# Mine-Craft: Data Mining and Analysis of Craft Beers in America

James Ingram Joshua Shusterman Spring 2018

# Contents

Background	3
Data Source	3
Our Goal	3
Initial Approach	3
Updated Approach	3
Data Cleaning and Pre-Processing	3
Exploratory Data Analysis	4
"Alcohol by Volume" variable	4
"International Bittering Units" variable	5
"Ounces" variable	6
"State" variable	7
Looking at ABV and IBU together	9
Classification Models	10
Decision Tree Models (RWeka J48)	10
Naïve Bayes Models	11
Support Vector Machine Models	13
Model Comparison	15
Conclusion	15
Appendix: R Code	16

# Background

Craft beers are gaining in popularity in the United States. From 2006 to 2016, craft beers grew their market share nearly 9% against "Big Beer." Local brewpubs are popular as restaurants that brew their own craft beer onsite, and may traditionally only distribute in small quantities directly to customers.

Beers are categorized into different styles based on a number of factors. Some of these include the brewing and fermenting process such as being brewed hot or cold, and what ingredients are used. This determines the beer's color, as well as alcohol content by volume (ABV) and international bittering units (IBU).

### **Data Source**

The dataset was chosen from Kaggle, a data science project hosting site that uses a competition format to share datasets, problems, and work toward solutions. One of the datasets hosted on the site is a set of craft beer data.<sup>2</sup>

The data is contained in two files. The first file contains data for over 2,000 beers, and the second file contains data for over 500 breweries. The datasets include the name of the beer, style of the beer, international bittering units (IBU), alcohol content by volume (ABV), and the volume of the beer.

#### Our Goal

The goal of this project is to perform exploratory data analysis and to create classification models that can predict the style of a beer based on its characteristics.

# Initial Approach

We began with the full dataset available on Kaggle. This dataset included 100 classes of beer styles. Since there are so many overlapping styles with similar ABV and IBU levels, only very low accuracy models could be generated from the full dataset.

# Updated Approach

Because the full dataset led to models with such low accuracy, we limited our approach to consider only three popular styles of beer that were selected based on results of our descriptive statistical analysis. A new dataset containing only the "American Amber", "American Double", and "American IPA" styles of beer was created.

# Data Cleaning and Pre-Processing

After removing the NAs from the dataframe, row numbers were reset. Beers\$Column1 and beers\$id were removed because they weren't necessary for analysis. The vector beers\$name was renamed to beers\$beer\_name to avoid confusion with

<sup>&</sup>lt;sup>1</sup> GuruFocus.com: Harding loevner commentary: Craft beer is going flat. can craft spirits continue the insurgency? (2017). . Chatham: Newstex. Retrieved from https://search-proquest-com.libezproxy2.syr.edu/docview/1933659154?accountid=14214

<sup>&</sup>lt;sup>2</sup> https://www.kaggle.com/nickhould/craft-cans/home

breweries\$name in subsequent dataframe merging. In the breweries dataframe, "X" was renamed to breweries\$brewery\_id (to match the column in "beers") and breweries\$name was changed to breweries\$brewery\_name. The two dataframes were merged using a left outer join on "brewery\_id". This resulted in the information from the breweries dataframe being merged with the beers dataframe where they had corresponding "brewery\_id"s. Once merged, the brewery\_id column was removed and the columns were re-ordered to create the unified dataframe.

While examining the data, it was noticed that most beer volumes were in multiples of 4 oz, however a few examples had 8.4, 16.9, and 19.2 oz volumes. We decided to transform these into 8, 16, and 20 ounces to conform them with the rest of the data. At this stage, our dataset was considered cleaned and ready for exploratory data analysis.

# **Exploratory Data Analysis**

First, we looked at the frequency of beer styles in our finished dataset. We then decided to explore the beer attributes by looking at their frequency distributions, ranges, and measures of central tendency, starting with "ABV".

