

# Brewer's Associate of American Independent Study

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## Introduction

According to a recent article in Forbes, MillerCoors (**which owns Blue Moon, Pilsener Urquell and numerous brands besides Miller and Coors products**), ranks No. 2 in volume of beer produced for sale. Constellation Brands is No. 3 with its various brands, including Corona and Modelo.

The traditional giants have bought some craft breweries in recent years, blurring the lines between craft and non-craft brewing companies. Anheuser-Busch bought 10 Barrel Brewing of Bend, Oregon, and Constellation Brands purchased Sand Diego-based Ballast Point for \$1 billion.

This signifies that Craft beers have grown in popularity since the early 2000's and have skyrocketed in 2010 (see Figure 1 below).

In the US alone there are well over 150 styles of beer with as many breweries in each state. Knowing the palate of potential customers can gain a brewery an edge in profit margins by staying close to what the customer wants. Even more important is that the various regions in the USA might actually have varying taste / alcohol content preferences from other regions.

Our client, Brewer's Association of America, wanted to conduct a study for would-be investors to help understand the best chances of product success in today's hyper-competative market.

```
#Load necessary libraries for the project.
```

```
library(ggplot2)
```

```
library(DataExplorer)
```

```
library(dplyr)
```

```
##
```

```
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
## filter, lag
```

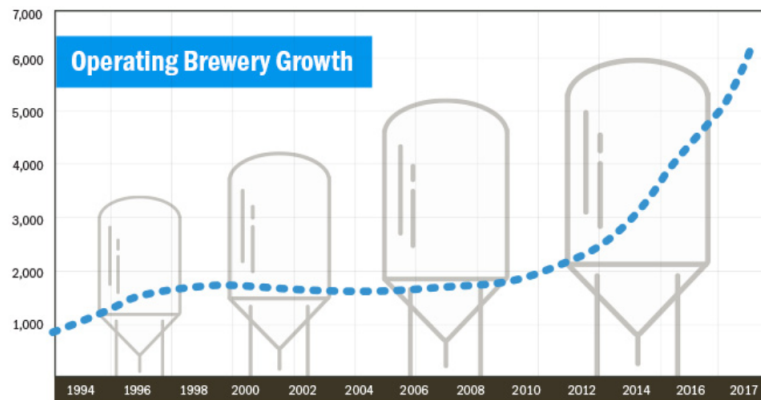


Figure 1:

```
## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
```

## Question 1

### Where and how many breweries are in the US?

The results indicate which states may be over populated with breweries and states that have not seen a wild growth in breweries.

```
#####
#####                               #####
#####      Load Data                #####
#####      Basic details             #####
#####                               #####
#####

#Read CSV file into R
Beers <- read.csv("Beers.csv", header=TRUE, sep=",", strip.white = TRUE)
Breweries <- read.csv("Breweries.csv", header=TRUE, sep=",", strip.white = TRUE)

#####
#####                               #####
#####                               #####
#####      Make State DB             #####
#####                               #####
#####

#Create State DB data frame
StateDB <- data.frame(state.name, state.abb, state.region)
#Rename columns for readability
colnames(StateDB)[colnames(StateDB)=='state.name'] <- 'StateName'
colnames(StateDB)[colnames(StateDB)=='state.abb'] <- 'State'
colnames(StateDB)[colnames(StateDB)=='state.region'] <- 'StateRegion'

#Add district of Columbia to StateDB Data Frame
DistrictColumbia <- data.frame("District of Columbia", "DC", "South")
names(DistrictColumbia) <- c("StateName", "State", "StateRegion")
StateDB <- rbind(StateDB, DistrictColumbia)

#head(StateDB,2) #Assuming this was for debugging purposes.

#####
#####                               #####
#####      Question 1                #####
#####      Breweries per state       #####
#####                               #####
#####

#count of breweries by state
BreweryCounts <- data.frame(table(Breweries$State))

#rename column names
```

```
colnames(BreweryCounts)[colnames(BreweryCounts)=='Var1'] <- 'State'
colnames(BreweryCounts)[colnames(BreweryCounts)=='Freq'] <- 'NumberOfBreweriesByState'

#Merge the StateDB and sort by count of breweries by state
BreweryCounts <- merge(BreweryCounts, StateDB, by.x="State", by.y="State")
BreweryCounts <- BreweryCounts[order(BreweryCounts$NumberOfBreweriesByState, decreasing=TRUE),c(3,2)]
BreweryCounts
```

