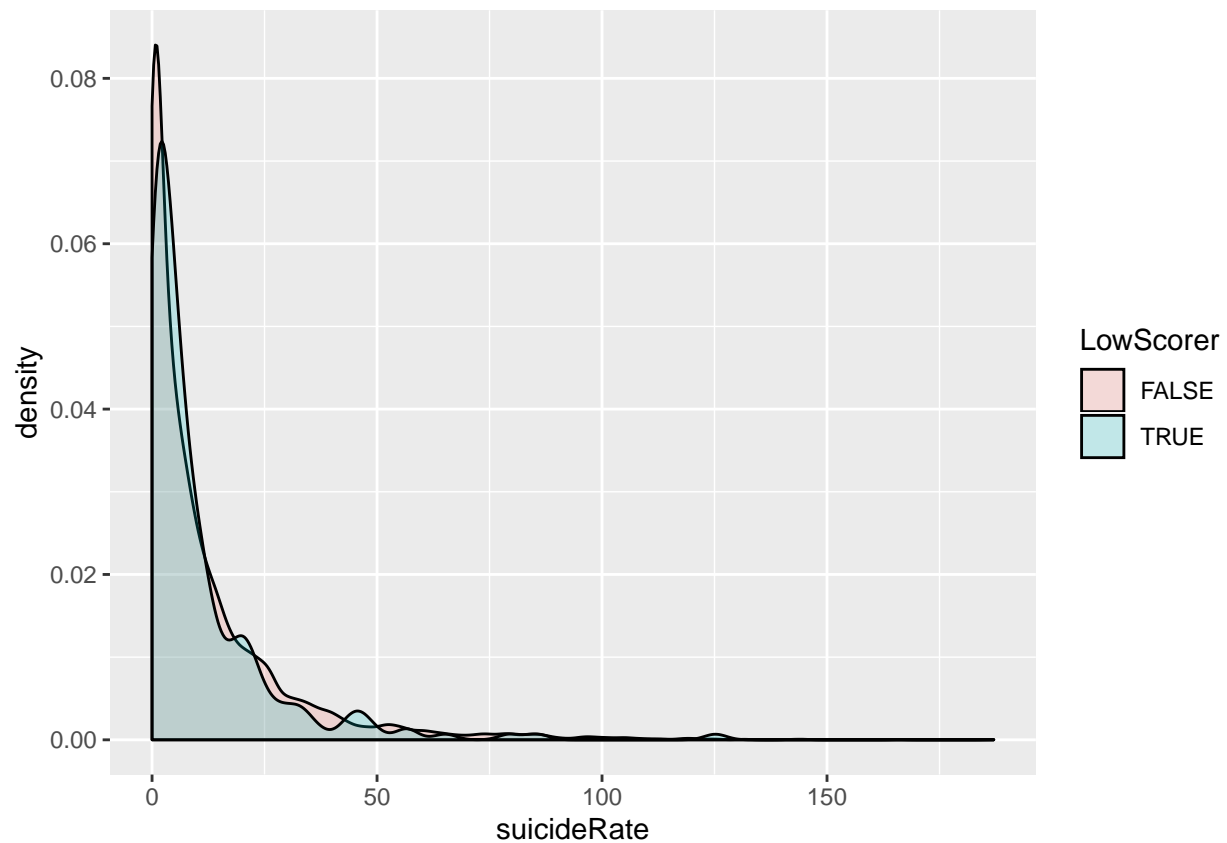
```
# Suicide based on countries with human development index, seperated into countries with 7.5 above/below
suicideByHDI <- sData %>%select(HDI.for.year,suicideRate = suicides.100k.pop) %>%drop_na() %>% group_by(HDI.for.year)

ggplot(suicideByHDI, aes(x=suicideRate,fill=HighScorer))+
  geom_density(alpha=0.2)
```





```
ggplot(suicideByHDI, aes(x=suicideRate,fill=LowScorer))+  
  geom_density(alpha=0.2)
```

