

Case Study 1

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R Markdown

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.5    v purrr  0.3.4
## v tibble  3.1.5    v dplyr  1.0.7
## v tidyr   1.1.3    v stringr 1.4.0
## v readr   2.0.1    v forcats 0.5.1
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
```

```
library(ggpubr)
library(class)
library(caret)
```

```
## Loading required package: lattice
```

```
##
```

```
## Attaching package: 'caret'
```

```
## The following object is masked from 'package:purrr':
```

```
##
```

```
## lift
```

```
library(e1071)
library(rvest)
```

```
##
```

```
## Attaching package: 'rvest'
```

```
## The following object is masked from 'package:readr':
```

```
##
```

```
## guess_encoding
```

```
beers <- read_csv("https://raw.githubusercontent.com/BivinSadler/MSDS_6306_Doing-Data-Science/Master/Un
```

```
## Rows: 2410 Columns: 7
```

```
## -- Column specification -----
```

```
## Delimiter: ","
```

```
## chr (2): Name, Style
```

```
## dbl (5): Beer_ID, ABV, IBU, Brewery_id, Ounces
```

```
##
```

```
## i Use 'spec()' to retrieve the full column specification for this data.
```

```
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
breweries <- read_csv("https://raw.githubusercontent.com/BivinSadler/MSDS_6306_Doing-Data-Science/Master
```

```
## Rows: 558 Columns: 4
```

```
## -- Column specification -----
```

```
## Delimiter: ","
```

```
## chr (3): Name, City, State
```

```
## dbl (1): Brew_ID
```

```
##
```

```
## i Use 'spec()' to retrieve the full column specification for this data.
```

```
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
##How many breweries are present in each state?
```

```
#This table outlines the quantity of breweries in each state and will be used as the basis of the follow
```

```
numbystate <- data.frame(breweries %>% count(State))
```

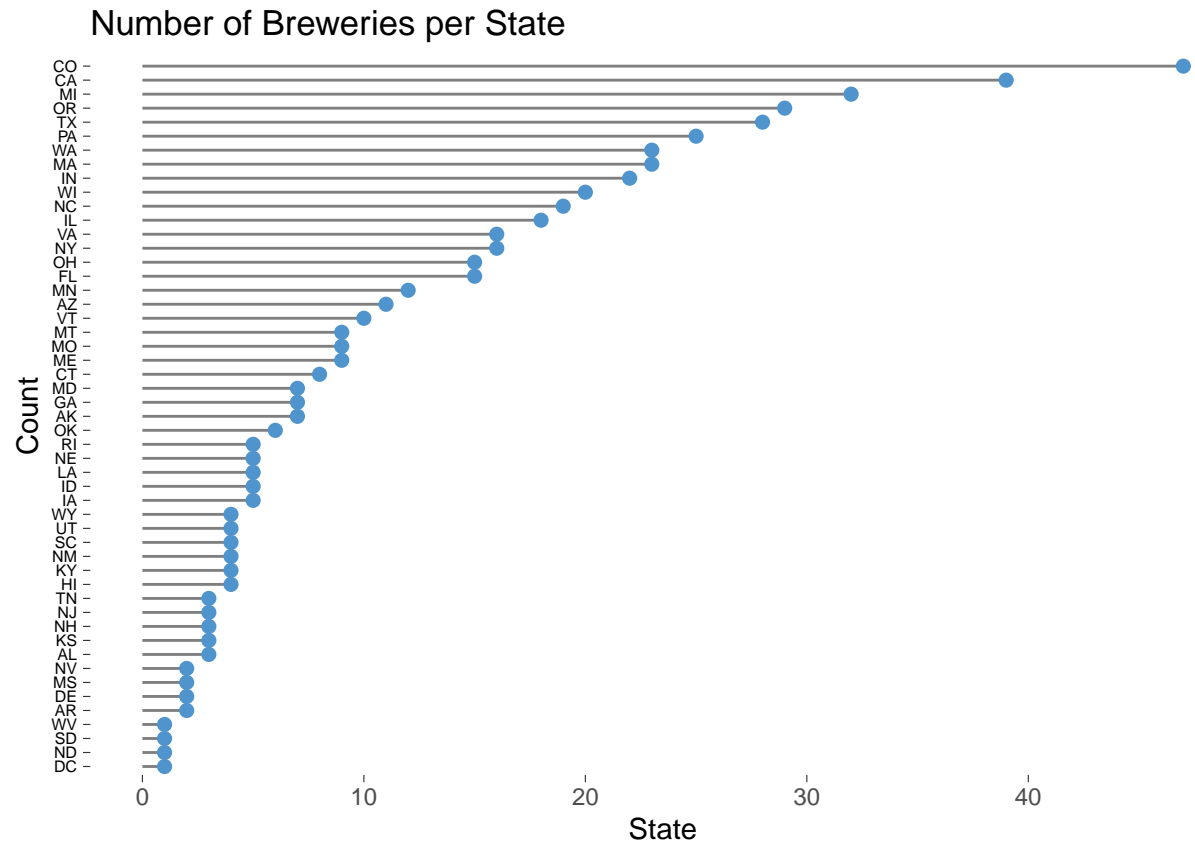
```
numbystate
```

```
##      State  n
## 1      AK   7
## 2      AL   3
## 3      AR   2
## 4      AZ  11
## 5      CA  39
## 6      CO  47
## 7      CT   8
## 8      DC   1
## 9      DE   2
## 10     FL  15
## 11     GA   7
## 12     HI   4
## 13     IA   5
## 14     ID   5
## 15     IL  18
## 16     IN  22
## 17     KS   3
## 18     KY   4
```

```
## 19    LA    5
## 20    MA   23
## 21    MD    7
## 22    ME    9
## 23    MI   32
## 24    MN   12
## 25    MO    9
## 26    MS    2
## 27    MT    9
## 28    NC   19
## 29    ND    1
## 30    NE    5
## 31    NH    3
## 32    NJ    3
## 33    NM    4
## 34    NV    2
## 35    NY   16
## 36    OH   15
## 37    OK    6
## 38    OR   29
## 39    PA   25
## 40    RI    5
## 41    SC    4
## 42    SD    1
## 43    TN    3
## 44    TX   28
## 45    UT    4
## 46    VA   16
## 47    VT   10
## 48    WA   23
## 49    WI   20
## 50    WV    1
## 51    WY    4
```

```
numbystate$State <- as.factor(numbystate$State)