##	StateName	NumberOfBreweriesByState
## 6	Colorado	47
## 5	California	39
## 23	Michigan	32
## 38	Oregon	29
## 44	Texas	28
## 39	Pennsylvania	25
## 20	Massachusetts	23
## 48	Washington	23
## 16	Indiana	22
## 49	Wisconsin	20
## 28	North Carolina	19
## 15	Illinois	18
## 35	New York	16
## 46	Virginia	16
## 10	Florida	15
## 36	Ohio	15
## 24	Minnesota	12
## 4	Arizona	11
## 47	Vermont	10
## 22	Maine	9
## 25	Missouri	9
## 27	Montana	9
## 7	Connecticut	8
## 1	Alaska	7
## 11	Georgia	7
## 21	Maryland	7
## 37	Oklahoma	6
## 13	Iowa	5
## 14	Idaho	5
## 19	Louisiana	5
## 30	Nebraska	5
## 40	Rhode Island	5
## 12	Hawaii	4
## 18	Kentucky	4
## 33	New Mexico	4
## 41	South Carolina	4
## 45	Utah	4
## 51	Wyoming	4
## 2	Alabama	3
## 17	Kansas	3
## 31	New Hampshire	3
## 32	New Jersey	3
## 43	Tennessee	3
## 3	Arkansas	2
## 9	Delaware	2

## 26	Mississippi	2
## 34	Nevada	2
## 8	District of Columbia	1
## 29	North Dakota	1
## 42	South Dakota	1
## 50	West Virginia	1

