

# Ch 8.2.1, 8.2.2: Bagging and Random Forests

## Lecture 25 - CMSE 381

Prof. Mengsen Zhang

Michigan State University  
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Dept of Computational Mathematics, Science & Engineering

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# Announcements

## Last time:

- 8.1 Decision Trees - regression
- 8.1 Decision Trees - classification

## This lecture:

- 8.2.1 Bagging
- 8.2.2 Random forest

## Announcements:

- Homework 7 Due Sunday

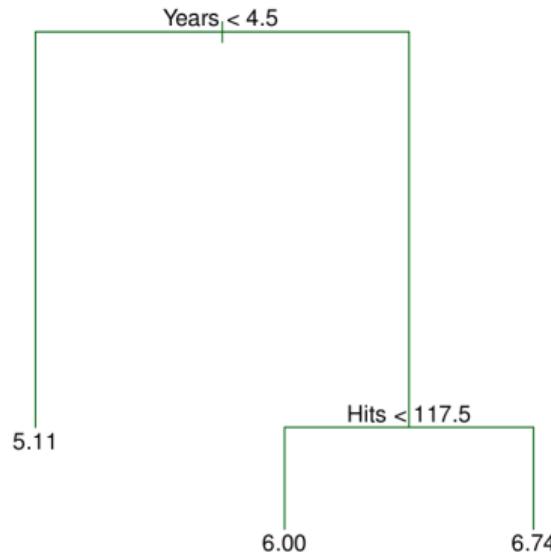
21	F	10/24	Polynomial & Step Functions	7.1-7.2	
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32	W	11/19	Unsupervised learning / clustering	12.1, 12.4	
33	F	11/21	Review		HW #9 Due Sun 11/23
	M	11/24	Midterm #3		
	W	11/26	Virtual: Project Office Hours		
	F	11/28	Thanksgiving		
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	F	12/5			Project Due
	M	12/8			
	W	12/10			
	F	12/12	No final exam		Honors Project Due

## Section 1

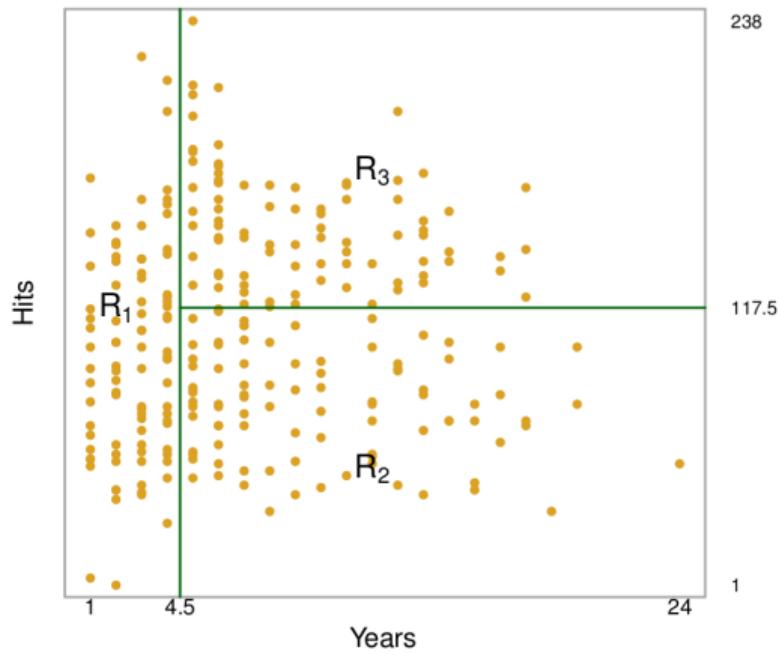
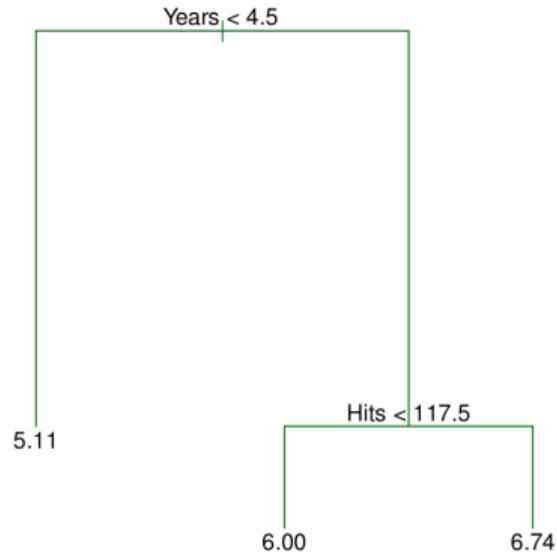
Previously