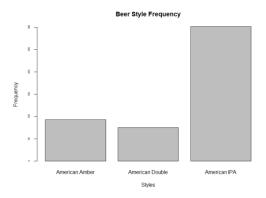


Figure 1: Frequency Distribution of Beer Styles in Data Set

# "Alcohol by Volume" variable

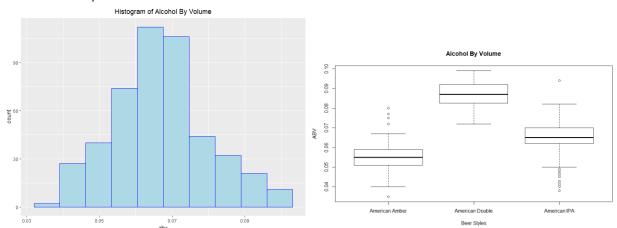


Figure 2: Frequency Distribution of Alcohol by Volume

Figure 3: Median and Range of Alcohol by Volume for Each Beer Style

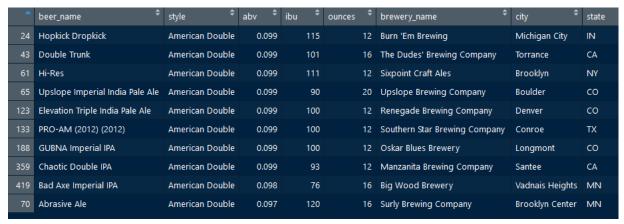


Table 1: Top 10 Beers Ranked by Highest Alcohol by Volume

In Figure 2 we can see that the average ABV is approximately 0.06. Table 1 indicates that all of the top 10 beers ranked by ABV are American Double style. While there is some overlap, we can see in Figure 3 that the three styles have distinct ABV levels, with American Amber having the lowest ABV and American Double having the highest.

# "International Bittering Units" variable

Next, we examined International Bittering Units (ibu). The results for analysis of IBU seem to match that of ABV pretty closely. In this case, however, there is an American IPA among the top 10 beers ranked by IBU.

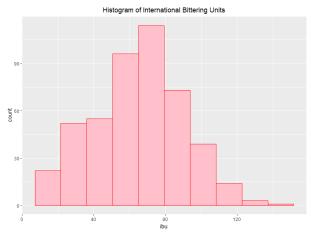


Figure 4: Frequency Distribution of International Bittering Units

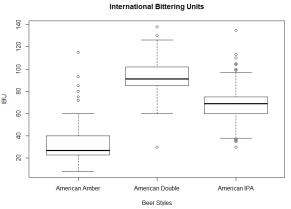


Figure 5: Median and Range of International Bittering Units for Each Beer Style

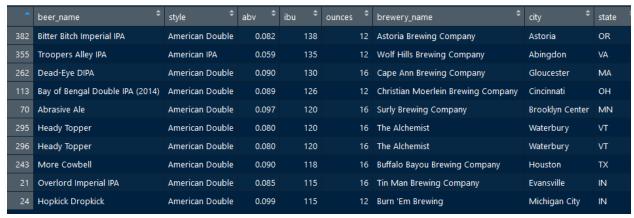
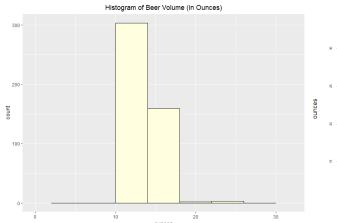


Table 2: Top 10 Beers Ranked by Highest International Bittering Units

Again, while there is some overlap, we can see that the three styles have distinct IBU levels. These seem to mirror the ABV level, which implies a correlation between ABV and IBU.

# "Ounces" variable

We next examined the distribution and range of the volume of alcohol (in ounces) among the beers. As expected, the 12 oz. size is the most common, with pints being the next common size. When we ranked the top ten beers by volume, there was a lot of diversity among styles, ABV levels, and IBU levels. There may not be a correlation between size of the can of beer and any other attribute. Ounces does not appear to be a useful attribute for our analysis.



**Figure 6: Frequency Distribution of Ounces** 

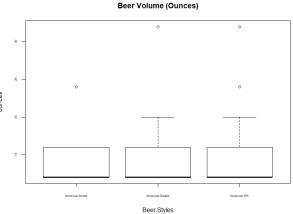


Figure 7: Median and Range for Volume (in Ounces) for Each Beer Style

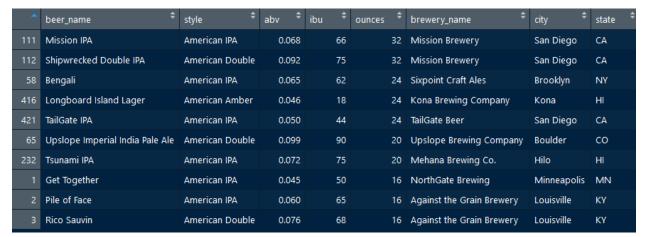


Table 3: Top Ten Beers Ranked by Highest Volume (in Ounces)

# "State" variable

We examined the relationship between the brewery state and the style of beer to see if this variable would be useful for classification purposes. First, we looked at the distribution of beers per state and then the presence or absence of the beer styles in each state.

