



MACHINE LEARNING WITH TREE-BASED MODELS IN R

# Introduction to regression trees

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Instructor



# Train a Regresion Tree in R

```
> rpart(formula = ____,  
        data = ____,  
        method = ____)
```

formula	is in the format: outcome ~ predictor1+predictor2+etc
data=	specifies the dataframe
method	"class" for classification tree "anova" for regression tree
control=	<i>optional</i> parameters for controlling the tree growth



# Train/Validation/Test Split

- training set
- validation set
- test set



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# Performance metrics for regression

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# Common metrics for regression

- Mean Absolute Error (MAE)

$$MAE = \frac{1}{n} \sum | actual - predicted |$$

- Root Mean Square Error (RMSE)

$$RMSE = \sqrt{\frac{1}{n} \sum (actual - predicted)^2}$$



# Evaluate a regression tree model

```
> pred <- predict(object = model, # model object
                  newdata = test) # test dataset

> library(Metrics)

# Compute the RMSE
> rmse(actual = test$response, # the actual values
        predicted = pred)      # the predicted values
[1] 2.278249
```



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**What are the  
hyperparameters for a  
decision tree?**

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# Decision tree hyperparameters

```
?rpart.control
```

```
rpart.control {rpart}
```

## Control for Rpart Fits

### Description

Various parameters that control aspects of the `rpart` fit.

### Usage

```
rpart.control(minsplit = 20, minbucket = round(minsplit/3), cp = 0.01,  
              maxcompete = 4, maxsurrogate = 5, usesurrogate = 2, xval = 10,  
              surrogatestyle = 0, maxdepth = 30, ...)
```



