

heearrt.R

harsh

2023-06-12

```
setwd('C:\\Users\\harsh\\OneDrive\\Desktop\\A1a,b')  
getwd()
```

```
## [1] "C:/Users/harsh/OneDrive/Desktop/A1a,b"
```

```
library(dplyr)
```

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

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

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

```
library(tidyr)  
library(readxl)  
library(readr)  
  
heart<-read.csv("heart.csv")  
  
dim(heart)
```

```
## [1] 303  14
```

```
names(heart)
```

```
## [1] "age"      "sex"      "cp"      "trtbps"   "chol"     "fbs"  
## [7] "restecg"  "thalachh" "exng"    "oldpeak"  "slp"      "caa"  
## [13] "thall"    "output"
```

```
is.na(heart)
```

[illegible]

file:///C:/Users/harsh/OneDrive/Desktop/A1a,b/heeartt.html

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```

```
any(is.na(heart))
```

```
## [1] FALSE
```

```
sum(is.na(heart))
```

```
## [1] 0
```

```
sort(colSums(is.na(heart)), decreasing = TRUE)
```

```
##      age      sex      cp  trtbps      chol      fbs  restecg  thalachh
##      0       0       0       0       0       0       0       0
##  exng  oldpeak    slp      caa    thall  output
##      0       0       0       0       0       0
```

```
sapply(heart, class)
```

```
##      age      sex      cp  trtbps      chol      fbs  restecg  thalachh
## "integer" "integer" "integer" "integer" "integer" "integer" "integer" "integer"
##  exng  oldpeak    slp      caa    thall  output
## "integer" "numeric" "integer" "integer" "integer" "integer"
```

```
library(caTools)
library(pROC)
```

```
## Type 'citation("pROC")' for a citation.
```

```
##
## Attaching package: 'pROC'
```

```
## The following objects are masked from 'package:stats':
##
##      cov, smooth, var
```

```
set.seed(123)

split <- sample.split(heart$output, SplitRatio = 0.7)
train <- subset(heart, split == TRUE)
test <- subset(heart, split == FALSE)

model <- glm(output ~ age+sex+cp+trtbps+chol+fbs+restecg+thalachh+exng+oldpeak+slp+caa+thall,
data = train, family = binomial)

predicted_probs <- predict(model, newdata = test, type = "response")
predicted_class <- ifelse(predicted_probs >= 0.5, 1, 0)

library(caret)
```

```
## Loading required package: ggplot2
```

```
## Loading required package: lattice
```

```
confusion <- confusionMatrix(factor(predicted_class), factor(test$output))  
print(confusion)
```

```
## Confusion Matrix and Statistics  
##  
##           Reference  
## Prediction  0  1  
##           0 27  8  
##           1 14 42  
##  
##           Accuracy : 0.7582  
##           95% CI : (0.6572, 0.8419)  
##    No Information Rate : 0.5495  
##    P-Value [Acc > NIR] : 3.058e-05  
##  
##           Kappa : 0.5052  
##  
##    McNemar's Test P-Value : 0.2864  
##  
##           Sensitivity : 0.6585  
##           Specificity : 0.8400  
##           Pos Pred Value : 0.7714  
##           Neg Pred Value : 0.7500  
##           Prevalence : 0.4505  
##           Detection Rate : 0.2967  
##    Detection Prevalence : 0.3846  
##           Balanced Accuracy : 0.7493  
##  
##           'Positive' Class : 0  
##
```

```
library(pROC)
```

```
roc_obj <- roc(test$output, predicted_probs)
```