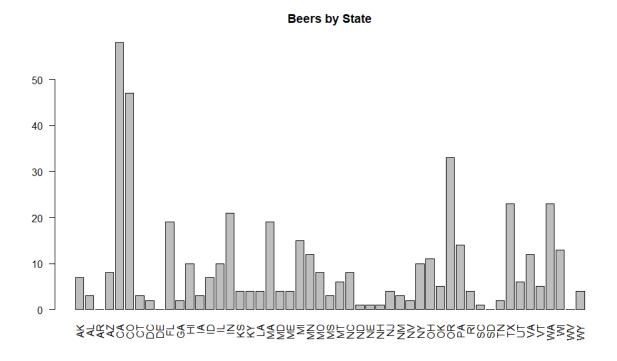


Figure 8: Count of Unique Beers Per State

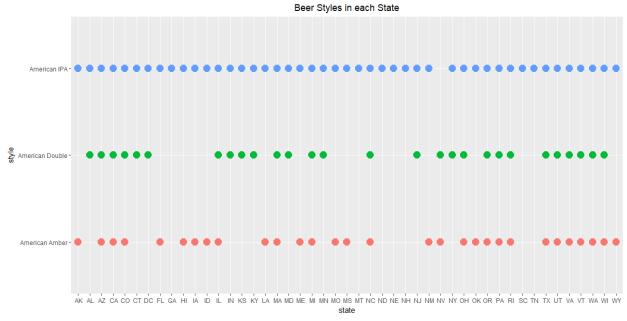


Figure 9: Presence or Absence of Each Beer Style Per State

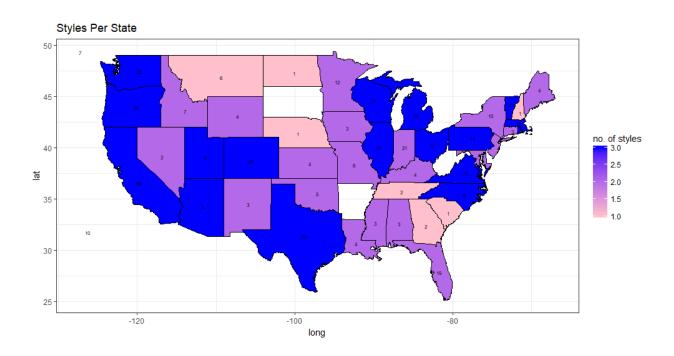


Figure 10: Beer Styles Per State (On US Map)

Very few states have only one style, and no style is unique to a particular state. This means that the state variable may not be useful in classification model building. Initial classification models were built including the "state" variable; however, they did not result in increased accuracy. Therefore, this variable was left out of our final models.

# Looking at ABV and IBU together

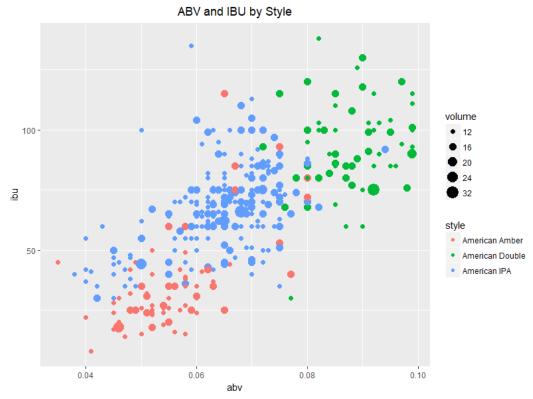


Figure 11: Scatterplot of Alcohol by Volume and International Bittering Units

Figure 11 shows that the chosen styles form three clusters based on ABV and IBU. For this reason, we chose to focus on only these two attributes to build the most parsimonious classification models.

