

Titanic - Decision Trees

[Code ▾](#)

```
library(rpart)
library(rattle)
library(dplyr)
library(RCurl)
library(ggplot2)
```

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1. Reading data

```
url <- getURL('https://raw.githubusercontent.com/frankwwu/R-Knots/master/Titanic/train.csv')
train <- read.csv(text = url)
url <- getURL('https://raw.githubusercontent.com/frankwwu/R-Knots/master/Titanic/test.csv')
test <- read.csv(text = url)
```

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2. Displaying data

```
str(train)
```

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```
'data.frame':   891 obs. of  12 variables:
 $ PassengerId: int   1  2  3  4  5  6  7  8  9 10 ...
 $ Survived   : int   0  1  1  1  0  0  0  0  1  1 ...
 $ Pclass     : int   3  1  3  1  3  3  1  3  3  2 ...
 $ Name       : Factor w/ 891 levels "Abbing, Mr. Anthony",...: 109 191 358 277 16 559 520 629 417 581 ...
 $ Sex        : Factor w/ 2 levels "female","male": 2 1 1 1 2 2 2 2 1 1 ...
 $ Age        : num   22 38 26 35 35 NA 54 2 27 14 ...
 $ SibSp      : int   1  1  0  1  0  0  0  3  0  1 ...
 $ Parch      : int   0  0  0  0  0  0  0  1  2  0 ...
 $ Ticket     : Factor w/ 681 levels "110152","110413",...: 524 597 670 50 473 276 86 396 345 133 ...
 $ Fare       : num   7.25 71.28 7.92 53.1 8.05 ...
 $ Cabin      : Factor w/ 148 levels "", "A10", "A14",...: 1 83 1 57 1 1 131 1 1 1 ...
 $ Embarked   : Factor w/ 4 levels "", "C", "Q", "S": 4 2 4 4 4 3 4 4 4 2 ...
```

```
str(test)
```

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```
'data.frame':   418 obs. of  11 variables:
 $ PassengerId: int  892 893 894 895 896 897 898 899 900 901 ...
 $ Pclass     : int   3  3  2  3  3  3  3  2  3  3 ...
 $ Name       : Factor w/ 418 levels "Abbott, Master. Eugene Joseph",...: 210 409 273 414 182 370 85 58 5 104 ..
 .
 $ Sex        : Factor w/ 2 levels "female","male": 2 1 2 2 1 2 1 2 1 2 ...
 $ Age        : num  34.5 47 62 27 22 14 30 26 18 21 ...
 $ SibSp      : int   0  1  0  0  1  0  0  1  0  2 ...
 $ Parch      : int   0  0  0  0  1  0  0  1  0  0 ...
 $ Ticket     : Factor w/ 363 levels "110469","110489",...: 153 222 74 148 139 262 159 85 101 270 ...
 $ Fare       : num   7.83 7 9.69 8.66 12.29 ...
 $ Cabin      : Factor w/ 77 levels "", "A11", "A18",...: 1 1 1 1 1 1 1 1 1 1 ...
 $ Embarked   : Factor w/ 3 levels "C", "Q", "S": 2 3 2 3 3 3 2 3 1 3 ...
```

3. Removing NAs

```
train <- train %>% na.omit()
test <- test %>% na.omit()
```

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4. Converting categorical variables to factors

```
train$Survived <- factor(train$Survived)
train$Pclass <- factor(train$Pclass)
test$Pclass <- factor(test$Pclass)
```

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5. Visualizing the training data

