

Data Science

R 강의: pruning of rpart tree

rpart 나무의 post pruning

■ cost-complexity pruning

- error 와 나무의 complexity 를 함께 고려 하여 가지치기
- 방법: full tree 생성 후
cost(or cost-complexity) 함수를 최소화 하도록
rpart 나무를 가지 치기
- $\text{cost} = \text{error} + \text{complexity}$
- $\text{complexity} = \text{cp} * \text{잎노드의 수}$
- cp: 잎노드 하나에 부과하는 penalty

■ 최적 cp

- 실험적으로 정함

최적 cp 값 검색

■ 방법

- Full tree 생성하고 이 tree 에 대하여
 - cp (complexity parameter) 를 0으로부터 시작하여 증가 시켜 가며 각 cp 값에 대하여
 - 최적 pruning 을 시도하고 pruning 된 나무에 대하여
 - xerror (cross validated relative error)를 측정합니다.
- xerror 가 최소인 경우의 cp 값이 최적 이므로
- 이 cp 값을 선택하여 full tree 를 pruning → final model
- Tree 생성 및 성능 측정시 Cross validation 방법 사용.
- xerror 는 순수한 cross validated error, 복잡도 포함 안되어 있음.
- cost 계산시 error 는 재대입 error = rel error 사용

http://mlwiki.org/index.php/Cost-Complexity_Pruning

printcp(rpart_model)

■ printcp(rpart_model): rpart 모델 출력

| Record | Report | | | | | | | | | | | | | | | | | | | | | | | | |
|--------|---|------------|-----------|------------|-----------|---------|-----------|---|------|---|------|------|----------|---|------|---|------|------|----------|---|------|---|------|------|----------|
| 1 | Summary Report for Decision Tree Model IrisDecisionTree | | | | | | | | | | | | | | | | | | | | | | | | |
| 2 | Call: rpart(formula = Species ~ SepalLengthCm + SepalWidthCm + PetalLengthCm + PetalWidthCm, data = the.data, minsplit = 20, minbucket = 7, xval = 10, maxdepth = 20, cp = 0, usesurrogate = 0, surrogatestyle = 0) | | | | | | | | | | | | | | | | | | | | | | | | |
| 3 | Model Summary Variables actually used in tree construction: [1] PetalLengthCm PetalWidthCm Root node error: 100/150 = 0.66667 n= 150 | | | | | | | | | | | | | | | | | | | | | | | | |
| 4 | Pruning Table | | | | | | | | | | | | | | | | | | | | | | | | |
| 5 | <table><tr><th>Level</th><th>CP</th><th>Num Splits</th><th>Rel Error</th><th>X Error</th><th>X Std Dev</th></tr><tr><td>1</td><td>0.50</td><td>0</td><td>1.00</td><td>1.17</td><td>0.050735</td></tr><tr><td>2</td><td>0.44</td><td>1</td><td>0.50</td><td>0.73</td><td>0.061215</td></tr><tr><td>3</td><td>0.00</td><td>2</td><td>0.06</td><td>0.10</td><td>0.030551</td></tr></table> | Level | CP | Num Splits | Rel Error | X Error | X Std Dev | 1 | 0.50 | 0 | 1.00 | 1.17 | 0.050735 | 2 | 0.44 | 1 | 0.50 | 0.73 | 0.061215 | 3 | 0.00 | 2 | 0.06 | 0.10 | 0.030551 |
| Level | CP | Num Splits | Rel Error | X Error | X Std Dev | | | | | | | | | | | | | | | | | | | | |
| 1 | 0.50 | 0 | 1.00 | 1.17 | 0.050735 | | | | | | | | | | | | | | | | | | | | |
| 2 | 0.44 | 1 | 0.50 | 0.73 | 0.061215 | | | | | | | | | | | | | | | | | | | | |
| 3 | 0.00 | 2 | 0.06 | 0.10 | 0.030551 | | | | | | | | | | | | | | | | | | | | |
| 6 | Leaf Summary node), split, n, loss, yval, (yprob) * denotes terminal node 1) root 150 100 Iris-setosa (0.33333333 0.33333333 0.33333333) 2) PetalLengthCm< 2.45 50 0 Iris-setosa (1.00000000 0.00000000 0.00000000) * 3) PetalLengthCm>=2.45 100 50 Iris-versicolor (0.00000000 0.50000000 0.50000000) 6) PetalWidthCm< 1.75 54 5 Iris-versicolor (0.00000000 0.90740741 0.09259259) * 7) PetalWidthCm>=1.75 46 1 Iris-virginica (0.00000000 0.02173913 0.97826087) * | | | | | | | | | | | | | | | | | | | | | | | | |

printcp(rpart_model)