### Classification Models

For classification models, we created a trainData dataframe containing only the beer style, ABV, and IBU. We also constructed a training (new\_train) and testing (new\_validation) data set with a 2/3 cut point.

# Decision Tree Models (RWeka J48)

The initial model was a pruned tree using the trainData set to classify the beer style. It had a size of 13, with 7 leaves. We performed 10-fold cross-validation to evaluate the model's accuracy (~88.7%). We attempted to improve the model by increasing the confidence level from 0.25 to 0.5. The same decision tree was produced with about the same accuracy. To discover the effect of pruning, an unpruned tree was created. The size of the unpruned tree was 23, with 12 leaves. This model produced an accuracy of ~87.6%. According to our calculated Information Gain results (to see the importance of each variable in the decision tree) ABV was used first and IBU second in the models.

```
abv <= 0.075
| ibu <= 35
| | ibu <= 28: American Amber (48.0)
| | ibu > 28
| | | abv <= 0.05: American IPA (9.0/3.0)
| | | abv > 0.05: American Amber (15.0/1.0)
| ibu > 35: American IPA (310.0/27.0)
| abv > 0.075
| abv <= 0.08
| | ibu <= 91: American IPA (18.0/9.0)
| | ibu > 91: American Double (5.0)
| abv > 0.08: American Double (64.0/2.0)

Number of Leaves : 7

Size of the tree : 13
```

Figure 12: J48 Decision Tree (default parameters)

```
pctCorrect pctIncorrect pctUnclassified kappa 88.6993603 11.3006397 0.0000000 0.7770953 meanAbsoluteError rootMeanSquaredError relativeAbsoluteError rootRelativeSquaredError 0.1091163 0.2529076 31.2206891 60.5559701
```

Figure 13: 10-fold CV Evaluation of J48 Decision Tree (default parameters)

```
Information Gain:
'abv' 'ibu'
0.7400649 0.5716396
```

Figure 14: Information Gain

## Naïve Bayes Models

### RWeka NB

Using the RWeka NB package, we created the following classification model:

Attribute	Class American Amber American (0.2)	Double (0.16)	American IPA (0.64)
abv			
mean	0.0555	0.0877	0.0648
std. dev.	0.0085	0.0067	0.0085
weight sum	93	75	301
precision	0.0011	0.0011	0.0011
ibu			
mean	33.9567	93.2494	67.5307
std. dev.	18.8665	17.3454	15.9636
weight sum	93	75	301
precision	1.4607	1.4607	1.4607

Figure 15: RWeka Naïve Bayes Model

10-fold CV evaluation demonstrated that the model's accuracy was about 87.6%.

pctCorrect	pctIncorrect	pctUnclassified	kappa
87.6332623	12.3667377	0.0000000	0.7677266
meanAbsoluteError	rootMeanSquaredError	relativeAbsoluteError	rootRelativeSquaredError
0.1137663	0.2575620	32.5511799	61.6704085

Figure 16: 10-fold CV Evaluation of RWeka Naïve Bayes Model

# Naïve Bayes using e1071

Next, we built a Naïve Bayes model using R's e1071 package.

```
Naive Bayes Classifier for Discrete Predictors

Call:
naiveBayes.default(x = X, y = Y, laplace = laplace)

A-priori probabilities:
Y
American Amber American Double American IPA
0.1982942 0.1599147 0.6417910

Conditional probabilities:
abv
Y
[,1] [,2]
American Amber 0.05562366 0.008449227
American Double 0.08769333 0.006655567
American IPA 0.06480731 0.008494473

ibu
Y
[,1] [,2]
American Amber 34.05376 18.90041
American Double 93.32000 17.53079
American IPA 67.63455 16.05260
```

Figure 17: e1071 Naïve Bayes Model

Using the model, we predicted the beer styles of the data from the trainData set. The model seems fairly accurate.

NB_pred	American Amber	American Double	American IPA
American Amber	74	0	33
American Double	0	69	3
American IPA	19	6	265

Figure 18: Confusion Matrix from NB Model "style" Prediction of "trainData" Data Set

Similar results were obtained when the model was applied to the "new\_train" and "new\_validation" data sets.