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```
ggplot(train, aes(Age, Fare, color=Survived)) +
  geom_point(alpha = 0.5) +
  facet_grid(Pclass~Sex) +
  ggtitle("Training Data")
```



6. Selecting features

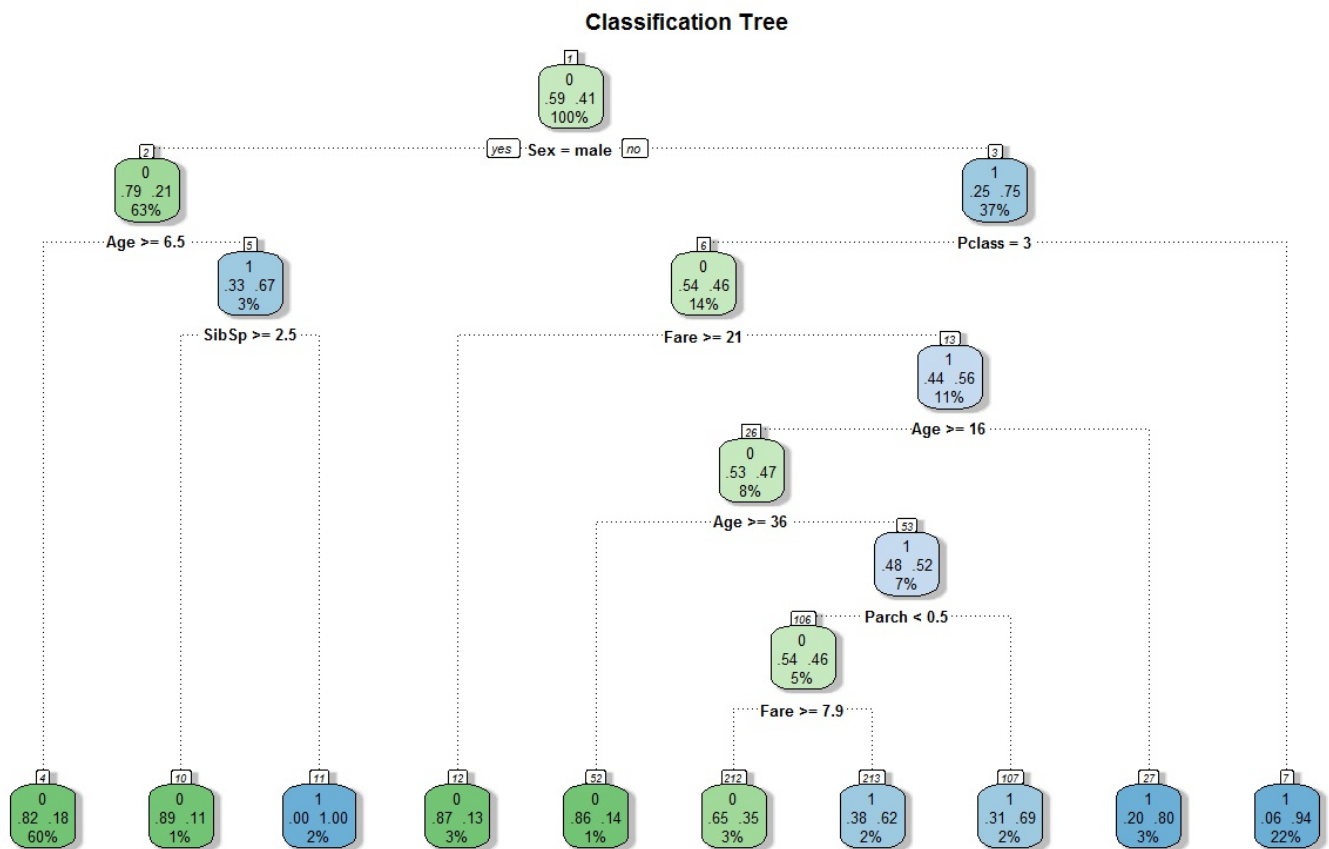
```
formula = Survived ~ Pclass + Sex + Age + SibSp + Parch + Fare + Embarked
```

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7. Creating the Classification Tree

```
set.seed(9)
tree <- rpart(formula, data=train, method="class")
fancyRpartPlot(tree, uniform=TRUE, main="Classification Tree")
```

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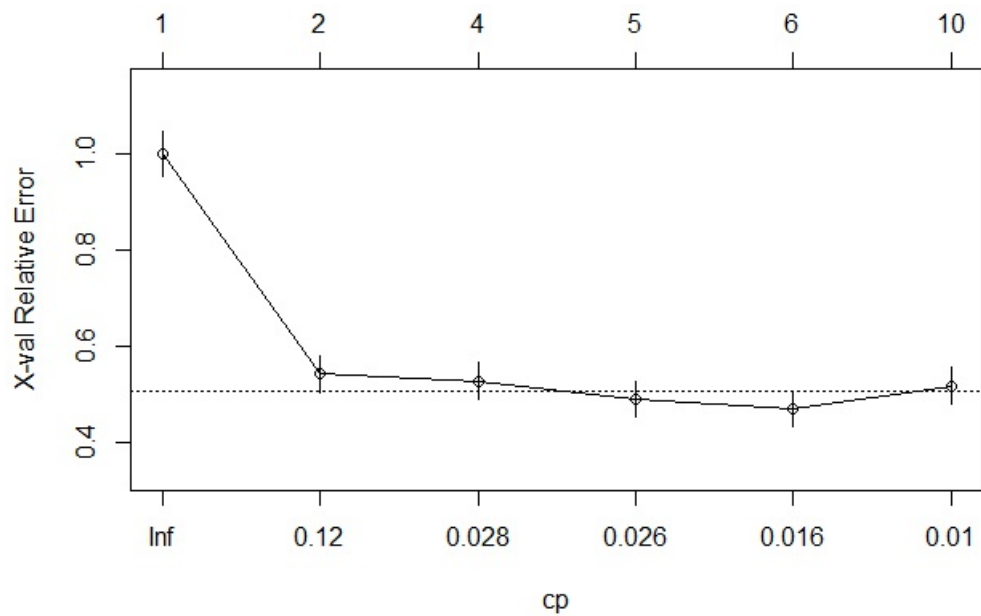
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8. Cross-Validation

To examine whether the tree model is over fitting, find the size of tree with the minimum error.