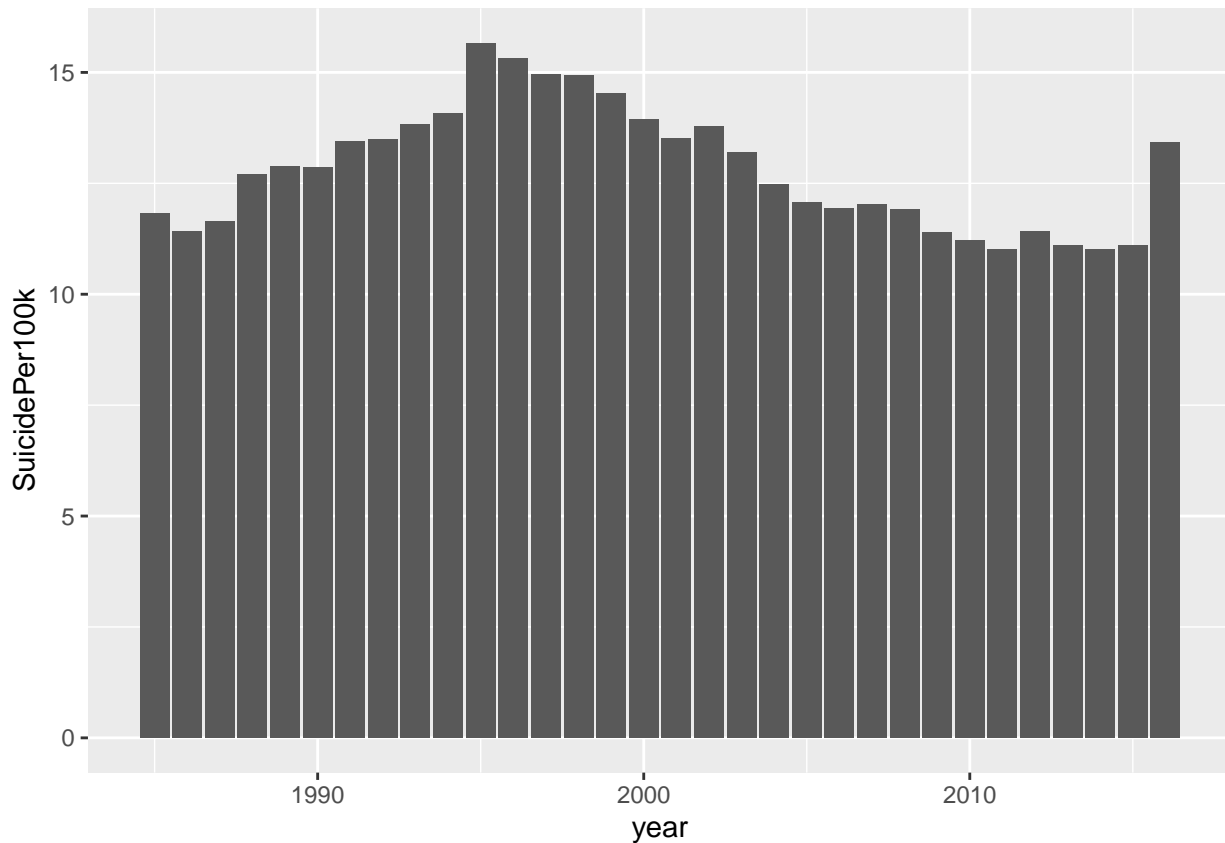


## Year Data

```
#Suicide based on year, went with plain bar graph no real insight gained
suicideByYear <- sData %>% select(year, suicides.100k.pop) %>% group_by(year) %>% summarise(SuicidePer100k = suicides.100k.pop / population)

yearBar<-ggplot(data=suicideByYear, aes(x=year, y=SuicidePer100k)) +
  geom_bar(stat="identity")

yearBar
```



## Clustering

```
summary(sData)
```

```
##          country          year          sex          age
## Austria   : 382   Min.   :1985 female:13910 15-24 years:4642
## Iceland   : 382 1st Qu.:1995 male  :13910 25-34 years:4642
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## Netherlands: 382 Mean   :2001                5-14 years :4610
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## Mean   : 242.6 Mean   : 1844794 Mean   : 12.82
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```

```
## Albania1994: 12 Max. :0.944 1,023,196,003,075: 12
## (Other) :27748 NA's :19456 (Other) :27748
## gdp_per_capita.... generation
## Min. : 251 Boomers :4990
## 1st Qu.: 3447 G.I. Generation:2744
## Median : 9372 Generation X :6408
## Mean : 16866 Generation Z :1470
## 3rd Qu.: 24874 Millenials :5844
## Max. :126352 Silent :6364
##
```

```
sData2 <- sData
library(dplyr)
library(cluster)
library(tidyverse)