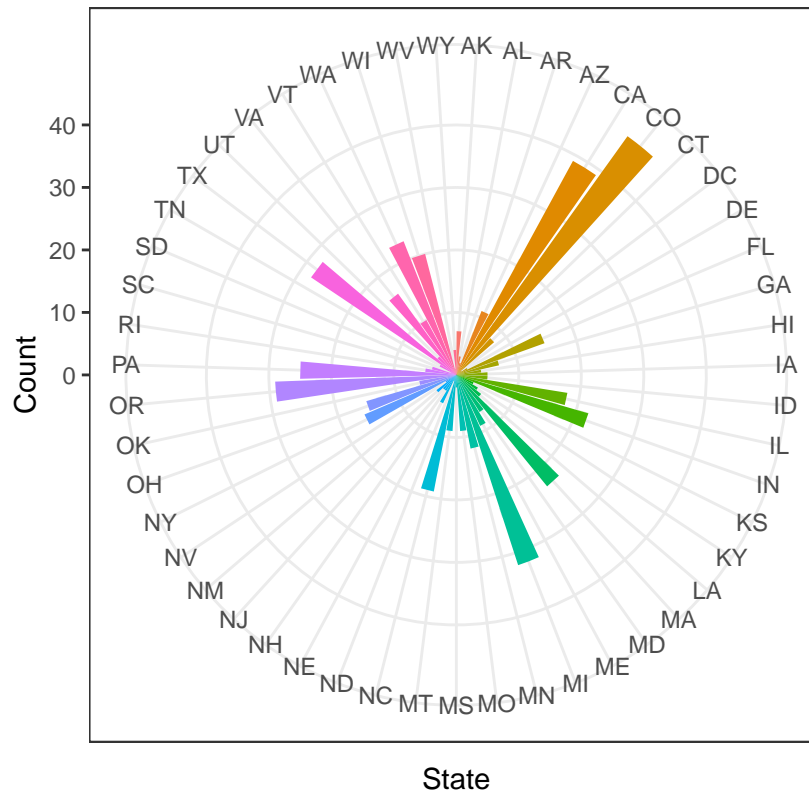
#bar graph depicting the number of breweries in each state
numbystate %>% ggplot(aes(x = reorder(State, n), y = n)) +
  geom_segment(aes(xend=State, yend=0), color = 'grey50') +
  geom_point(size=2, color="steelblue3") +
  coord_flip() +
  theme(legend.position = "none") +
  theme(axis.text.y = element_text(size = 6, color = "black")) +
  theme(panel.background = element_rect(fill = "white")) +
  theme(axis.ticks = element_line(size = .2)) +
  labs(x = "Count", y = "State") +
  ggtitle("Number of Breweries per State")
```



#Different chart (polar coordinates) giving a better visualization of relative amount of breweries in e

```
numbystate %>% ggplot(aes(x = State, y = n, fill = State)) +
  geom_bar(stat = 'identity') +
  coord_polar() + theme_bw() +
  theme(legend.position = "none") +
  labs(x = "State", y = "Count") + ggtitle("Number of Breweries per State")
```

Number of Breweries per State



A more accurate representation of brewery density is depicted by determining Breweries per Capita. D

```
statepop <- read_csv("https://raw.githubusercontent.com/j-dominguez9/Case-Study-1/main/statepop.csv")
```

```
## Rows: 50 Columns: 2
```

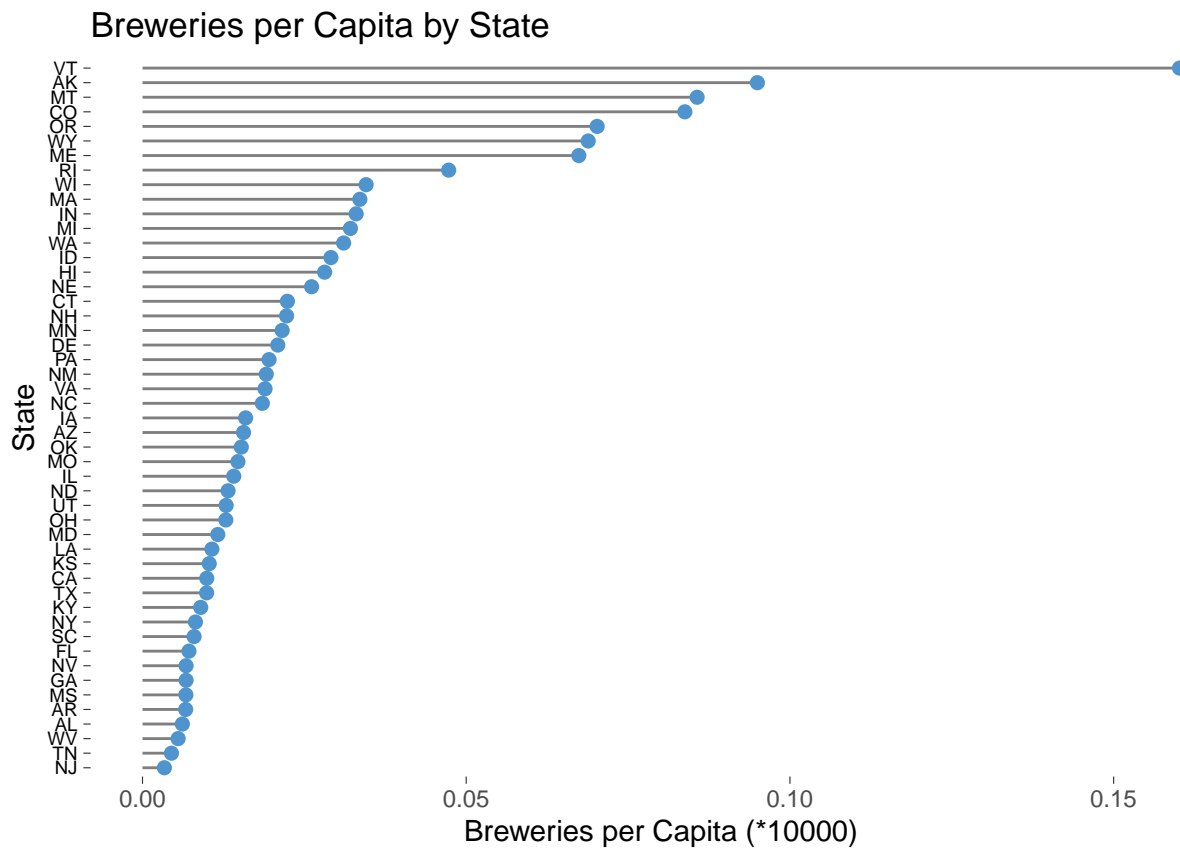
```
## -- Column specification -----
## Delimiter: ","
## chr (1): State
## dbl (1): population
```

```
##
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
```

```
#Join census data with previous table(numbystate)
breweriespercapita <- full_join(numbystate, statepop, by = "State") %>%
  mutate(bpc = (n/population)*10000) %>%
  mutate(State = as.factor(State))
```

```
#Plot
breweriespercapita %>% filter(!is.na(population)) %>% filter(!State == "SD") %>%
  ggplot(aes(x = reorder(State, bpc), y = bpc)) +
  geom_segment(aes(xend=State, yend=0), color = 'grey50') +
```

```
geom_point(size=2, color="steelblue3") +
coord_flip() +
theme(legend.position = "none") +
theme(axis.text.y = element_text(size = 7, color = "black")) +
theme(panel.background = element_rect(fill = "white")) +
theme(axis.ticks = element_line(size = .2)) +
ggtitle("Breweries per Capita by State") +
labs(x = "State", y = "Breweries per Capita (*10000)")
```



```
##Address the missing values in each column.
sum(is.na(breweries))
```

```
## [1] 0
```

#As we can see, the breweries data set holds no missing values, thus no need to eliminate any missing values.

