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title: "Case Study 1: Beers and Breweries""
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date: 01/24/2020
'''{r}
library(tidyverse)
library(ggplot2)
library(plotly)
library(forcats)
library(dplyr)
library(class)
library(caret)
library(maps)
library(mapproj)
#Reading Beers and Brewers files.
Beers = read.csv(file.choose(),header = TRUE)
Breweries=read.csv(file.choose(),header = TRUE)
count(Beers)
count(Breweries)
head(Beers)
head(Breweries)
str(Breweries)
#Q1. Count number of brewer in each State.
Breweries %>% group by(State) %>% summarize(count=n())
#Plot the number of brewer in each State.
Breweries %>% ggplot(aes(x=forcats::fct infreq(State,ordered = TRUE)))
+geom bar()+ggtitle("Number of Breweries in each State")+xlab("State")
#text(x=p, y=Breweries$State,labels =
Breweries$State,pos=3,cex=0.6,col="red")
#Q2. Mergering Beer and Breweries.
BB=merge(Beers, Breweries, by.x="Brewery_id", by.y = "Brew_ID")
#rename column name
colnames(BB)[2]<-"Beer Name"</pre>
colnames(BB)[8] <-"Brewery_Name"</pre>
head(BB, n=6)
tail(BB, n=6)
#Q3&4. Plot the median of ABV by grouping them on each State
BB %>% group by(State) %>% summarize(median=median(ABV,na.rm=TRUE))%>%
ggplot(aes(x=State,y=(median)))+geom bar(stat = "identity")+ggtitle("Median
Alcohol Content by State")
#Plot the median of IBU by grouping them on each State
BB %>% group_by(State) %>% summarize(median=median(IBU,na.rm=TRUE))%>%
ggplot(aes(x=State,y=median))+geom bar(stat = "identity")
+ggtitle("International Bitterness by State")
#Q5. Plot the max of IBU by grouping them on each State
BB %>% group_by(State) %>% summarize(max=max(ABV,na.rm=TRUE)) %>%
ggplot(aes(x=State,y=max))+geom_bar(stat = "identity")+ggtitle("Max ABV by
State")
#Plot the max of IBU by grouping them on each State
BB %>% group by(State) %>% summarize(max=max(IBU,na.rm=TRUE)) %>%
ggplot(aes(x=State,y=max))+geom_bar(stat = "identity")+ggtitle("Max IBU by
State")
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#Q6. Summary Statistics on IBU and ABV
SummaryStat= BB %>% group_by(State) %>%
summarize(mean=mean(ABV,na.rm=TRUE),sd=sd(ABV,na.rm=TRUE),range=max(ABV,na.rm=TRUE)-
min(ABV, na.rm=TRUE), IQR=IQR(ABV, na.rm=TRUE), count=n())
print(SummaryStat, n=51)
SummaryStat%>%arrange(desc(sd))%>% print(n=51)
#Q7. Correlation between IBU and ABV
install.packages("GGally")
library(GGally)
BB %>% select(ABV,IBU) %>% ggpairs()
#Q8. IPA vs Ale in IBU and ABV
#BBIPA= BB %>% filter(!is.na(IBU)) %>% filter(str detect(Style, "IPA")) %>%
select(Style, IBU, ABV)
#head(BBIPA)
BBAleIPA=BB %>% filter(!is.na(IBU)) %>% filter(str detect(Style, "Ale")|
str detect(Style,"IPA")) %>% select(Style, IBU, ABV, State)
head(BBAleIPA)
dummy <-str sub(BBAleIPA$Style,-3,-1)</pre>
BBAleIPA$Style <- ifelse(dummy == "Ale", "Ale", "IPA")
# Q8. New finding that will blow their socks off
colnames(BBAleIPA)[4]="abb"
head(BBAleIPA)
str view all(BBAleIPA$abb,"\\s")
trimws(BBAleIPA$abb, which=c("left"))
str(lookup)
str(BBAleIPA)
lookup = data.frame(abb = state.abb, State = state.name)
head(lookup)
BBAleIPA$abb=trimws(BBAleIPA$abb, which=c("left"))
BBAleIPA$abb <- as.factor(BBAleIPA$abb)</pre>
str(BBAleIPA)
BBAleIPA2= merge(BBAleIPA,lookup,by="abb")
head(BBAleIPA2)
BBAleIPACount=count(BBAleIPA2,State, sort = TRUE)
head(BBAleIPACount)
BBAleIPACount$region <- tolower(BBAleIPACount$State)</pre>
BBAleIPACount2= BBAleIPACount[-1]
states <- map data("state")</pre>
map.df <-merge(states, BBAleIPACount2, by="region", all.x=T)</pre>
map.df <-map.df[order(map.df$order),]</pre>
ggplot(map.df,aes(x=long,y=lat,group=group))+geom_polygon(aes(fill=n))
+geom path()
+scale fill gradientn(colours=rev(heat.colors(10)),na.value="grey90")
+coord_map()+ggtitle("Number of IPA and Ale in the State")
BBAleIPA$Style <- as.factor(BBAleIPA$Style)
str(BBAleIPA)
BBAleIPA %>% ggplot(aes(x=IBU,ABV,color=Style))+geom_point()+ggtitle(" IBU
vs ABV between IPA and Ale")
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summary(BBAleIPA)
trainIndices=sample(seg(1:944),round(0.7*944))
trainBeers=BBAleIPA[trainIndices,]
testBeers=BBAleIPA[-trainIndices,]
classifications= knn(trainBeers[,c(2,3)],testBeers[,c(2,3)],
(trainBeers$Style),prob=TRUE,k=13)
table(classifications,testBeers$Style)
confusionMatrix(table(classifications, testBeers$Style))
#naive Bayes
iterations = 100
masterAcc = matrix(nrow = iterations)
splitPerc = .7 #Training / Test split Percentage
for(j in 1:iterations)
 trainIndices=sample(seg(1:944),round(0.7*944))
 train=BBAleIPA[trainIndices,]
 test=BBAleIPA[-trainIndices,]
 model = naiveBayes(train[,c(2,3)],train$Style)
 table(predict(model, test[,c(2,3)]), test$Style)
 CM = confusionMatrix(table(predict(model, test[,c(2,3)]), test$Style))
 masterAcc[j] = CM$overall[1]
}
MeanAcc = colMeans(masterAcc)
MeanAcc
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