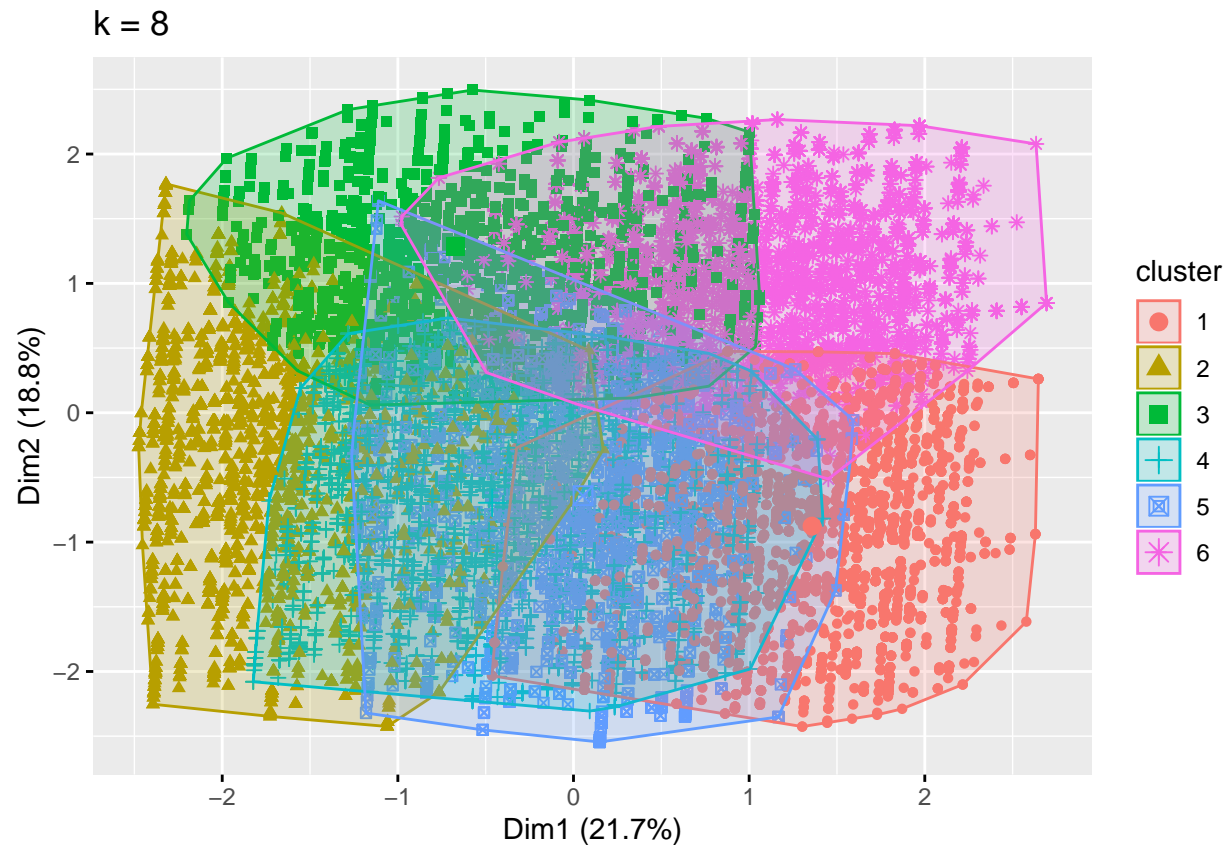
#drop values that are redundant or not useful for the model
drops <- c("country.year", "HDI.for.year", 'suicides_no', "gdp_for_year....", "generation")
sData2 <- sData2[ , !(names(sData2) %in% drops)]
#Drop na data
sData2 <- na.omit(sData2)

#bin variables based on quartiles
sData2$population<-cut(sData2$population, c(278,97498,430150,1486143,438025124))
sData2$suicides.100k.pop <- cut(sData2$suicides.100k.pop, c(0.00,0.92,5.99,16.62,224.97))
sData2$gdp_per_capita....<- cut(sData2$gdp_per_capita....,c(251,3447,9372,24874,126352))
sData2$year<-cut(sData2$year, c(1985,1995,2002,2008,2016))