■ Root node error

- error on root node

■ Pruning Table

- CP
 - 앞 노드 한 개당 penalty
- 각 CP 값에 대하여
 - Num Splits: number of splits in the tree
 - Rel(ative) error: resubstitution error rate scaled by Root node error, scaled by $\text{root_node_error}(= \text{resubstitution error} / \text{root_node_error})$
 - X error: Cross validated error rate also, scaled by root_node_error
 - X std dev: Standard error of CV

printcp(rpart_model)

■ Example

```
> printcp(rpartmod)
```

```
Classification tree:
```

```
rpart(formula = AHD ~ ., data = train, method = "class")
```

```
Variables actually used in tree construction:
```

```
[1] Ca          ChestPain MaxHR          Oldpeak    Slope          Thal
```

```
Root node error: 98/213 = 0.46009
```

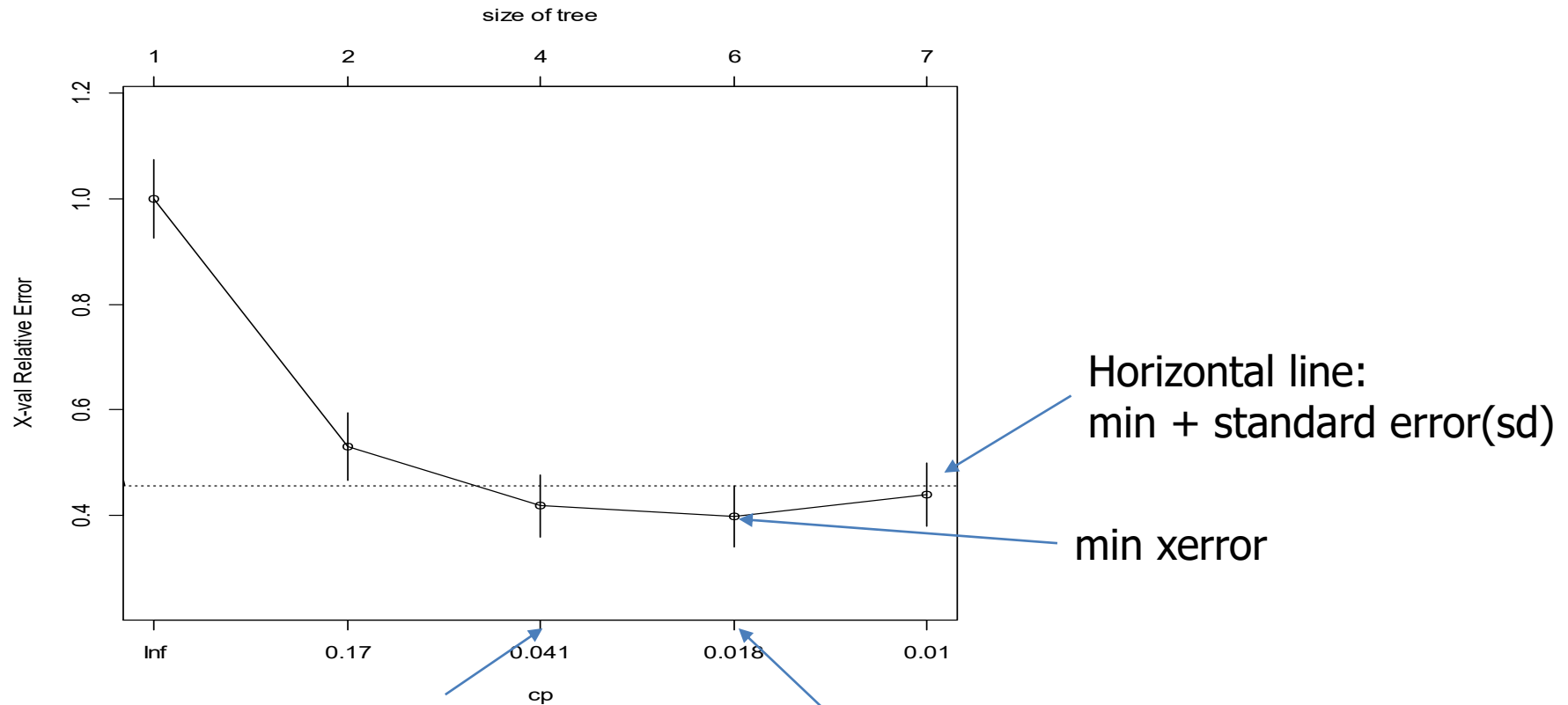
```
n= 213
```

| | CP | nsplit | rel error | xerror | xstd |
|---|----------|--------|-----------|---------|----------|
| 1 | 0.530612 | 0 | 1.00000 | 1.00000 | 0.074224 |
| 2 | 0.056122 | 1 | 0.46939 | 0.53061 | 0.063973 |
| 3 | 0.030612 | 3 | 0.35714 | 0.41837 | 0.058714 |
| 4 | 0.010204 | 5 | 0.29592 | 0.39796 | 0.057596 |
| 5 | 0.010000 | 6 | 0.28571 | 0.43878 | 0.059778 |

plotcp(rpart_model)

■ optimal cp value

- Choose the leftmost one below the horizontal line



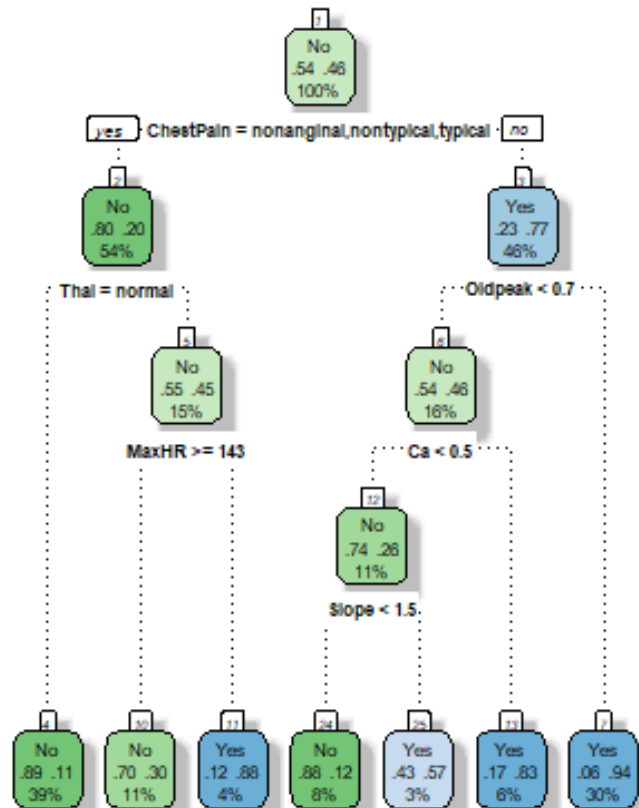
SE 범위 내에서 가장 작은
(간단한) 모델(cp = 0.041)을
최적모델(최적값)으로 선택