## Question 2

### Merging Data

The team merged both the beer and breweries data sets in order to get a better wholelistic view of the data and determine how they relate to each other. Below will show the first and last six rows of the data sets.

```
#####
#####
#####      Question 2      #####
#####      Merge Data Sets  #####
#####
#####

#join data on Brewery_id and Brew_ID
BeersAndBreweries <- merge(Beers, Breweries, by.x="Brewery_id", by.y="Brew_ID")

#list column names on the joined data frame
#colnames(BeersAndBreweries) #assuming this was for debugging purposes

#rename the name.x(Beer) and name.y(Brewery) after the merger
colnames(BeersAndBreweries)[colnames(BeersAndBreweries)=='Name.x'] <- 'BeerName'
colnames(BeersAndBreweries)[colnames(BeersAndBreweries)=='Name.y'] <- 'BreweryName'

# I don't think we don't need to create this again.
# #Create State DB data frame
# StateDB <- data.frame(state.name, state.abb, state.region)
# colnames(StateDB)[colnames(StateDB)=='state.name'] <- 'StateName'
# colnames(StateDB)[colnames(StateDB)=='state.abb'] <- 'State'
# colnames(StateDB)[colnames(StateDB)=='state.region'] <- 'StateRegion'
#
# #Add district of Columbia to StateDB Data Frame
# DistrictColumbia <- data.frame("District of Columbia","DC", "South")
# names(DistrictColumbia) <- c("StateName","State", "StateRegion")
# StateDB <- rbind(StateDB, DistrictColumbia)

#Merge data with State DB
BeersAndBreweries <- merge(BeersAndBreweries, StateDB, by="State", all = TRUE)

# dont think we need to show this.
# #find dimensions of data frames
# dim(BeersAndBreweries)
# dim(BeersAndBreweries)
```

```
#Show first and last 6 entries of merged files
head(BeersAndBreweries)
```

```
##      State Brewery_id      BeerName Beer_ID  ABV IBU
## 1      AK          103      King Street IPA    1667 0.060  70
## 2      AK          103          Amber Ale    2436 0.051  NA
## 3      AK          494      Polar Pale Ale    920 0.052  17
## 4      AK          459      Sunken Island IPA    349 0.068  NA
## 5      AK          103      King Street Pilsner    1706 0.055  NA
## 6      AK          459 Skilak Scottish Ale (2011)    348 0.058  NA
##
##              Style Ounces      BreweryName      City
## 1      American IPA      12 King Street Brewing Company Anchorage
## 2 American Amber / Red Ale      12 King Street Brewing Company Anchorage
## 3 American Pale Ale (APA)      12 Broken Tooth Brewing Company Anchorage
## 4      American IPA      12 Kenai River Brewing Company Soldotna
## 5      Czech Pilsener      12 King Street Brewing Company Anchorage
## 6      Scottish Ale      12 Kenai River Brewing Company Soldotna
##      StateName StateRegion
## 1      Alaska      West
## 2      Alaska      West
## 3      Alaska      West
## 4      Alaska      West
## 5      Alaska      West
## 6      Alaska      West
```

```
tail(BeersAndBreweries)
```

```
##      State Brewery_id      BeerName Beer_ID  ABV IBU
## 2405      WY          458 Saddle Bronc Brown Ale (2013)    1198 0.048  16
## 2406      WY          192      Pako's EyePA    393 0.068  60
## 2407      WY          192      Snow King Pale Ale    1606 0.060  55
## 2408      WY          458      Wagon Box Wheat Beer    1197 0.059  15
## 2409      WY          458      Indian Paintbrush IPA    1199 0.070  75
## 2410      WY          458 Bomber Mountain Amber Ale (2013)    1200 0.046  20
##
##              Style Ounces      BreweryName
## 2405      English Brown Ale      12 The Black Tooth Brewing Company
## 2406      American IPA      12 Snake River Brewing Company
## 2407 American Pale Ale (APA)      12 Snake River Brewing Company
## 2408 American Pale Wheat Ale      12 The Black Tooth Brewing Company
## 2409      American IPA      12 The Black Tooth Brewing Company
## 2410 American Amber / Red Ale      12 The Black Tooth Brewing Company
##
##      City StateName StateRegion
## 2405 Sheridan Wyoming      West
## 2406 Jackson Wyoming      West
## 2407 Jackson Wyoming      West
## 2408 Sheridan Wyoming      West
## 2409 Sheridan Wyoming      West
## 2410 Sheridan Wyoming      West
```