```
plotcp(tree)
```

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```
tree$cpstable[which.min(tree$cpstable[, "xerror"]), "CP"]
```

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```
[1] 0.01034483
```

```
printcp(tree)
```

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```
Classification tree:
rpart(formula = formula, data = train, method = "class")
```

```
Variables actually used in tree construction:
[1] Age    Fare    Parch  Pclass Sex    SibSp
```

```
Root node error: 290/714 = 0.40616
```

```
n= 714
```

| | CP | nsplit | rel error | xerror | xstd |
|---|----------|--------|-----------|---------|----------|
| 1 | 0.458621 | 0 | 1.00000 | 1.00000 | 0.045252 |
| 2 | 0.029310 | 1 | 0.54138 | 0.54138 | 0.038162 |
| 3 | 0.027586 | 3 | 0.48276 | 0.52759 | 0.037808 |
| 4 | 0.024138 | 4 | 0.45517 | 0.48966 | 0.036779 |
| 5 | 0.010345 | 5 | 0.43103 | 0.46897 | 0.036181 |
| 6 | 0.010000 | 9 | 0.38966 | 0.51724 | 0.037535 |

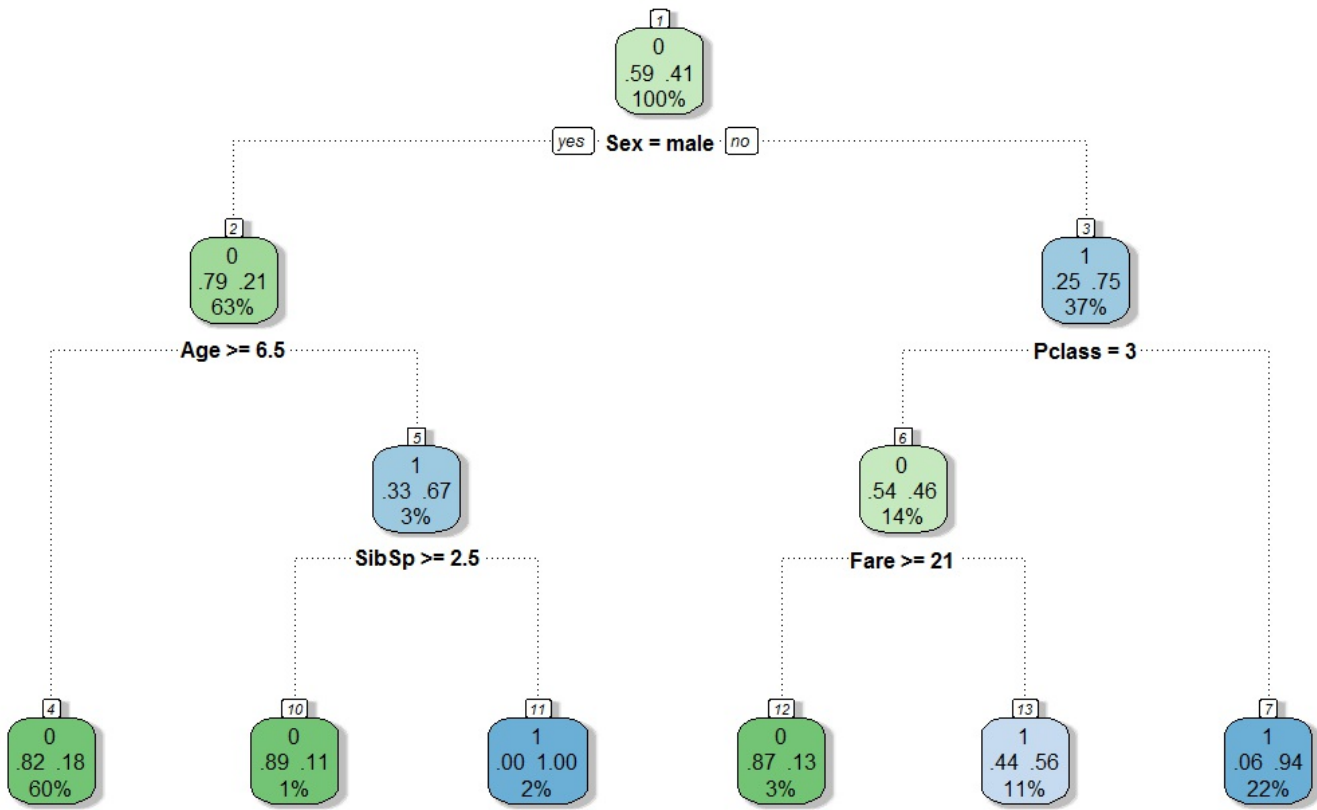
9. Pruning the Tree

Prune the over fitting notes.

```
trim <- tree$cpstable[which.min(tree$cpstable[, "xerror"]), "CP"]
ptree<- prune(tree, cp=trim)
fancyRpartPlot(ptree, uniform=TRUE, main="Pruned Classification Tree")
```

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Pruned Classification Tree



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10. Predicting with the test data

```
predict <- predict(ptree, test, type = "prob")
```

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11. Visualizing the result

```
test$Survived <- predict[,2]
ggplot(test, aes(Age, Fare, color=Survived)) +
  geom_point(alpha = 0.5) +
  facet_grid(Pclass~Sex) +
  ggtitle("Prediction with the Test Data ")
```

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