cp = 0.018 인 경우의
모델의 xerror 가
최소 이지만

Full tree vs pruned tree

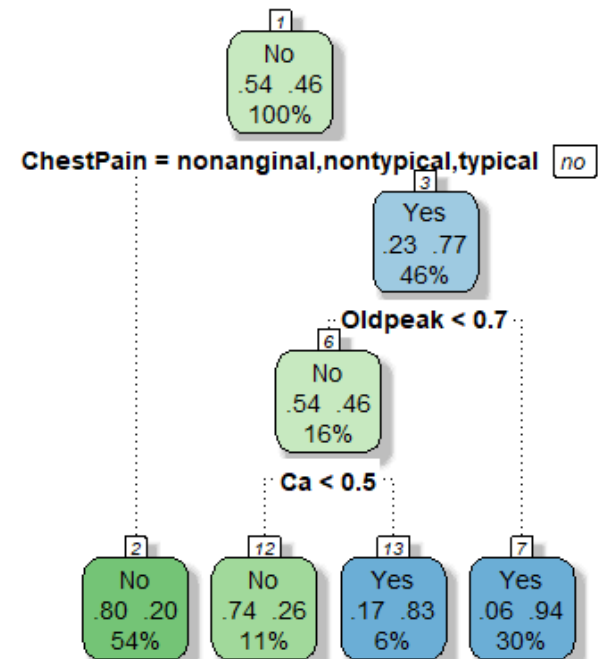
`prune(rpart_model, cp = 0.041)`

Full tree(acc = 0.71)



Rattle 2020-10-04 15:10:20 Friday

Pruned tree(acc = 0.73)



Rattle 2020-10-04 15:12:34 Friday

예제 코드 – pruning 예제

```
# rpart tree pruning
```

```
# Source
```

```
# https://rstudio-pubs-static.s3.amazonaws.com/332255\_d2b0ffd26ff64704a65573ba82a90ad6.html
```

```
# Data
```

```
# heart.csv
```

```
# https://www.kaggle.com/zhaoyingzhu/Heartcsv
```

```
# Heart disease data
```

```
# Target var : AHD with two values Yes and No
```

```
library(rpart)
```

```
library(caret)
```

```
library(e1071)
```

```
library(rattle)
```

```
library(tidyverse)
```

```
df<-read.csv('Heart.csv')
```

```
str(df)
```

```
head(df)
```

```
df$AHD <- as.factor(df$AHD)
```

```
# data partition
```

```
set.seed(10) # reproducibility setting
```

```
intrain<-createDataPartition(y=df$AHD, p=0.7, list=FALSE)
```

```
train<-df[intrain, ]
```

```
test<-df[-intrain, ]
```

```
# Full tree
```

```
rpartmod<-rpart(AHD~. , data=train, method="class")
```

```
fancyRpartPlot(rpartmod)
```

```
rpartpred<-predict(rpartmod, test, type='class')
```

```
confusionMatrix(rpartpred, test$AHD)
```

```
printcp(rpartmod)
```

```
plotcp(rpartmod)
```

```
# choose the model (cp= 0.041) leftmost under the horizontal line
```

```
# which xerror is within the range of standard error
```

```
# from min xerror of the model with cp = 0.18
```

```
ptree<-prune(rpartmod, cp = 0.041)
```

```
fancyRpartPlot(ptree)
```

```
rpartpred<-predict(ptree, test, type='class')
```

```
confusionMatrix(rpartpred, test$AHD)
```

■ rpart pruning

- <https://dzone.com/articles/decision-trees-and-pruning-in-r>
- https://rstudio-pubs-static.s3.amazonaws.com/332255_d2b0ffd26ff64704a65573ba82a90ad6.html

Note:

Q: Why are the cp values in plotcp() chart modified from the original table?

**A: printcp() gives the minimal cp for which the pruning happens.
plotcp() plots against the geometric mean**

<https://stackoverflow.com/questions/31546895/why-are-the-cp-values-in-plotcp-chart-modified-from-the-original-table>

Note: How to select the optimal cp value for pruning

1. Use the first level (i.e. least nsplit) with minimum xerror.
The first level only kicks in when there are multiple level having same, minimum xerror. This is the most common used method.
2. Use the first level where xerror falls into the ± 1 xstd range of $\min(\text{xerror})$, i.e., $\text{xerror} < \min(\text{xerror}) + \text{xstd}$, the level whose xerror is at or below horizontal line. This method takes into account the variability of xerror resulting from cross-validation.
Note: rel_error should NOT be used in pruning.
3. (A rarely used method) Use the first level where $\text{xerror} \pm \text{xstd}$ overlaps with $\min(\text{xerror}) \pm \text{xstd}$. i.e., the level whose lower limit is at or below horizontal line.