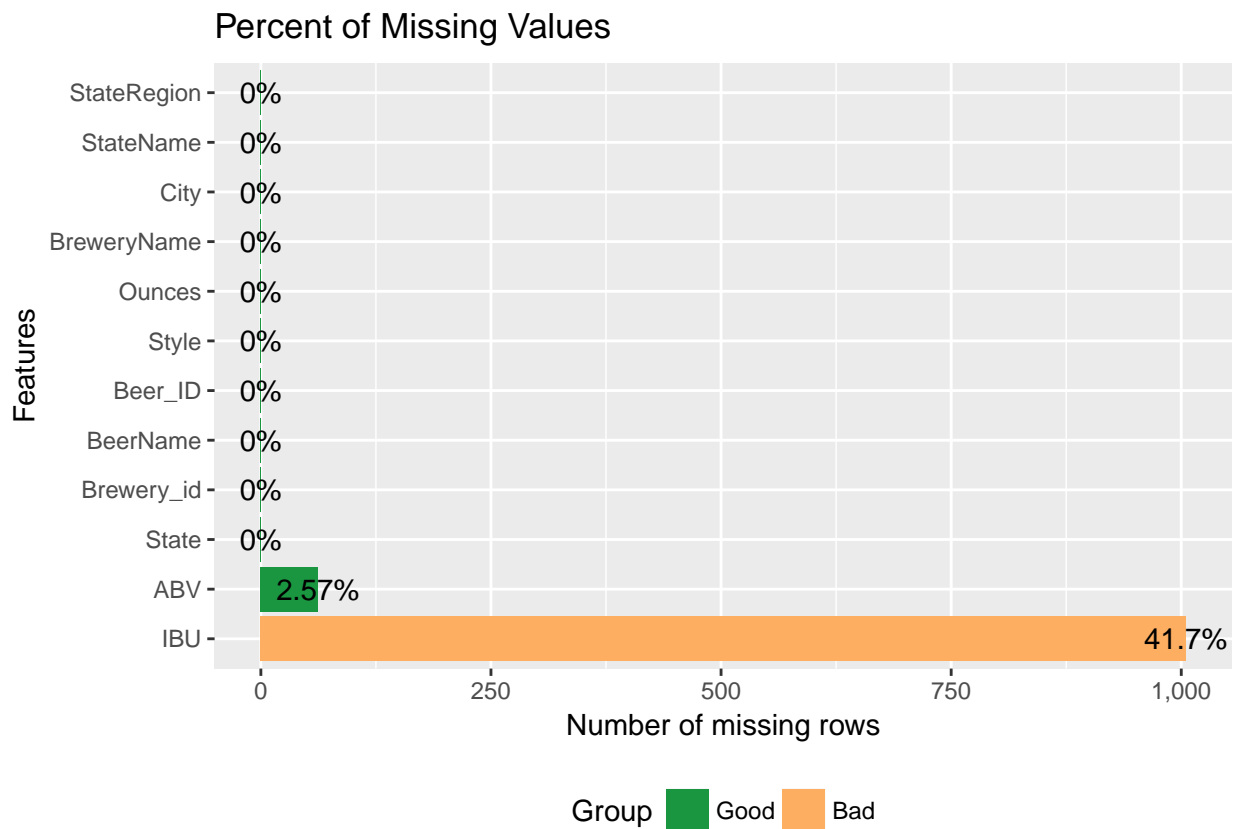
## Question 3

### Report Missing Values

To do a complete analysis the team needed to assess if the merged data set has any missing values. The team created a report for both a graphical and tabular representation of the results. The team utilized code from <http://www.gettinggeneticsdone.com/2011/02/summarize-missing-data-for-all.html> to count NA's in columns.

The data shows the variables with missing data are the IBU and ABV values.

```
#Graphical representation of missing values using 'DataExplorer' library
plot_missing(BeersAndBreweries, title = "Percent of Missing Values")
```



```
#Function to count all NA's in columns (sourced from the internet)
#http://www.gettinggeneticsdone.com/2011/02/summarize-missing-data-for-all.html
propmiss <- function(dataframe) {
  m <- sapply(dataframe, function(x) {
    data.frame(
      na_count=sum(is.na(x)),
      Obs=length(x),
      perc_missing=sum(is.na(x))/length(x)*100
    )
  })
  d <- data.frame(t(m))
  d <- sapply(d, unlist)
  d <- as.data.frame(d)
  d$variable <- row.names(d)
```

```

row.names(d) <- NULL
d <- cbind(d[ncol(d)], d[-ncol(d)])
return(d[order(d$na_count, decreasing=TRUE), ])
}

#show results of NA's counted
BeerColumnInventory_nacount <- propmiss(BeersAndBreweries)
BeerColumnInventory_nacount

```

```

##      variable na_count  Obs perc_missing
## 6          IBU    1005 2410    41.701245
## 5          ABV     62 2410    2.572614
## 1         State      0 2410    0.000000
## 2  Brewery_id      0 2410    0.000000
## 3    BeerName      0 2410    0.000000
## 4    Beer_ID      0 2410    0.000000
## 7         Style      0 2410    0.000000
## 8        Ounces      0 2410    0.000000
## 9  BreweryName      0 2410    0.000000
## 10         City      0 2410    0.000000
## 11   StateName      0 2410    0.000000
## 12 StateRegion      0 2410    0.000000

```

## Question 4

### Plotting Data

The team in order to look for trends plotted the ABV and IBU against the states to determine which states had the highest median value of each of the states by value.

