

```
cat("\f") # Clear the console
```

```
rm(list = ls()) # Clear the working environment
```

```
##### Load the required packages  
#####
```

```
library(ISLR)
```

```
library(e1071)
```

```
library(pROC)
```

```
library(dplyr)
```

```
##### Load the data  
#####
```

```
iris <- iris
```

```
glimpse(iris) # Metadata
```

```
str(iris)
```

```
head(iris) # See the first 6 values of the dataset
```

```
sum(is.na(iris)) # Check for NA's
```

```
attach(iris) #Optional Step
```

```
##### Subset the data  
#####
```

```
Random.seed <- c('Mersenne-Twister', 490)
```

```
set.seed(490) # Set seed for replication
```

```
index <- sample(1:nrow(iris), 0.5 * nrow(iris), replace = FALSE) # Create an index to split the data
```

```
# iris_train <- iris[index,]
```

```
# iris_test <- iris[-index,]
```

```
iris_train_x <- iris[index, 1:4]
```

```
iris_test_x <- iris[-index, 1:4]
```

```
iris_train_y <- iris[index, 5]
```

```
iris_test_y <- iris[-index, 5]
```

**#### \*\*\*\*\* For the TA: I need to understand why we chose to subset the dataset like we did. I have never**

**#### subsetted data like this and am struggling to see the necessity or requirement to do the split**

**#### like it is done in the lines. Is it done to adhere to the arguments for naiveBayes() only? \*\*\*\*\***

```
# table(Species)
```

```
model <- naiveBayes(iris_train_x, iris_train_y) # Apply the Naive Bayes Estimator
```

```
## Predictions for Training set
```

```
pred_train <- predict(model, iris_train_x, type = "class")
```

```
cfm_train <- table(pred_train, iris_train_y)
```

```
cfm_train
```

```
## Predictions for Testing set
```

```
pred_test <- predict(model, iris_test_x, type = "class")
```

```
cfm_test <- table(pred_test, iris_test_y)
```

```
cfm_test
```

```
## Predictions for the entire set
```

```
pred_all <- predict(model, iris[1:4], type = "class")
```

```
cfm_all <- table(pred_all, Species)
```

```
cfm_all
```

```
accuracy_train <- sum(diag(cfm_train))/sum(cfm_train) # Training Accuracy
```

```
accuracy_train
```

```
error_train <- 1 - accuracy_train # Training Error
```

```
error_train
```

```
accuracy_test <- sum(diag(cfm_test))/sum(cfm_test) # Testing Accuracy
```

```
accuracy_test
```

```
error_test <- 1 - accuracy_test # Testing Error
```

```
error_test
```

```
accuracy_all <- sum(diag(cfm_all))/sum(cfm_all) # Overall Error
```

```
accuracy_all
```

```