```
sum(is.na(beers))
```

```
## [1] 1072
```

#The beers dataset has 1072, so we must eliminate noted values.

```
beers_clean <- beers %>% filter(!is.na(Name)) %>% filter(!is.na(Beer_ID)) %>%
  filter(!is.na(ABV)) %>% filter(!is.na(IBU)) %>%
```

```
filter(!is.na(Brewery_id)) %>% filter(!is.na(Style)) %>%
filter(!is.na(Ounces))
```

#Removed 1007 rows due to missing values.

##Compute the median alcohol content and international bitterness unit for each state. Plot a bar chart

#In order to join the two data sets successfully, we will need to find a primary key and a foreign key

```
colnames(breweries)
```

```
## [1] "Brew_ID" "Name"      "City"      "State"
```

```
colnames(beers_clean)
```

```
## [1] "Name"      "Beer_ID"    "ABV"        "IBU"        "Brewery_id"
## [6] "Style"     "Ounces"
```

#As we can see, Brewery ID would be a good key to join them; however, we must make sure that they have

```
breweries$Brewery = breweries$Name
breweries$Brewery_id = breweries$Brew_ID
head(breweries)
```

```
## # A tibble: 6 x 6
##   Brew_ID Name                City                State Brewery                Brewery_id
##   <dbl> <chr>                  <chr>                <chr> <chr>                <dbl>
## 1      1 NorthGate Brewing    Minneapolis          MN    NorthGate Br~          1
## 2      2 Against the Grain Brewery Louisville          KY    Against the ~          2
## 3      3 Jack's Abby Craft Lagers Framingham          MA    Jack's Abby ~          3
## 4      4 Mike Hess Brewing Company San Diego            CA    Mike Hess Br~          4
## 5      5 Fort Point Beer Company  San Francisco        CA    Fort Point B~          5
## 6      6 COAST Brewing Company    Charleston           SC    COAST Brewin~          6
```

#Let's create a new dataframe with the relevant columns from 'breweries' before joining.

```
breweries_clean <- breweries %>% select(Brewery, City, State, Brewery_id)
join <- inner_join(breweries_clean, beers_clean, by = "Brewery_id")
```

#From the joined data frame, let's select the relevant columns.

```
medABVIBU <- join %>% select(State, ABV, IBU)
head(medABVIBU)
```

```
## # A tibble: 6 x 3
##   State ABV IBU
##   <chr> <dbl> <dbl>
## 1 MN    0.045  50
## 2 MN    0.049  26
## 3 MN    0.048  19
## 4 MN    0.06   38
## 5 MN    0.06   25
## 6 MN    0.056  47
```

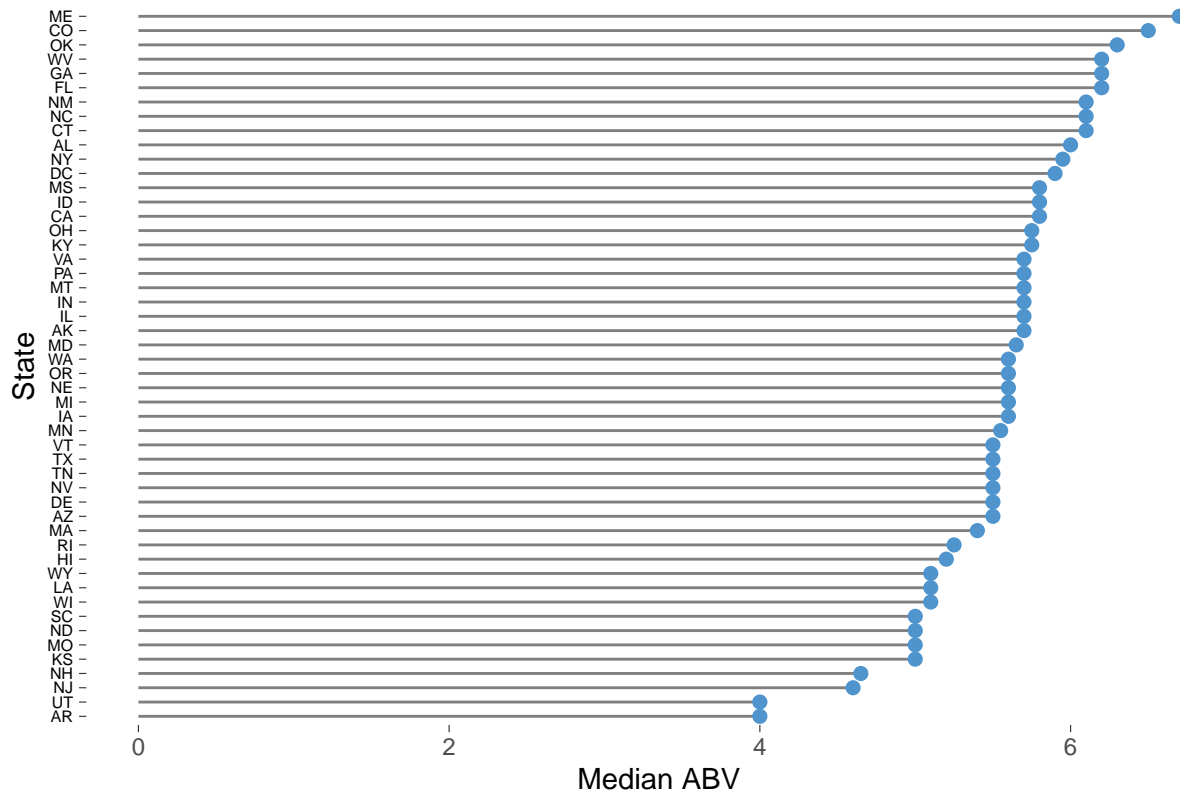
```
#Having the relevant columns to work with, lets create a new one with the median ABV of each state and
medABV <- medABVIBU %>% group_by(State) %>% mutate(medianABV = median(ABV)*100) %>% select(State, medianABV)
head(medABV)
```

```
## # A tibble: 6 x 2
## # Groups:   State [6]
##   State medianABV
##   <chr>      <dbl>
## 1 AK          5.7
## 2 AL           6
## 3 AR           4
## 4 AZ          5.5
## 5 CA          5.8
## 6 CO          6.5
```

```
##Data set is now ready to plot.
```

```
medABV %>% ggplot(aes(x = reorder(State, medianABV), y = medianABV)) +
geom_segment(aes(xend=State, yend=0), color = 'grey50') +
  geom_point(size=2, color="steelblue3") +
  coord_flip() +
  theme(legend.position = "none") +
  theme(axis.text.y = element_text(size = 6, color = "black")) +
  theme(panel.background = element_rect(fill = "white")) +
  theme(axis.ticks = element_line(size = .2)) +
  labs(x = "State", y = "Median ABV") +
  ggtitle("Median ABV by State")
```


Median ABV by State



##We will follow a similar process to derive median IBU by State.