- Process to analyze
  - Calculate the median values of ABV & IBU by state
  - Plot the data against states and sort by highest value

*As requested by the client all NA's have been removed*

```

#Make data frame with only State, ABV, IBU
DF_ABV_IBU <- BeersAndBreweries[,c("StateName", "ABV", "IBU")]
#head(DF_ABV_IBU) #I don't think we don't need to show this.

#remove any rows with a NA value using 'complete.cases'
DF_ABV_IBU_noNA <- DF_ABV_IBU[complete.cases(DF_ABV_IBU),]
#head(DF_ABV_IBU_noNA) #I don't think we don't need to show this.

#Calculate MEDIAN values for ABV&IBU by State
MEDIAN_ABV_IBU_by_State <- aggregate(DF_ABV_IBU_noNA[, 2:3], list(DF_ABV_IBU_noNA$StateName), median)
#head(MEDIAN_ABV_IBU_by_State) #I don't think we don't need to show this.

#Rename column names
colnames(MEDIAN_ABV_IBU_by_State)[colnames(MEDIAN_ABV_IBU_by_State)=='Group.1'] <- 'State'
colnames(MEDIAN_ABV_IBU_by_State)[colnames(MEDIAN_ABV_IBU_by_State)=='ABV'] <- 'Median_ABV'
colnames(MEDIAN_ABV_IBU_by_State)[colnames(MEDIAN_ABV_IBU_by_State)=='IBU'] <- 'Median_IBU'

```

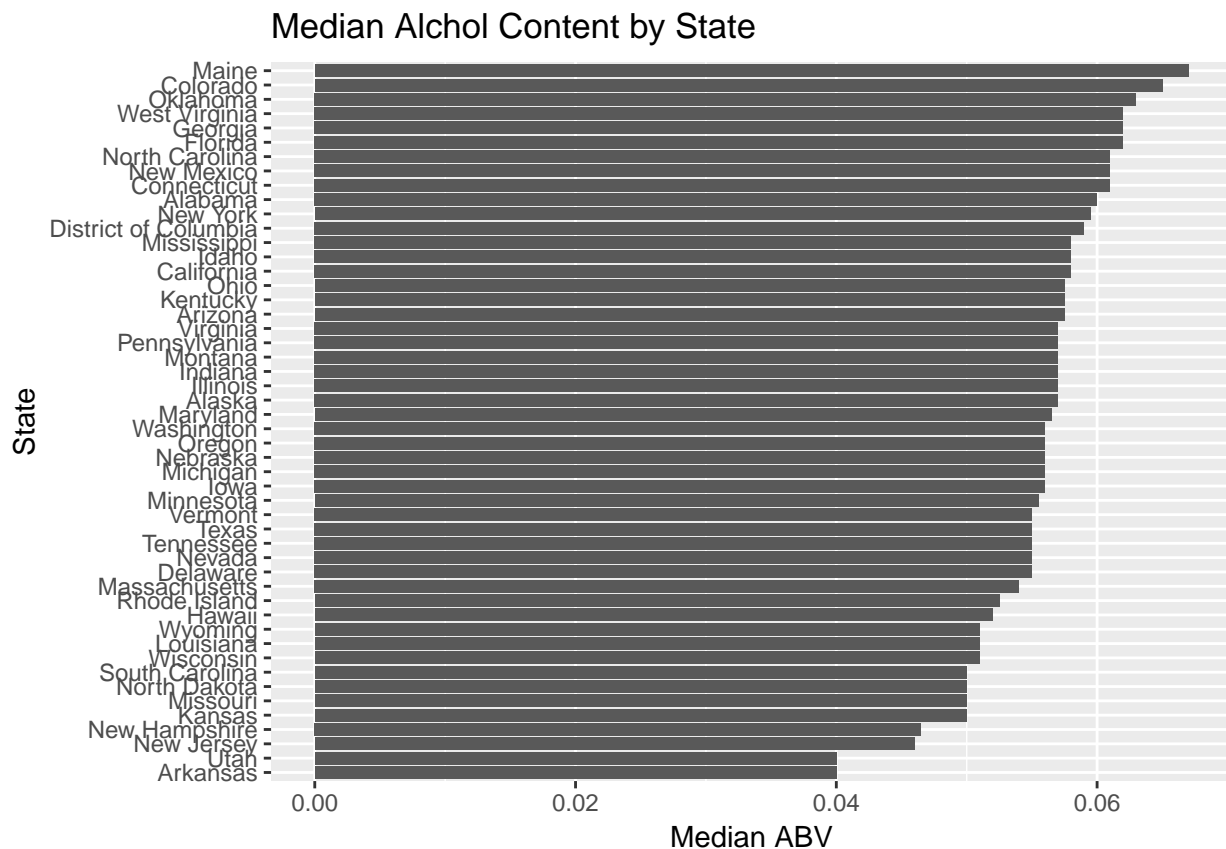
```

#Check data
#head(MEDIAN_ABV_IBU_by_State,10) #I don't think we don't need to show this.

##### Plot MEDIAN ABV By State #####
BarPlot_ABV_byState <- ggplot(data=MEDIAN_ABV_IBU_by_State,
                              aes(x=reorder(State, Median_ABV),
                                  y=Median_ABV)) +
  geom_bar(stat="identity")+
  coord_flip() +
  labs(x="State",
       y="Median ABV",
       title = "Median Alchol Content by State")

BarPlot_ABV_byState