error_all <- 1 - accuracy_all # Overall Accuracy
```

```
error_all
```

```
##### Calculate Sensitivity, Specificity, PPV & NPV
```

```
#####
```

```
##### TRAINING SET
```

```
#####
```

```
##### SETOSA
```

```
#####
```

```
macro_spec <- 0
```

```
macro_sens <- 0
```

```
train_sens <- cfm_train["setosa", "setosa"] / sum(cfm_train[, 'setosa'])
```

```
train_sens
```

```
train_spec <- sum(diag(cfm_train[2:3, 2:3])) / (sum(diag(cfm_train[2:3, 2:3])) +  
sum(cfm_train['setosa', 2:3]))
```

```
train_spec
```

```
ppv_train <- cfm_train['setosa', 'setosa'] / sum(cfm_train['setosa',])
```

```
ppv_train
```

```
npv_train <- sum(diag(cfm_train[2:3, 2:3])) / (sum(diag(cfm_train[2:3, 2:3])) +  
sum(cfm_train[2:3, 'setosa']))
```

```
npv_train
```

```
macro_sens <- macro_sens + train_sens
```

```
macro_spec <- macro_spec + train_spec
```

```
##### VERSICOLOR
```

```
#####
```

```
train_sens <- cfm_train["versicolor", "versicolor"] / sum(cfm_train[, 'versicolor'])
```

```
train_sens
```

```
train_spec <- sum(diag(cfm_train[c(1, 3), c(1, 3)])) / (sum(diag(cfm_train[c(1, 3), c(1, 3)])) +  
sum(cfm_train['versicolor', c(1, 3)]))
```

```
train_spec
```

```
ppv_train <- cfm_train['versicolor','versicolor']/sum(cfm_train['versicolor',])
```

```
ppv_train
```

```
npv_train <- sum(diag(cfm_train[c(1,3),c(1,3)]))/(sum(diag(cfm_train[c(1,3),c(1,3)])) +  
sum(cfm_train[c(1,3),'versicolor']))
```

```
npv_train
```

```
macro_sens <- macro_sens + train_sens
```

```
macro_spec <- macro_spec + train_spec
```

```
##### VIRGINICA  
#####
```

```
train_sens <- cfm_train["virginica","virginica"]/sum(cfm_train[, 'virginica'])
```

```
train_sens
```

```
train_spec <- sum(diag(cfm_train[1:2,1:2]))/(sum(diag(cfm_train[1:2,1:2])) +  
sum(cfm_train['virginica',1:2]))
```

```
train_spec
```

```
ppv_train <- cfm_train['virginica','virginica']/sum(cfm_train['virginica',])
```

```
ppv_train
```

```
npv_train <- sum(diag(cfm_train[1:2,1:2]))/(sum(diag(cfm_train[1:2,1:2])) +  
sum(cfm_train[1:2,'virginica']))
```

```
npv_train
```

```
macro_sens <- macro_sens + train_sens
```

```
macro_spec <- macro_spec + train_spec
```

```
macro_sens<- macro_sens/3
```

```
macro_spec<- macro_spec/3
```

```
macro_sens
```

```
macro_spec
```

```
##### TESTING SET
```

```
#####
```

```
##### SETOSA
```

```
#####
```

```
macro_spec<- 0
```

```
macro_sens<- 0
```

```
test_sens<- cfm_test["setosa","setosa"]/sum(cfm_test[, 'setosa'])