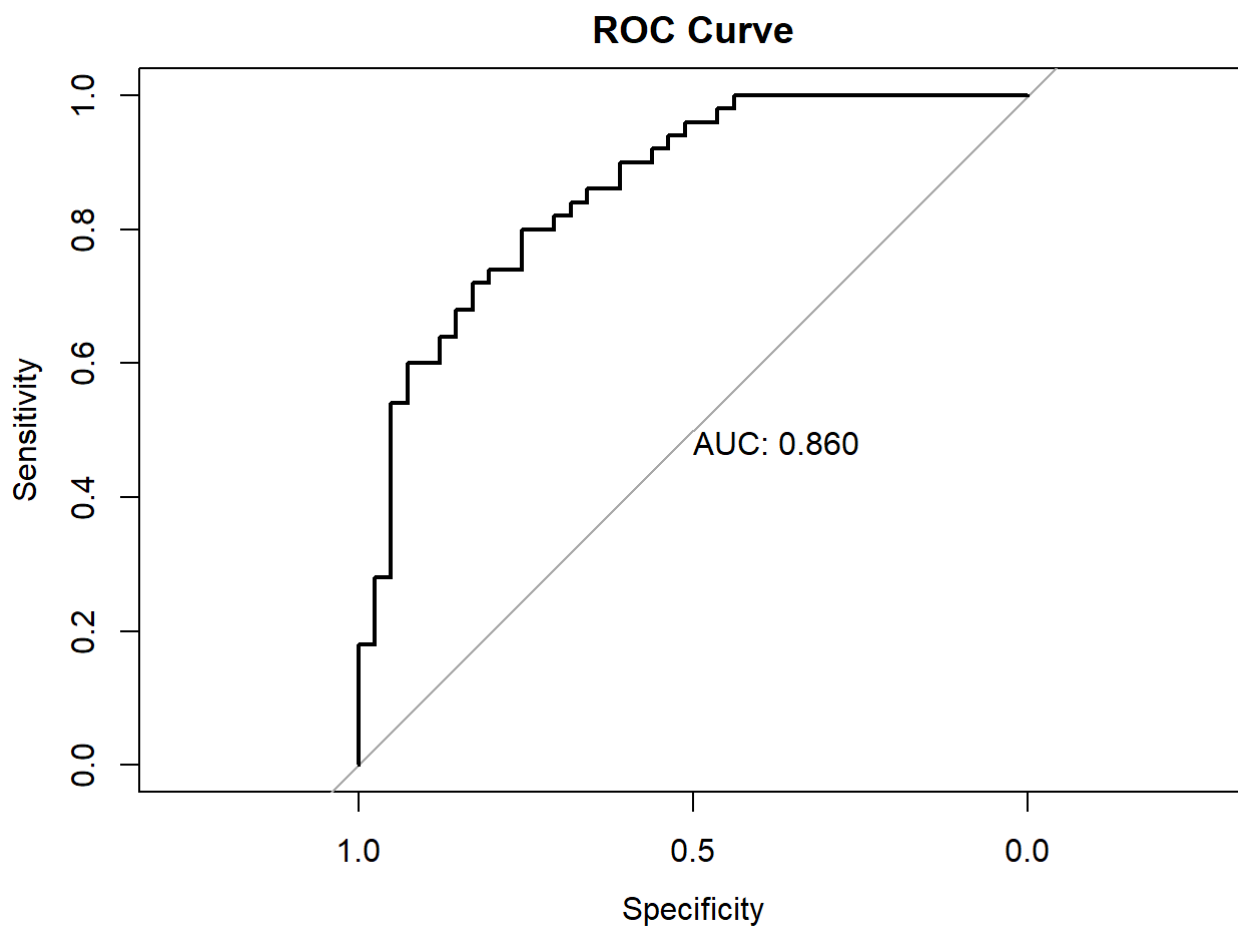
```
## Setting levels: control = 0, case = 1
```

```
## Setting direction: controls < cases
```

```
auc <- auc(roc_obj)  
print(paste("AUC-ROC:", auc))
```

```
## [1] "AUC-ROC: 0.86"
```

```
plot(roc_obj, main = "ROC Curve", print.auc = TRUE)
```



```
library(caTools)

set.seed(123)

split <- sample.split(heart$output, SplitRatio = 0.7)
train <- subset(heart, split == TRUE)
test <- subset(heart, split == FALSE)

library(rpart)

model <- rpart(output ~ ., data = train, method = "class")

predicted_probs <- predict(model, newdata = test, type = "prob")
predicted_class <- ifelse(predicted_probs[,2] >= 0.5, 1, 0)

confusion <- confusionMatrix(factor(predicted_class), factor(test$output))
print (confusion)
```

```
## Confusion Matrix and Statistics
##
##           Reference
## Prediction  0   1
##           0 28   9
##           1 13  41
##
##           Accuracy : 0.7582
##           95% CI : (0.6572, 0.8419)
##           No Information Rate : 0.5495
##           P-Value [Acc > NIR] : 3.058e-05
##
##           Kappa : 0.5074
##
##           McNemar's Test P-Value : 0.5224
##
##           Sensitivity : 0.6829
##           Specificity : 0.8200
##           Pos Pred Value : 0.7568
##           Neg Pred Value : 0.7593
##           Prevalence : 0.4505
##           Detection Rate : 0.3077
##           Detection Prevalence : 0.4066
##           Balanced Accuracy : 0.7515
##
##           'Positive' Class : 0
##
```

```
library(pROC)
```

```
roc_obj <- roc(test$output, predicted_probs[,2])
```

```
## Setting levels: control = 0, case = 1
## Setting direction: controls < cases
```

```
auc <- auc(roc_obj)
print(paste("AUC-ROC:", auc))
```

```
## [1] "AUC-ROC: 0.800487804878049"
```

```
plot(roc_obj, main = "ROC curve", print.auc = TRUE)
```