American Amber 28 0 8 American Double 0 18 1	NB_pred2	American Am	mber American	Double American	IPA
American Double 0 18 1	American Amb	er	28	0	8
	American Dou	ble	0	18	1
American IPA 6 3 93	American IPA		6	3	93

Figure 19: Confusion Matrix from NB Model "style" Prediction of "new\_validation" Data Set

# Support Vector Machine Models

## Linear SVM Model

A linear SVM model was created using "new\_train" data with a training accuracy of over 91%.

```
Support Vector Machines with Linear Kernel

312 samples
2 predictor
3 classes: 'American Amber', 'American Double', 'American IPA'

No pre-processing
Resampling: Bootstrapped (25 reps)
Summary of sample sizes: 312, 312, 312, 312, 312, ...
Resampling results:

Accuracy Kappa
0.914551 0.8327833

Tuning parameter 'C' was held constant at a value of 1
```

Figure 20: Linear SVM Model

Applying this model to predict the "new\_validation" data, we obtained results with ~91% accuracy, as can be seen in the confusion matrix and summary statistics below:

```
Confusion Matrix and Statistics
                   Reference
Prediction
                    American Amber American Double American IPA
  American Amber
                                  26
                                                    18
  American Double
                                                                   99
  American IPA
Overall Statistics
     Accuracy : 0.9108
95% CI : (0.8549, 0.9504)
No Information Rate : 0.6497
     P-Value [Acc > NIR] : 2.927e-14
 Kappa : 0.819
Mcnemar's Test P-Value : 0.1767
 Statistics by Class:
                       Sensitivity
 Specificity
Pos Pred Value
Neg Pred Value
                                         0.8966
                                                                  0.9474
0.9783
                                                                                         0.9083
0.9375
                                         0.9375
                                                                   0.9783
0.1338
0.1146
0.1210
0.9249
Prevalence
                                                                                         0.6497
Detection Rate
                                         0.1656
                                                                                         0.6306
Detection Prevalence
Balanced Accuracy
                                                                                         0.6943
                                         0.1847
```

Figure 21: Linear SVM Model Prediction Results

## Radial SVM Model

A non-linear (RBF) Support Vector Machine model was also created using the "new\_train" data set with varying levels of Confidence (0.25, 0.5, and 1.0). Accuracy levels of 90.9, 90.9, and 91.2% were obtained, respectively.

```
Support Vector Machines with Radial Basis Function Kernel

312 samples
2 predictor
3 classes: 'American Amber', 'American Double', 'American IPA'

No pre-processing
Resampling: Bootstrapped (25 reps)
Summary of sample sizes: 312, 312, 312, 312, 312, 312, ...
Resampling results across tuning parameters:

C Accuracy Kappa
0.25 0.9087421 0.8203480
0.50 0.9086468 0.8212055
1.00 0.9123640 0.8289506

Tuning parameter 'sigma' was held constant at a value of 2.861614
Accuracy was used to select the optimal model using the largest value.
The final values used for the model were sigma = 2.861614 and C = 1.
```

Figure 22: Radial SVM Model

Again, the model was applied to predict the beer styles from the "new\_validation" data set. The classification model was approximately 90.4% accurate as can be seen below:

```
Confusion Matrix and Statistics
                  Reference
Prediction
                   American Amber American Double American IPA
                                                               0
  American Amber
                               24
                                                 0
  American Double
                                                 18
                                                             100
  American IPA
Overall Statistics
                Accuracy: 0.9045
    95% CI : (0.8473, 0.9455)
No Information Rate : 0.6497
    P-Value [Acc > NIR] : 1.53e-13
                   Kappa : 0.8032
 Mcnemar's Test P-Value : 0.01694
Statistics by Class:
                      Class: American Amber Class: American Double Class: American IPA
                                                              0.8571
Sensitivity
                                      0.7059
                                                                                   0.9804
                                                                                   0.7818
Specificity
                                      1.0000
                                                              0.9779
                                                              0.8571
Pos Pred Value
                                      1.0000
                                                                                   0.8929
Neg Pred Value
                                      0.9248
                                                              0.9779
                                                                                   0.9556
                                                                                   0.6497
Prevalence
                                      0.2166
                                                              0.1338
Detection Rate
                                      0.1529
                                                              0.1146
                                                                                   0.6369
Detection Prevalence
                                      0.1529
                                                              0.1338
                                                                                   0.7134
Balanced Accuracy
                                      0.8529
                                                              0.9175
                                                                                   0.8811
```

Figure 23: Radial SVM Model Prediction Results

# Model Comparison

All three types of models produced similar results (approx. 88-91% accuracy), with SVM being the most accurate

# Conclusion

Once we chose to focus on only 3 styles of beer, and only 2 characteristics, we were able to create classification models that had much higher levels of accuracy than we had when we used the full dataset. When there are larger numbers of classes in the dataset, there is more overlap among the classes, which leads to lower accuracy in the classification models. The three classes that we chose had enough distinction to allow for high levels of accuracy.

