

Análisis Diabetes

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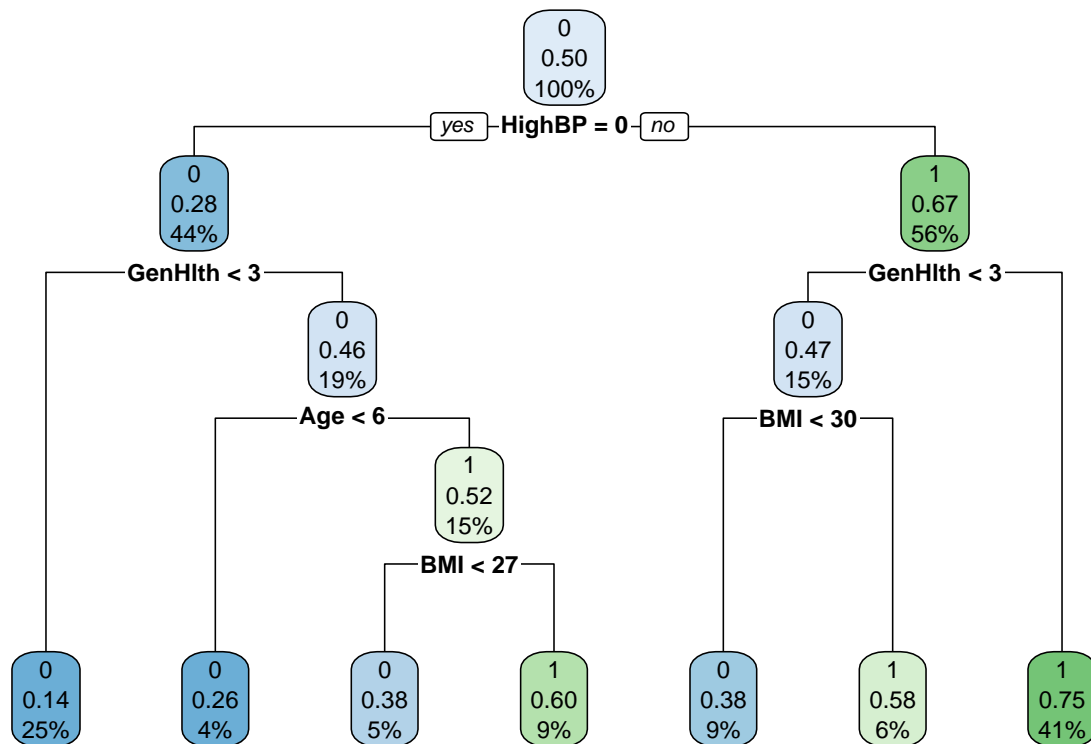
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R Markdown

```
data2 <- read.csv("data2.csv")
set.seed(123)
sample_data = sample.split(data2, SplitRatio = 0.7)
train_data <- subset(data2, sample_data == TRUE)
test_data <- subset(data2, sample_data == FALSE)

fit <- rpart(Diabetes_binary ~. , data=train_data, method = 'class')

rpart.plot(fit)
```



```
predict_unseen <- predict(fit, test_data, type = 'class')  
  
table_mat <- table(test_data$Diabetes_binary, predict_unseen)  
table_mat
```

```
##      predict_unseen  
##           0       1  
##    0 7304 3942  
##    1 2324 8923
```

```
accuracy_Test <- sum(diag(table_mat)) / sum(table_mat)  
accuracy_Test
```

```
## [1] 0.7214244
```