# First decision tree example

	Hits	Years	LogSalary
1	81	14	6.163315
2	130	3	6.173786
3	141	11	6.214608
4	87	2	4.516339
5	169	11	6.620073
...	...	...	...
317	127	5	6.551080
318	136	12	6.774224
319	126	6	5.953243
320	144	8	6.866933
321	170	11	6.907755

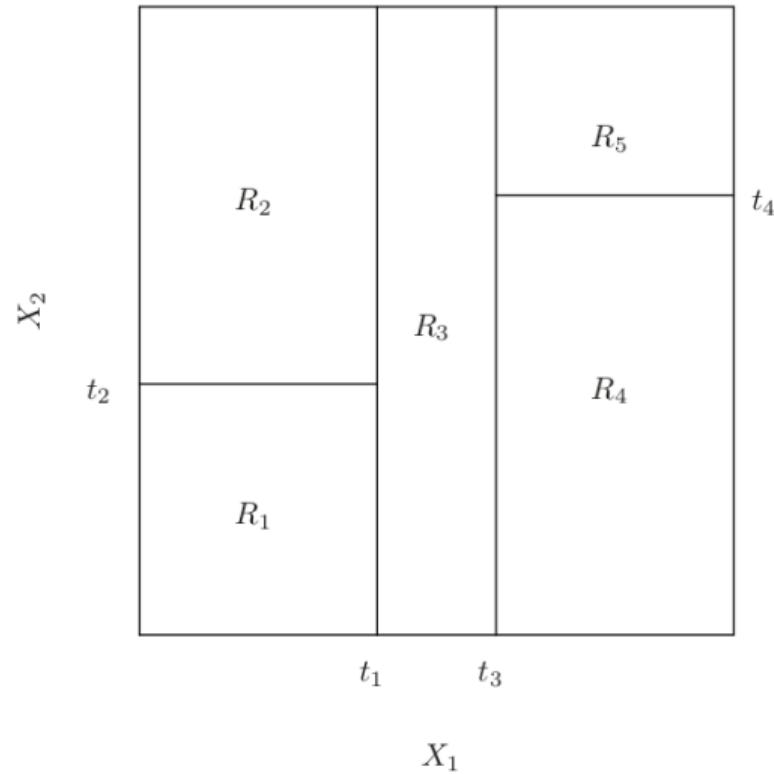


# Viewing Regions Defined by Tree



## How do we actually get the tree? Two steps

- ① We divide the predictor space – that is, the set of possible values for  $X_1, X_2, \dots, X_p$  — into  $J$  distinct and non-overlapping regions,  $R_1, R_2, \dots, R_J$ .
- ② For every observation that falls into the region  $R_j$ , we make the same prediction = the mean of the response values for the training observations in  $R_j$ .



# Recursive binary splitting

**Goal:**

Find boxes  $R_1, \dots, R_J$  that minimize

$$\sum_{j=1}^J \sum_{i \in R_j} (y_i - \hat{y}_{R_j})^2$$

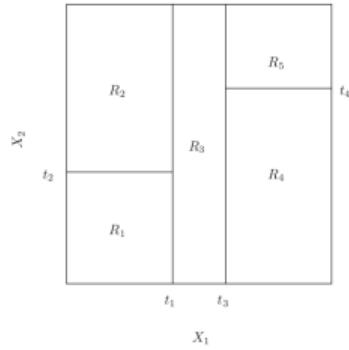
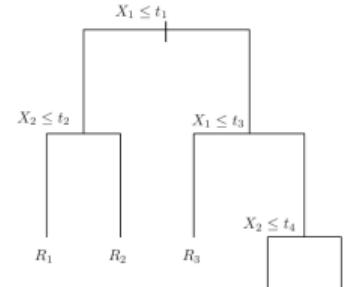
$\hat{y}_{R_j}$  = mean response for training observations in  $j$ th box

Pick  $s$  so that splitting into  $\{X \mid X_j < s\}$  and  $\{X \mid X_j \geq s\}$  results in largest possible reduction in RSS:

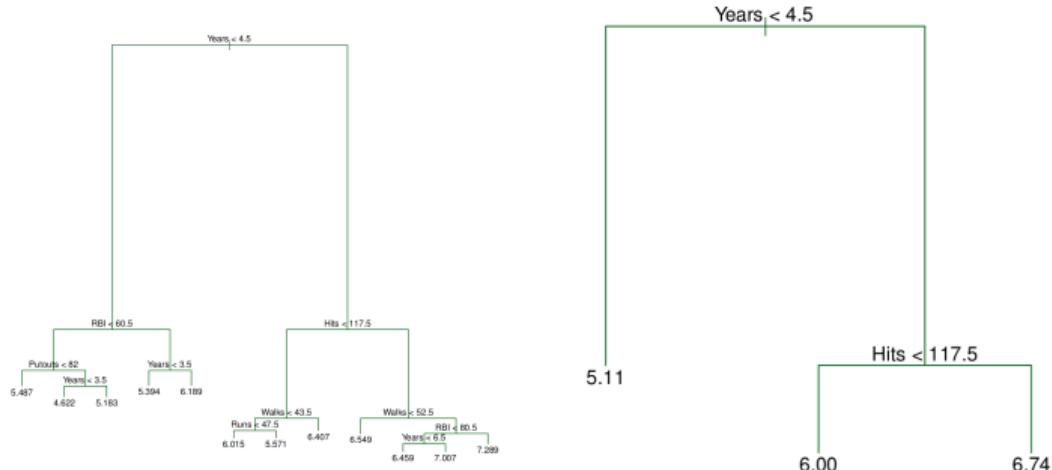
$$R_1(j, s) = \{X \mid X_j < s\}$$

$$R_2(j, s) = \{X \mid X_j \geq s\}$$

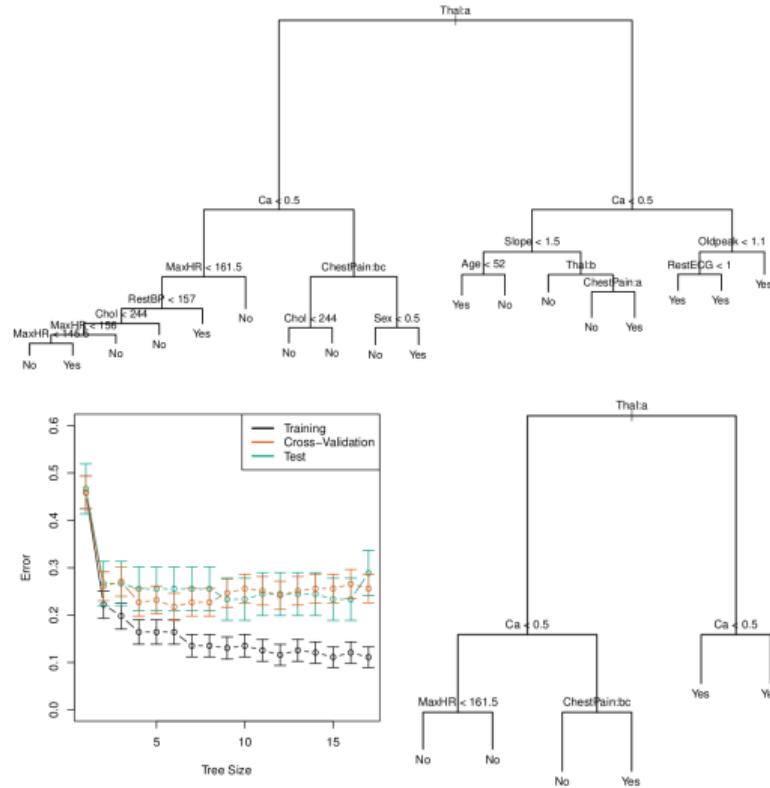
$$\sum_{i|x_i \in R_1(j, s)} (y_i - \hat{y}_{R_1})^2 + \sum_{i|x_i \in R_2(j, s)} (y_i - \hat{y}_{R_2})^2$$



# Pruning



# Classification version

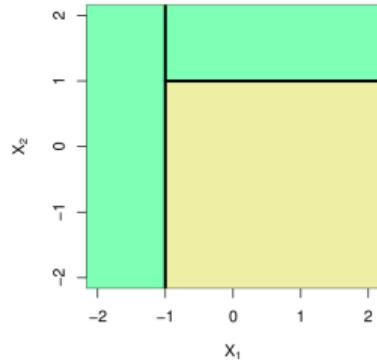
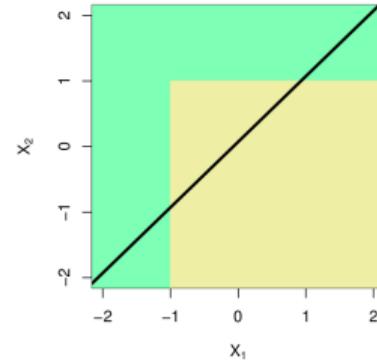
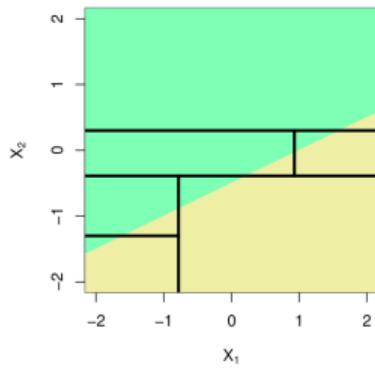
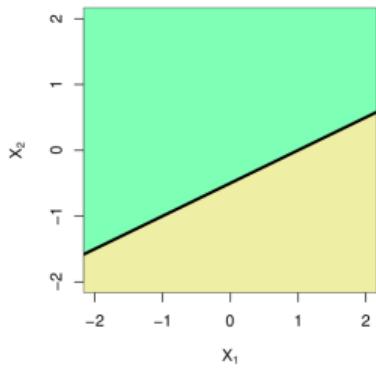