```

```
test_sens
```

```
test_spec<- sum(diag(cfm_test[2:3,2:3]))/(sum(diag(cfm_test[2:3,2:3])) + sum(cfm_test['setosa',2:3]))
```

```
test_spec
```

```
ppv_test<- cfm_test['setosa','setosa']/sum(cfm_test['setosa',])
```

```
ppv_test
```

```
npv_test<- sum(diag(cfm_test[2:3,2:3]))/(sum(diag(cfm_test[2:3,2:3])) + sum(cfm_test[2:3,'setosa']))
```

```
npv_test
```

```
macro_sens<- macro_sens + test_sens
```

```
macro_spec<- macro_spec+ test_spec
```

```
##### VERSICOLOR  
#####
```

```
test_sens<- cfm_test["versicolor","versicolor"]/sum(cfm_test[, 'versicolor'])
```

```
test_sens
```

```
test_spec<- sum(diag(cfm_test[c(1,3),c(1,3)]))/(sum(diag(cfm_test[c(1,3),c(1,3)])) +  
sum(cfm_test['versicolor',c(1,3)]))
```

```
test_spec
```

```
ppv_test<- cfm_test['versicolor','versicolor']/sum(cfm_test['versicolor',])
```

```
ppv_test
```

```
npv_test<- sum(diag(cfm_test[c(1,3),c(1,3)]))/(sum(diag(cfm_test[c(1,3),c(1,3)])) +  
sum(cfm_test[c(1,3), 'versicolor']))
```

```
npv_test
```

```
macro_sens<- macro_sens+ train_sens
```

```
macro_spec<- macro_spec+ train_spec
```

```
##### VIRGINICA  
#####
```

```
test_sens<- cfm_test["virginica","virginica"]/sum(cfm_test[, 'virginica'])
```

```
test_sens
```

```
test_spec<- sum(diag(cfm_test[1:2,1:2]))/(sum(diag(cfm_test[1:2,1:2])) + sum(cfm_test['virginica',1:2]))
test_spec
```

```
ppv_test<- cfm_test['virginica','virginica']/sum(cfm_test['virginica',])
ppv_test
```

```
npv_test<- sum(diag(cfm_test[1:2,1:2]))/(sum(diag(cfm_test[1:2,1:2])) + sum(cfm_test[1:2,'virginica']))
npv_test
```

```
macro_sens<- macro_sens + train_sens
macro_spec<- macro_spec + train_spec
```

```
macro_sens<- macro_sens/3
macro_spec<- macro_spec/3
macro_sens
macro_spec
```

```
##### FULL DATA SET
#####
```

```
##### SETOSA
#####
```

```
macro_spec<- 0
macro_sens<- 0
```

```
sens<- cfm_all["setosa","setosa"]/sum(cfm_all[, 'setosa'])
sens
```



```
spec<- sum(diag(cfm_all[2:3,2:3]))/(sum(diag(cfm_all[2:3,2:3])) + sum(cfm_all['setosa',2:3]))
```

```
spec
```

```
ppv<- cfm_all['setosa','setosa']/sum(cfm_all['setosa',])
```

```
ppv
```

```
npv<- sum(diag(cfm_all[2:3,2:3]))/(sum(diag(cfm_all[2:3,2:3])) + sum(cfm_all[2:3,'setosa']))
```

```
npv
```

```
macro_sens<- macro_sens + sens
```

```
macro_spec<- macro_spec + spec
```

```
##### VERSICOLOR
```

```
#####
```

```
sens<- cfm_all["versicolor","versicolor"]/sum(cfm_all[, 'versicolor'])