```
medIBU <- medABVIBU %>%
  group_by(State) %>%
  mutate(medianIBU = median(IBU)) %>%
  select(State, medianIBU) %>%
  arrange(State) %>%
  distinct(State, medianIBU)

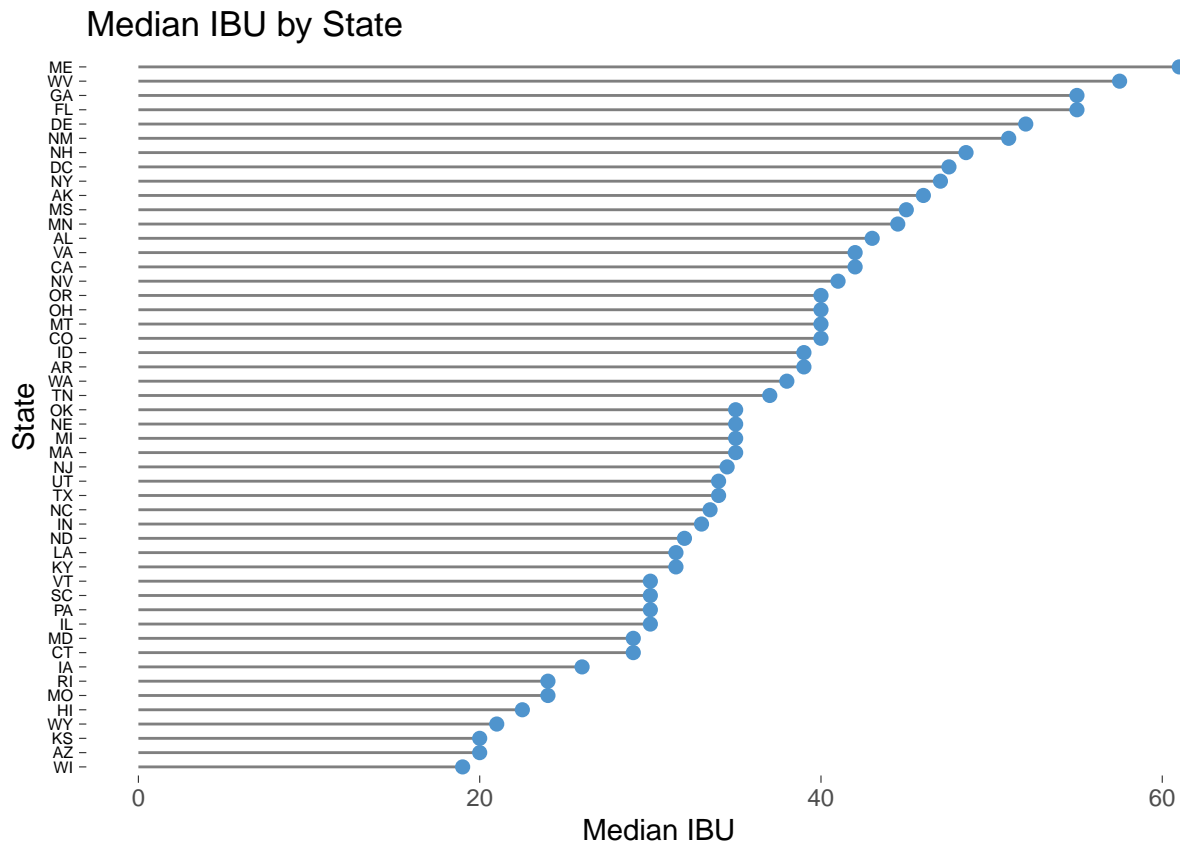
head(medIBU)
```

```
## # A tibble: 6 x 2
## # Groups:   State [6]
##   State medianIBU
##   <chr>      <dbl>
## 1 AK          46
## 2 AL          43
## 3 AR          39
## 4 AZ          20
## 5 CA          42
## 6 CO          40
```

#And plot in a similar manner.

```
medIBU %>% ggplot(aes(x = reorder(State, medianIBU), y = medianIBU)) +
```

```
geom_segment(aes(xend=State, yend=0), color = 'grey50') +
geom_point(size=2, color="steelblue3") +
coord_flip() +
theme(legend.position = "none") +
theme(axis.text.y = element_text(size = 6, color = "black")) +
theme(panel.background = element_rect(fill = "white")) +
theme(axis.ticks = element_line(size = .2)) +
labs(x = "State", y = "Median IBU") +
ggtitle("Median IBU by State")
```



##Which state has the maximum alcoholic (ABV) beer? Which state has the most bitter (IBU) beer?

#We will create a new data frame with relevant columns to explore this question.

```
maxABVIBU <- join %>% select(Brewery, Name, State, ABV, IBU)
```

#We need to identify the highest ABV level

```
max(maxABVIBU$ABV)
```

```
## [1] 0.125
```

```
maxABVIBU %>% filter(ABV == "0.125")
```

```
## # A tibble: 1 x 5
```

```
##   Brewery      Name      State  ABV    IBU
```

```
##      <chr>                <chr>          <chr> <dbl> <dbl>
## 1 Against the Grain Brewery London Balling KY    0.125    80
```

We can see that the beer with the highest ABV belongs to the state of Kentucky (KY) with an ABV of 1.

#We follow the same process for IBU.

