```
library(caret)
library(gbm)
data("scat")
preProcValues <- preProcess(scat, method = c("knnImpute","center","scale"))</pre>
library('RANN')
train_processed <- predict(preProcValues, scat)</pre>
unique(train_processed$Species)
#Question 1
train_processed$Species<-ifelse(scat$Species=='coyote',0,ifelse(scat$Species=='bobcat',1,2))
#Question 2
#https://www.listendata.com/2015/06/r-keep-drop-columns-from-data-frame.html
train processed = subset(train processed, select = -c(Month, Year, Site, Location))
#Question 3
sum(is.na(train processed))
#No null values
#Question 4
dmy <- dummyVars(" ~ .", data = train_processed,fullRank = T)</pre>
train_transformed <- data.frame(predict(dmy, newdata = train_processed))</pre>
#Question 5
train_transformed$Species<-as.factor(train_transformed$Species)</pre>
set.seed(100)
index <- createDataPartition(train_transformed$Species, p=0.75, list=FALSE)
trainSet <- train_transformed[ index,]</pre>
testSet <- train_transformed[-index,]</pre>
str(trainSet)
```

```
#Feature selection using rfe in caret
control <- rfeControl(functions = rfFuncs,
             method = "repeatedcv",
             repeats = 3,
             verbose = FALSE)
outcomeName<-'Species'
predictors<-names(trainSet)[!names(trainSet) %in% outcomeName]</pre>
Species_Pred_Profile <- rfe(trainSet[,predictors], trainSet[,outcomeName],rfeControl = control)
Species_Pred_Profile
#Taking only the top 4 predictors
predictors <- c("d15N", "d13C", "Mass", "CN")
names(getModelInfo())
model_gbm<-train(trainSet[,predictors],trainSet[,outcomeName],method='gbm')</pre>
model_rf<-train(trainSet[,predictors],trainSet[,outcomeName],method='rf')</pre>
model_nnet<-train(trainSet[,predictors],trainSet[,outcomeName],method='nnet')</pre>
model NB <- train(trainSet[,predictors],trainSet[,outcomeName],method='nb')
fitControl <- trainControl(</pre>
 method = "repeatedcv",
 number = 5,
 repeats = 5)
### Using tuneGrid ####
modelLookup(model='gbm')
#Creating grid
grid <-
expand.grid(n.trees=c(10,20,50,100,500,1000),shrinkage=c(0.01,0.05,0.1,0.5),n.minobsinnode
= c(3,5,10), interaction.depth = c(1,5,10)
# training the model
model_gbm<-train(trainSet[,predictors],trainSet[,outcomeName],method='gbm',trControl=fitCont
rol,tuneGrid=grid)
# summarizing the model
print(model_gbm)
# Visualizing the models
```

```
plot(model_gbm)
### Using tuneLength ###
#using tune length
model_gbm<-train(trainSet[,predictors],trainSet[,outcomeName],method='gbm',trControl=fitCont
rol,tuneLength=10)
print(model_gbm)
# visualize the models
plot(model gbm)
#Checking variable importance for GBM
#Variable Importance
varImp(object=model_gbm)
#Plotting Varianle importance for GBM
plot(varImp(object=model_gbm),main="GBM - Variable Importance")
#Checking variable importance for RF
varImp(object=model_rf)
#Plotting Varianle importance for Random Forest
plot(varImp(object=model rf),main="RF - Variable Importance")
#Checking variable importance for NNET
varImp(object=model nnet)
#nnet variable importance
#Plotting Variable importance for Neural Network
plot(varImp(object=model_nnet),main="NNET - Variable Importance")
```