#mutate all variables into numeric, drop NA values, normalize the dataset
sData2 <- mutate_all(sData2, function(x) as.numeric(x))
sData2 <- na.omit(sData2)
sData2 <- normalize.Dataset(sData2)

clusters <- kmeans(sData2,centers=6,nstart=50)
library(factoextra)
```

```
## Welcome! Related Books: `Practical Guide To Cluster Analysis in R` at https://goo.gl/13EFCZ
p3 <- fviz_cluster(clusters,geom="point", sData2) + ggtitle('k = 8')
p3
```



## Naive Bayes Full Data Set

```
#model without HDI
library(e1071)

#Cut sData2 into a train/test datasets, mutate train/test back into factor,
train <- sData2[1:18467,]
test <- sData2[18468:nrow(sData2),]
train <- mutate_all(train, function(x) as.factor(x))
test <- mutate_all(test, function(x) as.factor(x))

#create model using all values stored in sData2
bayesModel <- naiveBayes(as.factor(suicides.100k.pop)~country + year+sex+age + population + gdp_per_cap,
                        data = train)

#predict using bayesmodel
pred.raw <- predict(bayesModel, test, type = "class")

#create confusion matrix based on how well it predicts suicide.100k.pop then calculate accuracy
confusion <- table(predict(bayesModel, test),
                    test$suicides.100k.pop,
                    dnn=c("prediction","truth"))

confusion
```

```
##                truth
## prediction      0 0.333333333333333 0.666666666666667 1
##    0            455                171                0    0
##    0.333333333333333 93                761            470 108
##    0.666666666666667 27                299            493 183
##    1              10                116            367 1064
```

```
sum(diag(confusion)/nrow(test))
```

```
## [1] 0.6006065
```

```
#60% accuracy not bad considering there is 4 potential options
```

## Bayes With HDI

```
#Model with HDI but a lot less rows
#reference sData(original dataset) to use with the new set used for Naive Bayes
bayesWithHDI <- sData
#Drop values that won't be used in the model
drops <- c("country.year", "suicides_no", "gdp_for_year...", "generation")
bayesWithHDI <- bayesWithHDI[, !(names(bayesWithHDI) %in% drops)]
#drop any NAs
bayesWithHDI <- na.omit(bayesWithHDI)

#Use this to bin continuous values into categorical values, uses their quartiles as binning cuts/breaks
bayesWithHDI$population <- cut(bayesWithHDI$population, c(278, 97498, 430150, 1486143, 438025124))
bayesWithHDI$suicides.100k.pop <- cut(bayesWithHDI$suicides.100k.pop, c(0.00, 0.92, 5.99, 16.62, 224.97))
bayesWithHDI$gdp_per_capita... <- cut(bayesWithHDI$gdp_per_capita..., c(251, 3447, 9372, 24874, 126352))
bayesWithHDI$year <- cut(bayesWithHDI$year, c(1985, 1995, 2002, 2008, 2016))
bayesWithHDI$HDI.for.year <- cut(bayesWithHDI$HDI.for.year, c(.4830, .7130, .7790, .8550, .9440))

#mutate dataset into numeric, omit any nas again, normalize dataset
bayesWithHDI <- mutate_all(bayesWithHDI, function(x) as.numeric(x))
bayesWithHDI <- na.omit(bayesWithHDI)
bayesWithHDI <- normalize.Dataset(bayesWithHDI)

#make train/test datasets, revert back to factor
trainHDI <- bayesWithHDI[1:6000,]
testHDI <- bayesWithHDI[6001:nrow(bayesWithHDI),]
trainHDI <- mutate_all(trainHDI, function(x) as.factor(x))
testHDI <- mutate_all(testHDI, function(x) as.factor(x))

#naive bayes model this time uses HDI
modelHDI <- naiveBayes(suicides.100k.pop~country+year+sex+age + population + gdp_per_capita... + HDI.f
                        data = trainHDI)

#predict class labels for test dataset based on suicides.100k
pred.raw <- predict(modelHDI, testHDI, type = "class")
confusion <- table(predict(modelHDI, testHDI),
                    testHDI$suicides.100k.pop,
                    dnn=c("prediction", "truth"))

confusion
```

```
##
## prediction      0 0.333333333333333 0.666666666666667 1
## 0              84              39              0 0
## 0.333333333333333 7              155             96 8
## 0.666666666666667 1              102             75 45
## 1              0              18              64 120
```

```
#Find the accuracy of the model
sum(diag(confusion)/nrow(testHDI))
```

```
## [1] 0.5331695
```

```
#only 53% lower than without, probably due to dropping a ton of data to use HDI
```