```
max(maxABVIBU$IBU)
```

```
## [1] 138
```

```
maxABVIBU %>% filter(IBU == "138")
```

```
## # A tibble: 1 x 5
```

```
##   Brewery      Name      State  ABV  IBU
##   <chr>        <chr>    <chr> <dbl> <dbl>
## 1 Astoria Brewing Company Bitter Bitch Imperial IPA OR    0.082    138
```

#The beer with the highest IBU belongs to the state of Oregon (OR) with an IBU of 138. The beer is "Bit

##Comment on the summary statistics and distribution of the ABV variable.

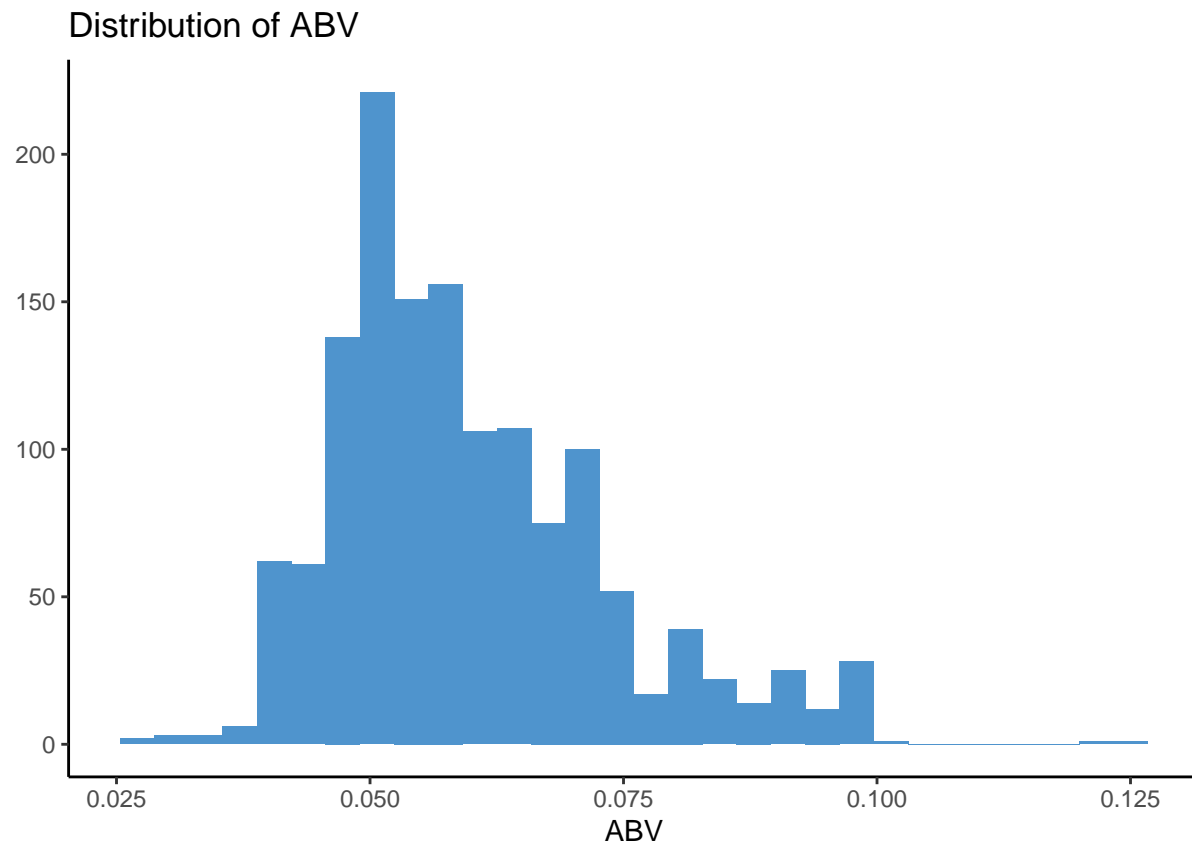
```
summary(maxABVIBU$ABV)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.02700 0.05000 0.05700 0.05992 0.06800 0.12500
```

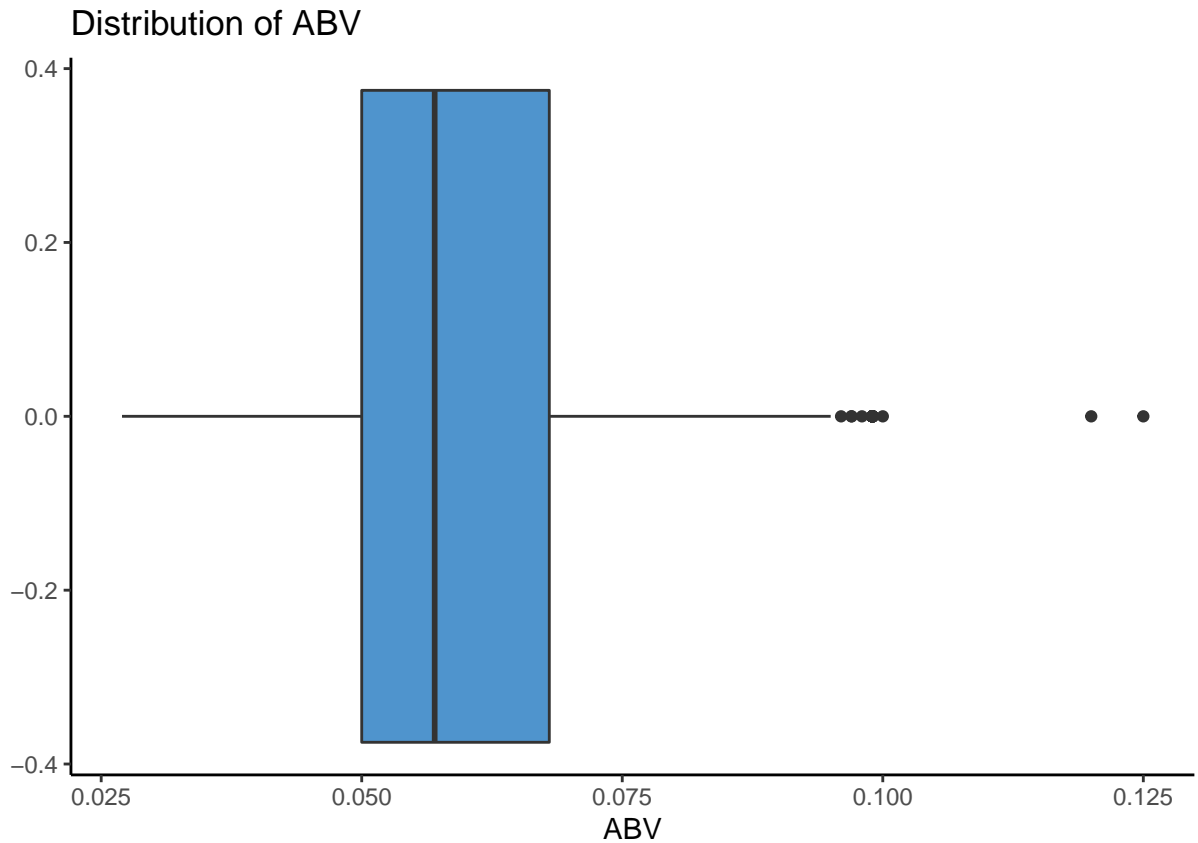
#Histogram

```
maxABVIBU %>% ggplot(aes(x = ABV)) +
  geom_histogram(fill = "steelblue3") +
  theme_classic() +
  labs(x = "ABV", y = "") +
  ggtitle("Distribution of ABV")
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```



```
#Boxplot
maxABVIBU %>% ggplot(aes(x = ABV)) +
  geom_boxplot(fill = "steelblue3") +
  theme_classic() +
  labs(x = "ABV", y = "") +
  ggtitle("Distribution of ABV")
```