# Appendix: R Code

```
# Install Packages and Load Libraries
packages <- c("tidyverse", "RWeka","rJava","caret","e1071","kernlab", "randomForest", "class",</pre>
"cluster", "ggplot2", "ggmap", "maps", "mapdata", "maptools", "rstudioapi")
package.check <- lapply(packages, FUN = function(x){</pre>
  if(!require(x, character.only=TRUE)){
   install.packages(x, dependencies=TRUE)
   library(x, character.only=TRUE)
})
# Data Load and Pre-Processing
setwd(dirname(getActiveDocumentContext()$path))
beers <- read.csv("beers_new.csv")</pre>
breweries <- read.csv("breweries.csv")</pre>
# examine datasets
str(beers)
sum(is.na(beers)) # find total NAs in beers data set
sum(is.na(breweries)) # find total NAs in breweries data set
# omit NAs
beers <- na.omit(beers)</pre>
rownames(beers) <- NULL # resets row counts after NAs are removed
# remove beers$Column1 and beers$id
beers <- beers[, -1,]</pre>
beers <- beers[, -3]</pre>
head(beers) # examine structure of beers dataframe
# rename beers$name to beers$beer_name
cnames <- colnames(beers)</pre>
cnames[3] <- "beer_name"</pre>
colnames(beers) <- cnames</pre>
colnames(beers)
# rename breweries$X to breweries$brewery id and breweries$name to breweries$brewery name
cnames <- colnames(breweries)</pre>
cnames[1] <- "brewery_id"</pre>
cnames[2] <- "brewery_name"</pre>
colnames(breweries) <- cnames</pre>
colnames(breweries)
# merge beers and breweries data sets on brewery id
df <- merge(x = beers, y = breweries, by = "brewery id")</pre>
# re-order column headings and remove brewery id from dataframe
df \leftarrow df[,c(4,5,2,3,6,7,8,9)]
View(head(df,25))
# standardize ounces
df$ounces[df$ounces==8.4]<-8
```

```
df$ounces[df$ounces==16.9]<-16</pre>
df$ounces[df$ounces==19.2]<-20</pre>
# create new csv file containing cleaned data set
write.csv(df, file="clean_beer_data.csv")
###############################
# Exploratory Data Analysis
# frequency of each beer style
barplot(table(df$style), main="Beer Style Frequency", xlab="Styles", ylab="Frequency", cex.axis=0.5)
# ABV
# display frequency distribution of abv
g.abv <- ggplot(df, aes(x=abv))</pre>
g.abv <- g.abv + geom histogram(bins=10, color = "blue", fill="light blue")</pre>
g.abv <- g.abv + ggtitle("Histogram of Alcohol By Volume")</pre>
g.abv <- g.abv + theme(plot.title = element_text(hjust = 0.5))</pre>
g.abv
# sort beers with highest alcohol by volume (display top 10)
sortedAbv <- df[order(-df$abv), ]</pre>
View(head(sortedAbv,10))
# Look at ABV by Beer Style
plot(x=df$style, y=df$abv, xlab="Beer Styles", ylab="ABV", cex.axis=1, main="Alcohol By Volumne")
# display frequency distribution of ibu
g.ibu <- ggplot(df, aes(x=ibu))</pre>
g.ibu <- g.ibu + geom_histogram(bins=10, color = "red", fill="pink")</pre>
g.ibu <- g.ibu + ggtitle("Histogram of International Bittering Units")</pre>
g.ibu <- g.ibu + theme(plot.title = element_text(hjust = 0.5))</pre>
g.ibu
# sort beers with highest international bitter units (display top 10)
sortedIbu <- df[order(-df$ibu), ]</pre>
View(head(sortedIbu, 10))
# Look at IBU by Beer Style
plot(x=df$style, y=df$ibu, xlab="Beer Styles", ylab="IBU", cex.axis=1, main="International Bittering
Units")
# display frequency distribution of ounces
g.ounces <- ggplot(df, aes(x=ounces))</pre>
g.ounces <- g.ounces + geom_histogram(binwidth = 4, color = "black", fill="light yellow")</pre>
g.ounces <- g.ounces + ggtitle("Histogram of Beer Volume (in Ounces)") + xlim(0,32)</pre>
g.ounces <- g.ounces + theme(plot.title = element_text(hjust = 0.5))</pre>