```

```
sens
```

```
spec<- sum(diag(cfm_all[c(1,3),c(1,3)]))/(sum(diag(cfm_all[c(1,3),c(1,3)])) +  
sum(cfm_all['versicolor',c(1,3)]))
```

```
spec
```

```
ppv<- cfm_all['versicolor','versicolor']/sum(cfm_all['versicolor',])
```

```
ppv
```

```
npv<- sum(diag(cfm_all[c(1,3),c(1,3)]))/(sum(diag(cfm_all[c(1,3),c(1,3)])) +  
sum(cfm_all[c(1,3), 'versicolor']))
```

```
npv
```

```
macro_sens<- macro_sens + sens
```

```
macro_spec<- macro_spec + spec
```

```
##### VIRGINICA
```

```
#####
```

```
sens<- cfm_all["virginica","virginica"]/sum(cfm_all[, 'virginica'])
```

```
sens
```

```
spec<- sum(diag(cfm_all[1:2,1:2]))/(sum(diag(cfm_all[1:2,1:2])) + sum(cfm_all['virginica',1:2]))
```

```
spec
```

```
ppv<- cfm_all['virginica','virginica']/sum(cfm_all['virginica',])
```

```
ppv
```

```
npv<- sum(diag(cfm_all[1:2,1:2]))/(sum(diag(cfm_all[1:2,1:2])) + sum(cfm_all[1:2,'virginica']))
```

```
npv
```

```
macro_sens<- macro_sens + sens
```

```
macro_spec<- macro_spec + spec
```

```
macro_sens<- macro_sens/3
```

```
macro_spec<- macro_spec/3
```

```
macro_sens
```

```
macro_spec
```

```
##### ROC CURVE
```

```
#####
```

```
prob <- predict(model, iris_test_x, type = "raw")
```

```
setosa_labels <- rep(0, length(iris_test_y))
```

```
versicolor_labels <- rep(0, length(iris_test_y))
```

```
virginica_labels <- rep(0, length(iris_test_y))
```

```
for(f in 1:length(iris_test_y)){
```

```
  if(iris_test_y[f] == "setosa"){
```

```
    setosa_labels[f] <- 1
```

```
  } else if(iris_test_y[f] == "versicolor"){
```

```
    versicolor_labels[f] <- 1
```

```
  } else if(iris_test_y[f] == "virginica"){
```

```
    virginica_labels[f] <- 1
```

```
  }
```

```
}
```

```
setosa_roc <- roc(setosa_labels, prob[, 'setosa'], auc.polygon = TRUE, max.auc.polygon = TRUE, print.auc = TRUE, show.thres = TRUE)
```

```
setosa_smooth_roc <- smooth(setosa_roc, method = "density")
```

```
plot(setosa_smooth_roc, col = 'red', xaxt = 'n', xlab = "False Positive Rate (1 - Specificity)", ylab = "True Positive Rate (Sensitivity)")
```

```
par(new = TRUE)
```

```
virginica_roc <- roc(virginica_labels, prob[, 'virginica'], auc.polygon = TRUE, max.auc.polygon = TRUE, print.auc = TRUE, show.thres = TRUE)
```

```
virginica_smooth_roc <- smooth(virginica_roc, method = "density")
```

```
plot(virginica_smooth_roc, col = 'green', xaxt='n', xlab="", ylab = "")
```

```
par(new = TRUE)