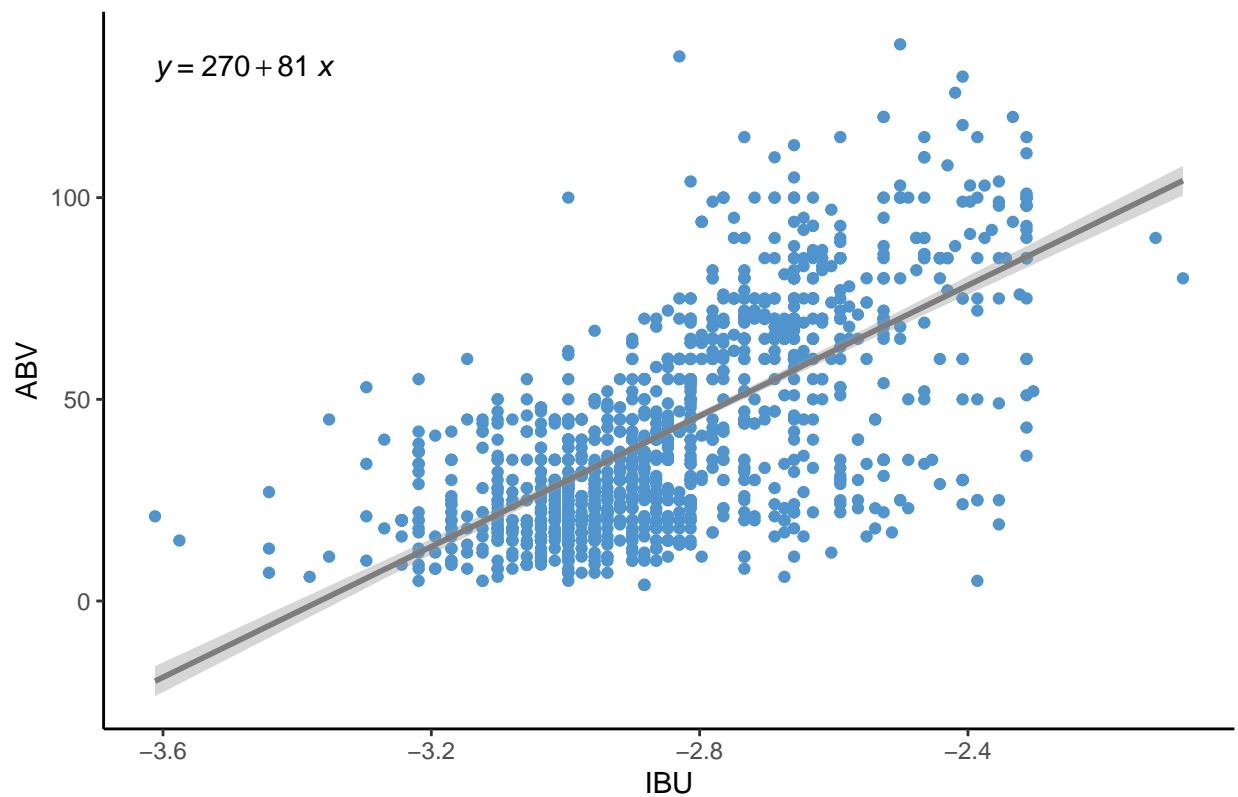


#As we saw earlier, the distributions of both the IBU and ABV columns were right skewed, it'd be helpful

```
maxABVIBU %>% ggplot(aes(x = log(ABV), y = IBU)) +  
  geom_point(color = "steelblue3") +  
  geom_smooth(method = "lm", color = "grey49") +  
  stat_regline_equation() +  
  theme_classic() +  
  labs(x = "IBU", y = "ABV") +  
  ggtitle("Relationship between IBU and ABV")
```

```
## 'geom_smooth()' using formula 'y ~ x'
```

Relationship between IBU and ABV



##From a visual inspection as well as a simple linear regression model, we can say that there is an app

```
cor.test(x = log(maxABVIBU$ABV), y = maxABVIBU$IBU)
```

```
##
## Pearson's product-moment correlation
##
## data: log(maxABVIBU$ABV) and maxABVIBU$IBU
## t = 34.032, df = 1401, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.6430317 0.7004025
## sample estimates:
##      cor
## 0.6727271
```

#This Pearson's correlation provides overwhelming evidence that there is a positive linear relationship

##Budweiser would also like to investigate the difference with respect to IBU and ABV between IPAs (Ind

```
sum(grepl("Ale", join$Style))
```

```
## [1] 559
```

```
sum(grepl("(India | IPA)", join$Style))
```

```
## [1] 395
```

```
IPAs <- join %>% filter(grepl("(India | IPA)", Style)) %>% filter(!grepl("Lager", Style))  
Ales <- join %>% filter(grepl("Ale", Style)) %>% filter(!grepl("(India | IPA)", Style))
```