g.ounces
# sort beers with largest volume in ounces (display top 10)
sortedOunces <- df[order(-df$ounces), ]</pre>
View(head(sortedOunces, 10))
# Look at ounces by Beer Style
plot(x=df$style, y=df$ounces, xlab="Beer Styles", ylab="ounces", cex.axis=0.5, main="Beer Volume
(Ounces)")
# relationship between abv and ibu
volume <- as.factor(df$ounces)</pre>
qplot(data=df, x=abv, y=ibu, color=style, size=volume, main="ABV and IBU by Style") +
theme(plot.title = element text(hjust = 0.5))
# Frequency of beers by state
stateBeers <- barplot(table(df$state), main="Beers by State", las=2)</pre>
```

```
# relationship between style and state
qplot(data=df, x=state, y=style, main="Beer Styles in each State", color=df$style, size=3) +
theme(legend.position="none") + theme(plot.title = element_text(hjust = 0.5))
# map visualization of styles per state
# function from: https://favorableoutcomes.wordpress.com/2012/10/19/create-an-r-function-to-convert-
state-codes-to-full-state-name/
stateFromLower <-function(x) {</pre>
st.codes<-data.frame(
  state=as.factor(c("AK", "AL", "AR", "AZ", "CA", "CO", "CT", "DC", "DE", "FL", "GA",
                             , "IA", "ID", "IL", "IN", "KS", "KY", "LA", "MA", "MD", "ME"
                        "MI", "MN", "MO", "MS", "MT", "NC", "ND", "NE", "NH", "NJ", "NM", "NV", "NY", "OH", "OK", "OR", "PA", "PR", "RI", "SC", "SD", "TN", "TX", "UT", "VA", "VT", "WA", "WI", "WV", "WY")),
    full=as.factor(c("alaska", "alabama", "arkansas", "arizona", "california", "colorado",
                       "connecticut", "district of columbia", "delaware", "florida", "georgia",
                       "hawaii", "iowa", "idaho", "illinois", "indiana", "kansas", "kentucky",
                       "louisiana", "massachusetts", "maryland", "maine", "michigan", "minnesota",
                       "missouri", "mississippi", "montana", "north carolina", "north dakota",
                       "nebraska", "new hampshire", "new jersey", "new mexico", "nevada",
                       "new york", "ohio", "oklahoma", "oregon", "pennsylvania", "puerto rico",
                       "rhode island", "south carolina", "south dakota", "tennessee", "texas",
                       "utah", "virginia", "vermont", "washington", "wisconsin",
                       "west virginia","wyoming"))
  )
  st.x<-data.frame(state=x)</pre>
  refac.x<-st.codes$full[match(st.x$state,st.codes$state)]</pre>
  return(refac.x)
}
states<-map data("state")</pre>
df$state<-gsub("[[:space:]]", "", df$state)</pre>
df$region<-stateFromLower(df$state)</pre>
# text data for maps
counts<-as.data.frame(table(df$state)) # no. of observations per state
colnames(counts)<-c("state.abb","count")</pre>
txt <- data.frame(state.center, state.abb)</pre>
d1<-txt
d2<-counts
lab<-merge(d1,d2, by = "state.abb", all=FALSE)</pre>
rm(counts,txt,d1,d2)
plot.data <- inner_join(states, agg, by = "region")</pre>
df$styles<-as.character(df$style)</pre>
df.new<-within(df,{no.styles<-ave(styles,region,FUN=function(x) length(unique(x)))})</pre>
agg<-subset(df.new,select=c("region","no.styles"))</pre>
agg<-unique(agg)</pre>
agg$no.styles<-as.numeric(paste(agg$no.styles))</pre>
plot.data <- inner_join(states, agg, by = "region")</pre>
ggplot(data = plot.data, mapping = aes(x = long, y = lat, group = group)) +
  coord fixed(1.3) + geom polygon(data = plot.data, aes(fill = no.styles), color = "white") +
  geom polygon(color = "black", fill = NA) +theme bw() +labs( title="Styles Per State") +