```

```
versicolor_roc<- roc(versicolor_labels, prob[, 'versicolor'], auc.polygon = TRUE, max.auc.polygon = TRUE,  
print.auc = TRUE, show.thres = TRUE)
```

```
versicolor_smooth_roc<- smooth(versicolor_roc, method = "density")
```

```
plot(versicolor_smooth_roc, col = 'blue', xaxt='n', xlab="", ylab = "")
```

```
y_labels<- c(setosa_labels,versicolor_labels,virginica_labels)
```

```
y_probs<- c(prob[, "setosa"],prob[, "versicolor"],prob[, "virginica"])
```

```
par(new = TRUE)
```

```
micro_roc<- roc(y_labels,y_probs, auc.polygon = TRUE, max.auc.polygon = TRUE, print.auc = TRUE,  
show.thres = TRUE)
```

```
micro_smooth_roc<- smooth(micro_roc, method = "density")
```

```
plot(micro_smooth_roc, col = 'black', lty = 'dotdash', xaxt='n', xlab="", ylab = "")
```

```
par(new=TRUE)
```

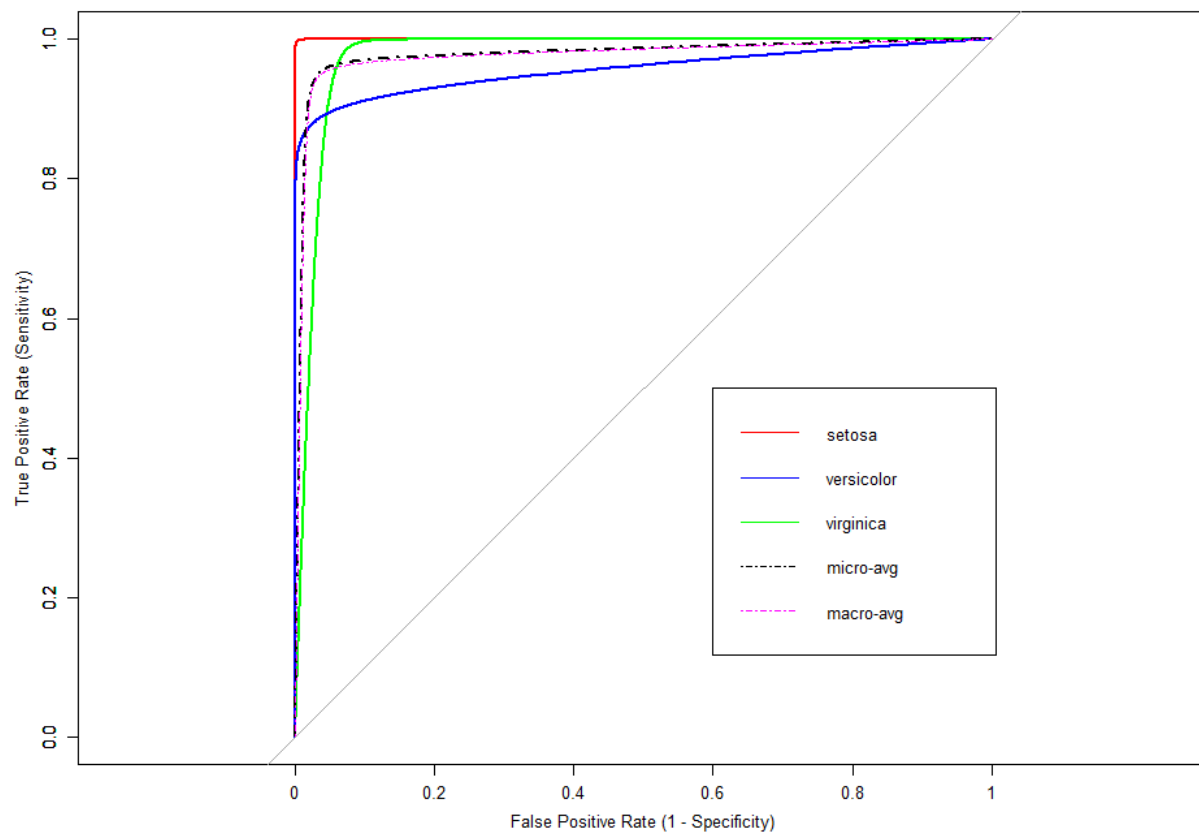
```
macro_sensitivity<- (setosa_smooth_roc$sensitivities + versicolor_smooth_roc$sensitivities +  
virginica_smooth_roc$sensitivities) / 3
```

```
macro_specificity<- (setosa_smooth_roc$specificities + versicolor_smooth_roc$specificities +  
virginica_smooth_roc$specificities) / 3
```

```
lines(macro_specificity, macro_sensitivity, type='l', xlim = rev(range(macro_specificity)), col='magenta',  
lty=4)
```

```
axis(1, at=(5:0) * 0.2, labels=(0:5) * 0.2, pos=c(-0.04,0))
```

```
legend(0.4, 0.5, legend = c('setosa', 'versicolor', 'virginica', 'micro-avg', 'macro-avg'), col = c('red', 'blue',  
'green', 'black', 'magenta'), lty = c(1,1,1,4,4))
```



```
##### AUC
#####
```

```
probs_train <- predict(model, iris_train_x, type = "raw") # Training AUC
```

```
multiclass.roc(iris_train_y, probs_train)
```

```
multiclass.roc(iris_test_y, prob) # Testing AUC
```

```
probs_all <- predict(model, iris[1:4], type = "raw") # Overall AUC
```

```
multiclass.roc(Species, probs_all)
```

**Results** -> The model identifies and classifies the species accordingly with a high rate of certainty, especially for Setosa.

		Train	Test	All
Accuracy		0.946	0.946	0.946
AUC		0.99	0.99	0.99
Macro Sensitivity		0.95	0.92	0.94
Macro Specificity		0.97	0.98	0.97
Sensitivity	Setosa	1.00	1.00	1.00
	Versicolor	0.95	0.84	0.90
	Virginica	0.89	1.00	0.94
Specificity	Setosa	1.00	1.00	1.00
	Versicolor	0.94	1.00	0.97
	Virginica	0.97	0.92	0.95
PPV	Setosa	1.00	1.00	1.00
	Versicolor	0.88	1.00	0.93
	Virginica	0.96	0.84	0.90
NPV	Setosa	1.00	1.00	1.00
	Versicolor	0.97	0.92	0.95
	Virginica	0.93	1.00	0.96