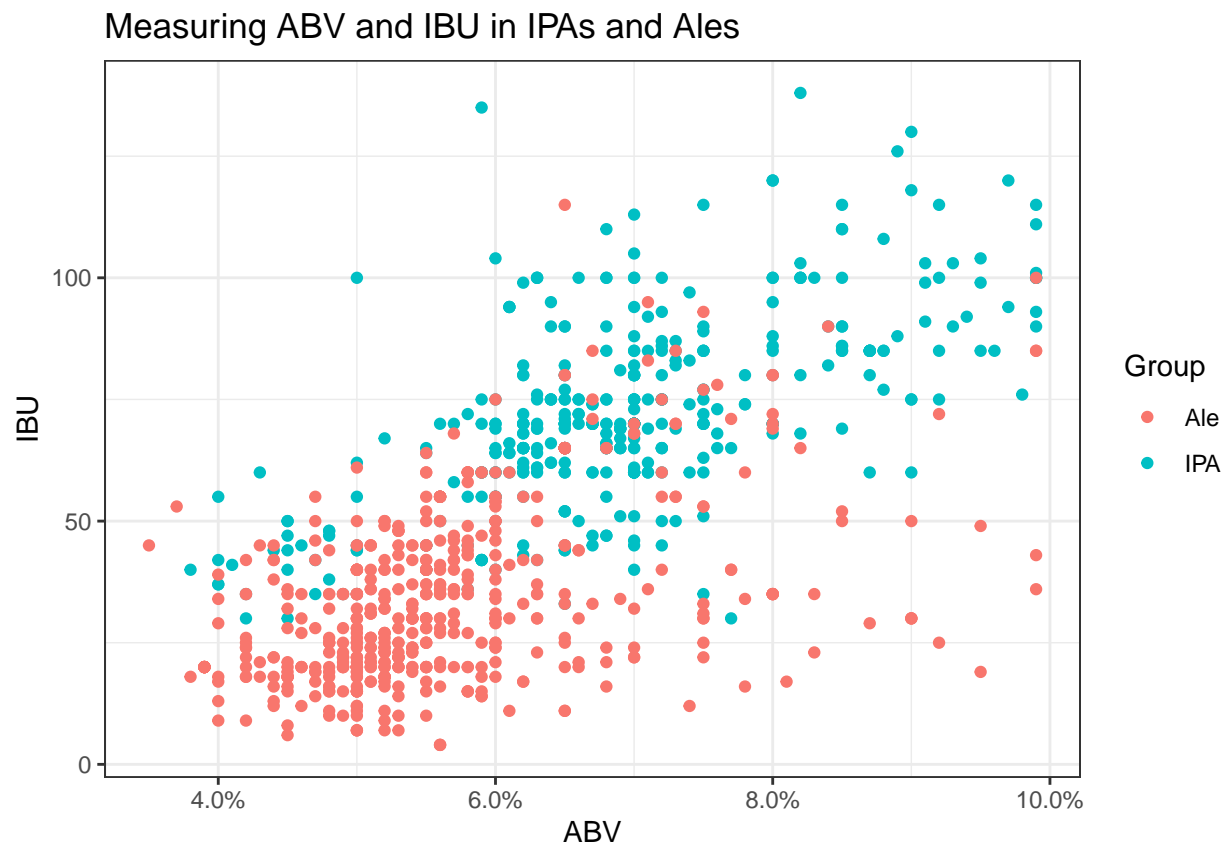
```
x <- data.frame(Group = "Ale", c(1:552)) %>% select(Group)  
final_Ales <- cbind(x, Ales)
```

```
y <- data.frame(Group = "IPA", c(1:392)) %>% select(Group)  
final_IPAs <- cbind(y, IPAs)
```

```
IPA_Ales <- full_join(final_IPAs, final_Ales)
```

```
## Joining, by = c("Group", "Brewery", "City", "State", "Brewery_id", "Name", "Beer_ID", "ABV", "IBU",
```

```
IPA_Ales %>% ggplot(aes(x = ABV, y = IBU, color = Group)) +  
  geom_point() +  
  scale_x_continuous(labels = scales::percent) +  
  theme_bw() +  
  ggtitle("Measuring ABV and IBU in IPAs and Ales")
```



```
## k-NN model. We first run a simple model to determine the accuracy of the model with k = 3.
```

```
confusionMatrix(table(knn.cv(IPA_Ales[,8:9], IPA_Ales$Group, k = 3), IPA_Ales$Group))
```

```
## Confusion Matrix and Statistics
```

```
##
```

```
##
```

```
##      Ale IPA
```

```
## Ale 486  70
```

```
## IPA  66 322
```

```
##
```

```
##              Accuracy : 0.8559
```

```
##              95% CI : (0.8319, 0.8777)
```

```
## No Information Rate : 0.5847
```

```
## P-Value [Acc > NIR] : <2e-16
```

```
##
```

```
##              Kappa : 0.7029
```

```
##
```

```
## McNemar's Test P-Value : 0.797
```

```
##
```

```
##              Sensitivity : 0.8804
```

```
##              Specificity : 0.8214
```

```
## Pos Pred Value : 0.8741
```

```
## Neg Pred Value : 0.8299
```

```
##              Prevalence : 0.5847
```

```
## Detection Rate : 0.5148
```

```
## Detection Prevalence : 0.5890
```

```
## Balanced Accuracy : 0.8509
```

```
##
```

```
## 'Positive' Class : Ale
```

```
##
```

```
##We must establish the optimal number for k, with regards to accuracy.
```

```
set.seed(1)
```

```
iterations = 500
```

```
numks = 50
```

```
masterAcc = matrix(nrow = iterations, ncol = numks)
```

```
for(j in 1:iterations)
```

```
{
```

```
  for(i in 1:numks)
```

```
  {
```

```
    CM = confusionMatrix(table(IPA_Ales$Group, knn.cv(IPA_Ales[,8:9], IPA_Ales$Group, k = i)))
```

```
    masterAcc[j,i] = CM$overall[1]
```

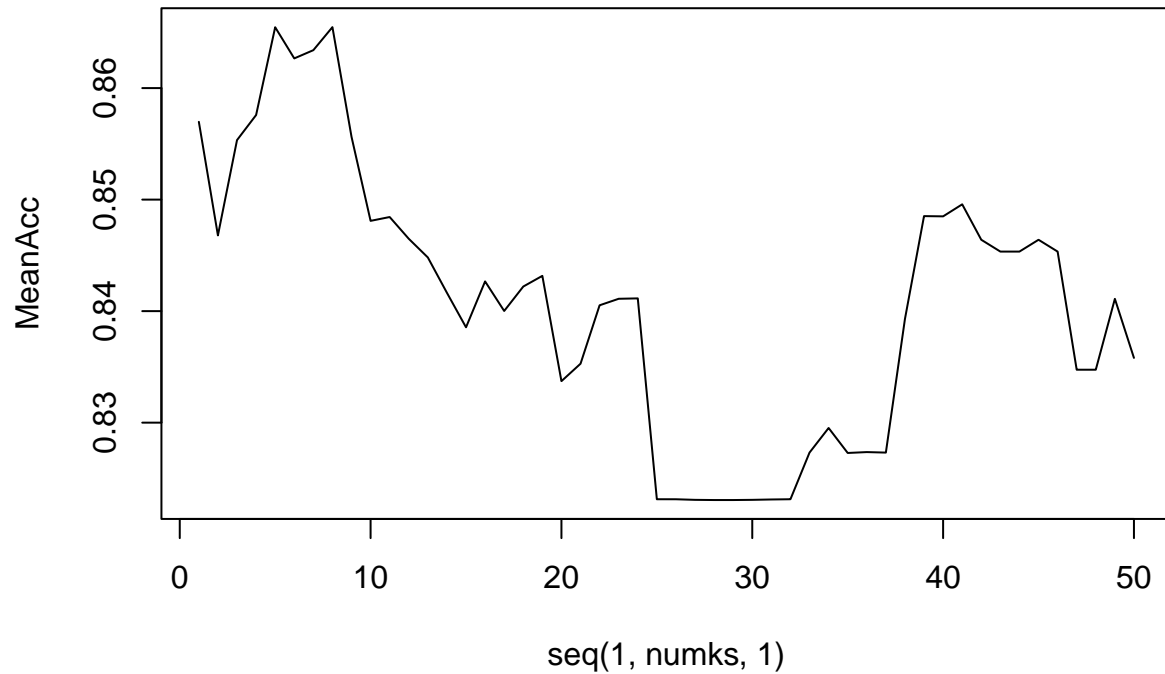
```
  }
```

```
}
```

```
MeanAcc = colMeans(masterAcc)
```



```
plot(seq(1,numks,1),MeanAcc, type = "l")
```



```
which.max(MeanAcc)
```

```
## [1] 8
```

```
max(MeanAcc)
```

```
## [1] 0.8654661
```

```
confusionMatrix(table(IPA_Ales$Group, knn.cv(IPA_Ales[,8:9], IPA_Ales$Group, k = 8)))
```

```
## Confusion Matrix and Statistics
```

```
##
```

```
##
```

```
##      Ale IPA
```

```
## Ale 498  54
```

```
## IPA  71 321
```

```
##
```

```
##              Accuracy : 0.8676
```

```
##              95% CI : (0.8443, 0.8886)
```

```
##      No Information Rate : 0.6028
```

```
##      P-Value [Acc > NIR] : <2e-16
```

```
##
##           Kappa : 0.7256
##
## Mcnemar's Test P-Value : 0.1524
##
##           Sensitivity : 0.8752
##           Specificity : 0.8560
##           Pos Pred Value : 0.9022
##           Neg Pred Value : 0.8189
##           Prevalence : 0.6028
##           Detection Rate : 0.5275
##           Detection Prevalence : 0.5847
##           Balanced Accuracy : 0.8656
##
##           'Positive' Class : Ale
##
```