  scale fill gradientn("no. of styles",colors=c("pink","blue" )) +
  theme(axis.text = element blank(),
        axis.line = element blank(),
        axis.ticks = element blank(),
        panel.border = element blank(),
```

```
panel.grid = element_blank(),
       axis.title = element_blank())+
 geom_text(data = lab, aes(x = x, y = y, label = count, group = NULL), size = 2)+theme_bw()
# Classification Models
# create data set with only style, abv, and ibu
trainData <- df[,-1] #remove beer name
trainData <- trainData[,-4:-7] #remove ounces, brewery name, city, and state
# hold-out method to measure performance
randIndex <- sample(1:dim(trainData)[1])</pre>
cutpoint2 3 <- floor(2*dim(trainData)[1]/3)</pre>
new_train <- trainData[randIndex[1:cutpoint2_3], ]</pre>
new_validation <- trainData[randIndex[(cutpoint2_3+1):dim(trainData)[1]],]</pre>
train_labels <- new_train$style</pre>
# Decision Tree Modeling (J48 from RWeka)
# J48 decision tree model with default values
model_dt <- J48(style~.,data=trainData)</pre>
model dt
# use 10-fold cross-validation to evaluate model dt
e <- evaluate Weka classifier(model dt, numFolds=10, seed=1, class=TRUE)
e$details
# improve model with increased confidence
model dt2 <- J48(style~.,data=trainData, control=Weka control(U=FALSE, M=2, C=0.5))</pre>
model dt2
# use 10-fold cross-validation to evaulate model dt2
e2 <- evaluate_Weka_classifier(model_dt2, numFolds=10, seed=1, class=TRUE)
e2$details
# examine the effect of pruning by creating unpruned tree
model dt3 <- J48(style~.,data=trainData, control=Weka_control(U=TRUE, M=2))
model dt3
# use 10-fold cross-validation to evaulate model dt3
e3 <- evaluate_Weka_classifier(model_dt3, numFolds=10, seed=1, class=TRUE)
e3$details
# evaluate information gain
ig <- InfoGainAttributeEval(style~.,data=trainData)</pre>
cat("Information Gain:\n\n'abv' 'ibu' \n", ig)
# Naive Bayes Classification Model
# RWeka NB Model with defualt values
NB <- make Weka classifier("weka/classifiers/bayes/NaiveBayes")
model nb <- NB(style~.,data=trainData)</pre>
model nb
```

```
# use 10-fold cross-validation to evaulate the model_nb
e_nb <- evaluate_Weka_classifier(model_nb, numFolds=10, seed=1, class=TRUE)</pre>
NB_results <- e_nb$details
NB_results
# Naive Bayes using e1071
NB_model <- naiveBayes(style~.,data=trainData)</pre>
NB model
NB_pred <- predict(NB_model, trainData)</pre>
table(NB_pred, trainData$style)
NB model2 <- naiveBayes(style~.,data=new train)
NB pred2 <- predict(NB model2, new validation)</pre>
table(NB_pred2, new_validation$style)
# Support Vector Machines Model
# SVM linear model using caret package
set.seed(1818)
train_control <- trainControl(method="cv", number=10)</pre>
fit_svm_linear <- train(style~.,data=new_train,</pre>
                       method="svmLinear",
                       traControl = train_control,
                       na.action=na.pass)
# cross-validation performance
fit_svm_linear
pred_svm_linear <- predict(fit_svm_linear, newdata=new_validation)</pre>
table(pred_svm_linear)
confusionMatrix(pred_svm_linear, new_validation$style)
# SVM RBF model using caret package
fit_svm_rbf <- train(style~.,data=new_train,</pre>
                    method="svmRadial",
                    traControl = train_control,
                     na.action=na.pass)
# cross-validation performance
fit svm rbf
pred_svm_rbf <- predict(fit_svm_rbf, newdata=new_validation)</pre>
table(pred_svm_rbf)
confusionMatrix(pred_svm_rbf, new_validation$style)
```