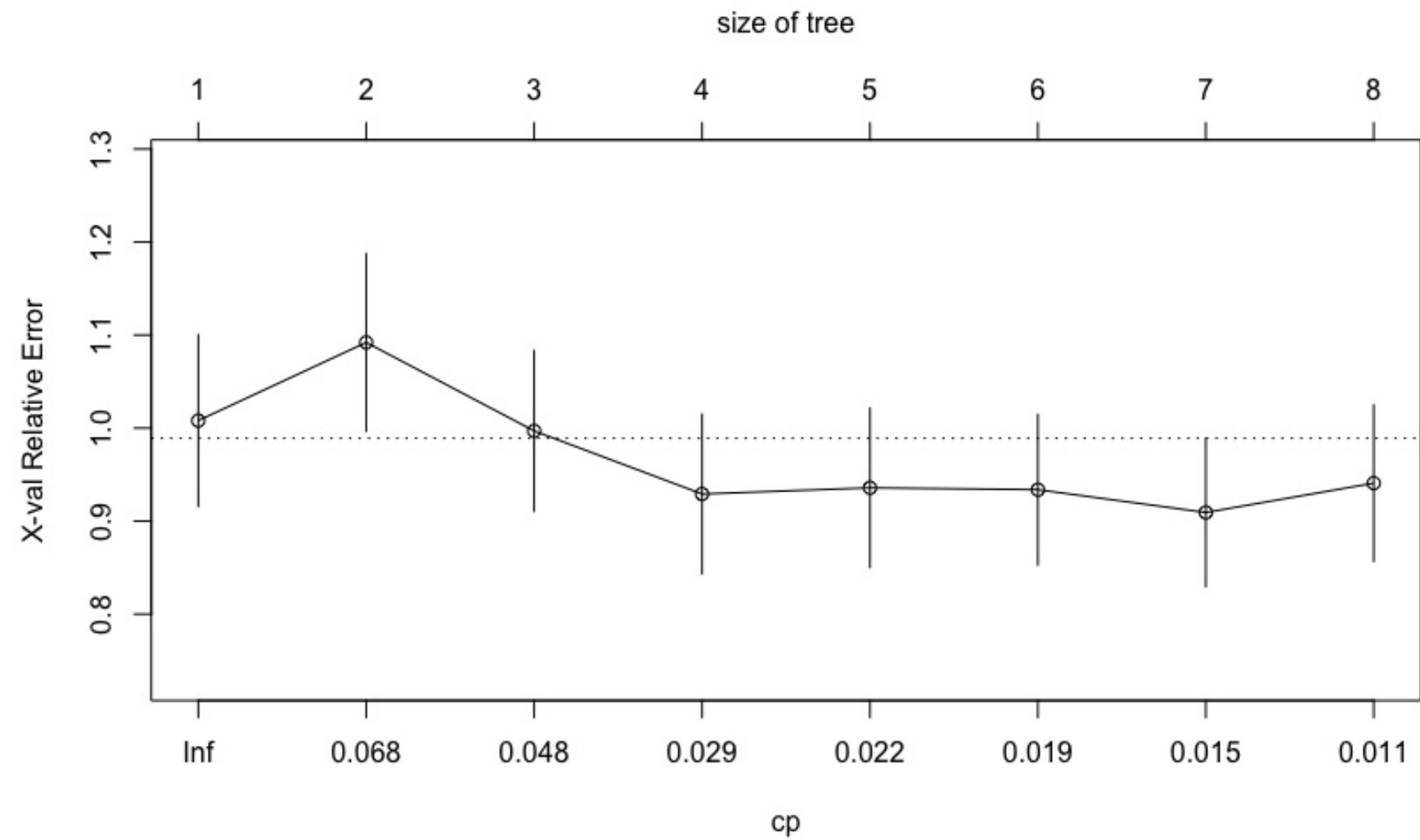
# Decision tree hyperparameters

- `minsplit`: minimum number of data points required to attempt a split
- `cp`: complexity parameter
- `maxdepth`: depth of a decision tree



# Cost-Complexity Parameter (CP)

```
> plotcp(grade_model)
```





# Cost-Complexity Parameter (CP)

```
> print(model$cpstable)
```

	CP	nsplit	rel error	xerror	xstd
1	0.06839852	0	1.0000000	1.0080595	0.09215642
2	0.06726713	1	0.9316015	1.0920667	0.09543723
3	0.03462630	2	0.8643344	0.9969520	0.08632297
4	0.02508343	3	0.8297080	0.9291298	0.08571411
5	0.01995676	4	0.8046246	0.9357838	0.08560120
6	0.01817661	5	0.7846679	0.9337462	0.08087153
7	0.01203879	6	0.7664912	0.9092646	0.07982862
8	0.01000000	7	0.7544525	0.9407895	0.08399125

```
# Prune the model (to optimized cp value)  
# Returns the optimal model
```

```
model_opt <- prune(tree = model, cp = cp_opt)
```



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# Grid Search for model selection

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# Grid Search

- What is a model hyperparameter?
- What is a "grid"?
- What is the purpose of a grid search?
- How is the best model chosen?





# Set up the grid

```
# Establish a list of possible values for minsplit and maxdepth
```

```
> minsplit <- seq(1, 30, 5)  
> maxdepth <- seq(5, 40, 10)
```

```
# Create a data frame containing all combinations
```

```
> hyper_grid <- expand.grid(minsplit = minsplit,  
                             maxdepth = maxdepth)
```

```
> hyper_grid[1:10,]  
  minsplit maxdepth  
1         1         5  
2         6         5  
3        11         5  
4        16         5  
5        21         5  
6        26         5  
7         1        15  
8         6        15  
9        11        15  
10       16        15
```



# Grid Search in R: Train models

```
# create an empty list to store models

models <- list()

# execute the grid search

> for (i in 1:nrow(hyper_grid)) {

  # get minsplit, maxdepth values at row i
  minsplit <- hyper_grid$minsplit[i]
  maxdepth <- hyper_grid$maxdepth[i]

  # train a model and store in the list
  models[[i]] <- rpart(formula = response ~ .,
                      data = train,
                      method = "anova",
                      minsplit = minsplit)

}
```



# Grid Search in R: Evaluate models

```
# create an empty vector to store RMSE values
rmse_values <- c()

# compute validation RMSE for
for (i in 1:length(models)) {

  # retrieve the i^th model from the list
  model <- models[[i]]

  # generate predictions on grade_valid
  pred <- predict(object = model,
                  newdata = valid)

  # compute validation RMSE and add to the
  rmse_values[i] <- rmse(actual = valid$response,
                       predicted = pred)
}
```



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