```



```

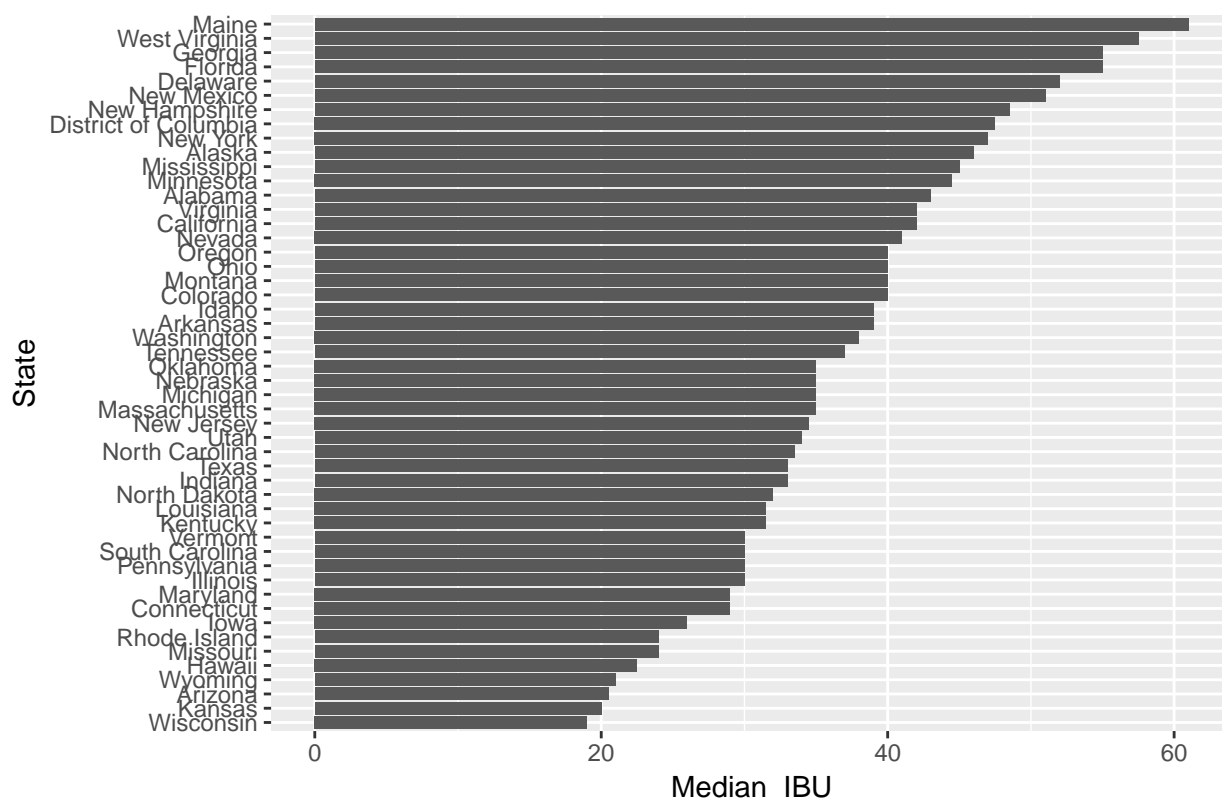
##### Plot MEDIAN IBU By State #####
BarPlot_IBU_byState <- ggplot(data=MEDIAN_ABV_IBU_by_State,
                              aes(x=reorder(State, Median_IBU),
                                  y=Median_IBU)) +
  geom_bar(stat="identity")+
  coord_flip()+
  labs(x="State",
       y="Median IBU",
       title = "Median Bitterness Content by State")

BarPlot_IBU_byState