#Naive Bayes

```
nbIPA_Ales <- IPA_Ales %>% select(Group, ABV, IBU)
naiveBayes(Group~., data = nbIPA_Ales )
```

```
##
## Naive Bayes Classifier for Discrete Predictors
##
## Call:
## naiveBayes.default(x = X, y = Y, laplace = laplace)
##
## A-priori probabilities:
## Y
##      Ale      IPA
## 0.5847458 0.4152542
##
## Conditional probabilities:
##      ABV
## Y      [,1]      [,2]
## Ale 0.05655616 0.01112430
## IPA 0.06914286 0.01216069
##
##      IBU
## Y      [,1]      [,2]
## Ale 34.33333 17.97471
## IPA 71.94898 19.54567
```

```
iterations = 100
masterAcc = matrix(nrow = iterations)
splitPerc = .8 #Training / Test split Percentage
for(j in 1:iterations)
{
  trainIndices = sample(1:dim(nbIPA_Ales)[1],round(splitPerc * dim(nbIPA_Ales)[1]))
  train = nbIPA_Ales[trainIndices,]
  test = nbIPA_Ales[-trainIndices,]
  model = naiveBayes(train[,2:3],train$Group)
  table(test$Group,predict(model,test[,2:3]))
}
```

```

CM = confusionMatrix(table(test$Group,predict(model,test[,2:3])))
masterAcc[j] = CM$overall[1]
}
MeanAcc = colMeans(masterAcc)
MeanAcc

```

```
## [1] 0.8419577
```

```
### t-test
```

```

Ales_ABV <- final_Ales %>% select(ABV, Group)
IPAs_ABV <- final_IPAs %>% filter(!grepl("Lager", Style)) %>% select(ABV, Group)

ABVjoin <- full_join(Ales_ABV, IPAs_ABV)

```

```
## Joining, by = c("ABV", "Group")
```

```
t.test(log(ABV) ~ Group, data = ABVjoin)
```

```

##
## Welch Two Sample t-test
##
## data: log(ABV) by Group
## t = -16.91, df = 849.95, p-value < 2.2e-16
## alternative hypothesis: true difference in means between group Ale and group IPA is not equal to 0
## 95 percent confidence interval:
## -0.2260548 -0.1790364
## sample estimates:
## mean in group Ale mean in group IPA
## -2.889941 -2.687395

```

```
t.test(IBU ~ Group, data = IPA_Ales)
```

```

##
## Welch Two Sample t-test
##
## data: IBU by Group
## t = -30.118, df = 797.55, p-value < 2.2e-16
## alternative hypothesis: true difference in means between group Ale and group IPA is not equal to 0
## 95 percent confidence interval:
## -40.06727 -35.16402
## sample estimates:
## mean in group Ale mean in group IPA
## 34.33333 71.94898

```

```
##NaiveBayes State, Style (IPA, Ales)
```

```

nb2 <- IPA_Ales %>% select(State, Group) %>% mutate(State = as.factor(State))
model <- naiveBayes(Group~., data = nb2)
predict(model, data.frame(State = "NH"), type = 'raw')

```

```
##           Ale           IPA
## [1,] 0.3556701 0.6443299
```

```
####NB State, City, Style (all). Find most popular Style by City
df <- join %>% select(Style, City, State)
model <- naiveBayes(Style~., data = df)
predict(model, data.frame(State = "CO", City = "Buena Vista"))
```

```
## [1] American IPA
## 90 Levels: Abbey Single Ale Altbier ... Witbier
```

```
####NB Find most popular Style by State
df2 <- df %>% select(Style, State)
model <- naiveBayes(Style~State, data = df2)
predict(model, data.frame(State = "CO"))
```

```
## [1] American Pale Ale (APA)
## 90 Levels: Abbey Single Ale Altbier ... Witbier
```

```
#### NB Find most popular City for Style
```

```
df1 <- df %>% select(Style, City)
model <- naiveBayes(City~Style, data = df1)
pred <- predict(model, df1$Style)
CM <- confusionMatrix(as.factor(pred), as.factor(df1$City))
predict(model, data.frame(Style = "American IPA"))
```

```
## [1] San Diego
## 281 Levels: Abingdon Abita Springs Afton Albuquerque Anchorage ... York
```

```
####NB find most popular State for style
```

```
model <- naiveBayes(State~Style, data = df2)
pred <- predict(model, df2$Style)
CM <- confusionMatrix(as.factor(pred), as.factor(df2$State))
predict(model, data.frame(Style = "American IPA"))
```

```
## [1] CA
## 50 Levels: AK AL AR AZ CA CO CT DC DE FL GA HI IA ID IL IN KS KY LA MA ... WY
```

```
content <- read_html("https://vinepair.com/articles/best-highest-rated-beer-every-state/")
tables <- content %>% html_table(fill = TRUE)
first_table <- tables[[1]]
first_table <- first_table[-1,]
topbeer <- first_table %>% rename(State = X1, Beer = X2)
topbeer <- topbeer %>% cbind(data.frame(state = state.abb[match(topbeer$State, state.name)]))
topbeer[is.na(topbeer)] <- "DC"
topbeer <- topbeer %>%
  select(state, Beer) %>%
  rename(Style = state) %>%
  separate(Beer, into = c("Name", "Brewery", "Style"), sep = "\n") %>%
  separate(Style, into = c("Style", "ABV"), sep = "([|])")
```

```
## Warning: Expected 2 pieces. Missing pieces filled with 'NA' in 1 rows [39].
```

```
head(topbeer)
```

```
##      State                Name                Brewery
## 1      AL              El Gordo    Good People Brewing Company
## 2      AK              Blessed    Anchorage Brewing Company
## 3      AZ White Russian Imperial Stout    Sun Up Brewing Co.
## 4      AR              BDCS        Ozark Beer Co.
## 5      CA      Pliny The Younger Russian River Brewing Company
## 6      CO      Medianoche - Coconut    WeldWerks Brewing Co.
##                Style      ABV
## 1 Stout - Russian Imperial  13.90%
## 2 Stout - American Imperial  14.00%
## 3 Stout - American Imperial   9.20%
## 4 Stout - American Imperial  10.20%
## 5              IPA - Imperial  10.25%
## 6 Stout - American Imperial  14.10%
```

```
#Tableau Workbook URL: https://public.tableau.com/shared/4QFJCDFGZ?:display\_count=n&origin=viz\_share\_l
```