```



Median Bitterness Content by State



## Question 5

### States with highest ABV and IBU

For a quick reference the team identified the states with the highest ABV and IBU recorded within the data set.

*#Find MAX ABV with State*

```
MAX_ABV_byState <- head(BeersAndBreweries[order(BeersAndBreweries$ABV, na.last = TRUE, decreasing=TRUE)], 1)
MAX_ABV_byState
```

```
##      StateName  ABV
## 533   Colorado 0.128
```

*#Find MAX IBU with State, column has missing values*

```
MAX_IBU_byState <- head(BeersAndBreweries[order(BeersAndBreweries$IBU, na.last = TRUE, decreasing=TRUE)], 1)
MAX_IBU_byState
```

```
##      StateName IBU
## 1824    Oregon 138
```

## Question 6

### Summary of ABV

As part of the analysis the team has provided the summary results for the ABV variable.

```
#####
#####
#####      Question 6      #####
##### Summary of ABV Variable #####
#####
#####

#Summary Stats of the ABV variable
SUMMARY_ABV <- summary(BeersAndBreweries$ABV)

#Show ABV Summary
SUMMARY_ABV

##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.     NA's
## 0.00100 0.05000 0.05600 0.05977 0.06700 0.12800     62
```

## Question 7

### Relationship between ABV & IBU

To determine any relationships between ABV and IBU a scatterplot was created. In addition the data was color coded by the region of the brewery.

```
#Merge data with State DB
DF_ABV_IBU_noNA <- merge(DF_ABV_IBU_noNA, StateDB, by="StateName", all = TRUE)

#Remove rows with NA's
DF_ABV_IBU_noNA <- DF_ABV_IBU_noNA[complete.cases(DF_ABV_IBU_noNA),]

# I don't think we need to show this.
#Check merge and top of the file
# propmiss(DF_ABV_IBU_noNA)
# head(DF_ABV_IBU_noNA)

#Scatter plot ABV vs IBU and color by StateRegion
ABUvsIBU <- qplot(ABV, IBU,
                  xlab = "ABV (Alcohol Content)",
                  ylab = "IBU (Bitterness)",
                  main= "ABV vs IBU",
                  colour=StateRegion,
                  data=DF_ABV_IBU_noNA)

#Show Scatter Plot
ABUvsIBU
```