## Evaluating the splits:

- $\hat{p}_{mk}$  = proportion of training observations in  $R_m$  from the  $k$ th class
- Error:  $E = 1 - \max_k(\hat{p}_{mk})$
- Gini index:

$$G = \sum_{k=1}^K \hat{p}_{mk}(1 - \hat{p}_{mk})$$

# Linear models vs trees



# What will you learn today?

- What does bagging of decision trees accomplish?
- How do you use out of bag error estimation for decision trees?
  - ▶ You should be able to describe this for both regression and classification trees.
- What problem of bagging does using random forest address?
- What is the relationship between random forest and bagging?

## Section 2

### 8.2.1 Bagging

# Use ensemble of trees to reduce variance

## Want to do (but can't):

Build separate models from independent training sets, and average resulting predictions:

- $\hat{f}^1(x), \dots, \hat{f}^B(x)$  for  $B$  separate training sets
- Return the average

$$\hat{f}_{avg}(x) = \frac{1}{B} \sum_{b=1}^B \hat{f}^b(x)$$

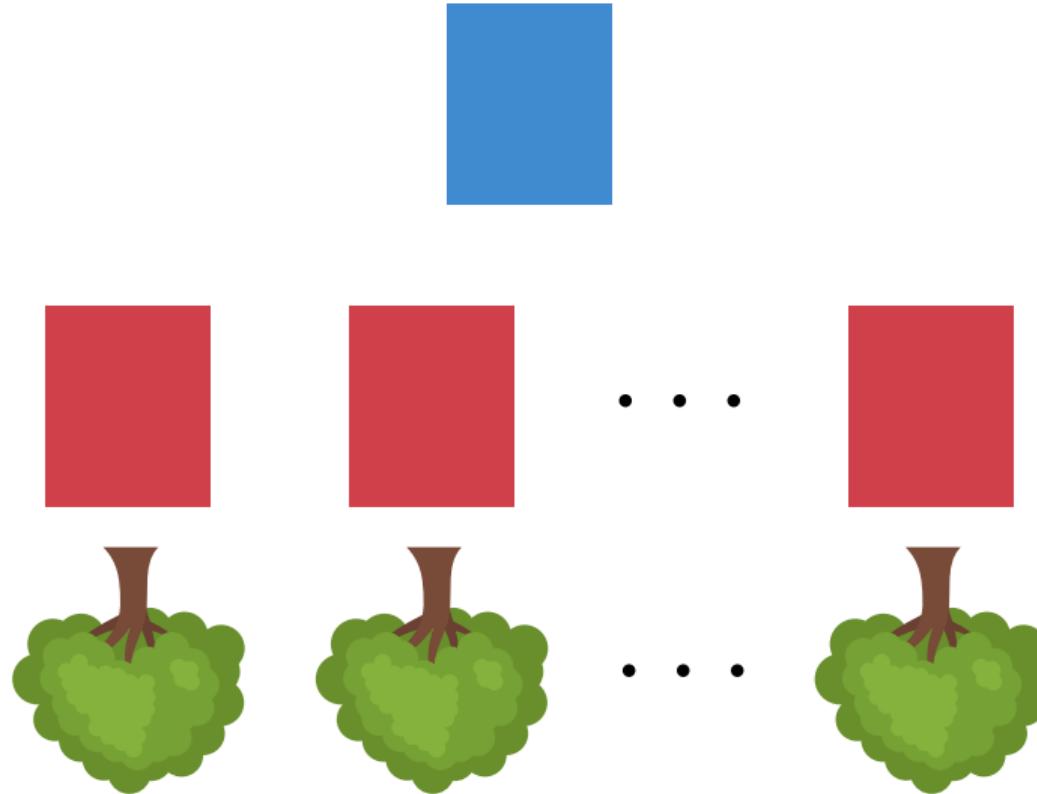
## Bootstrap modification:

- Work with fixed data set
- Take  $B$  samples from this data set (with replacement)
- Train method on  $b$ th sample to get  $\hat{f}^{*b}(x)$
- Return average of predictions (regression)

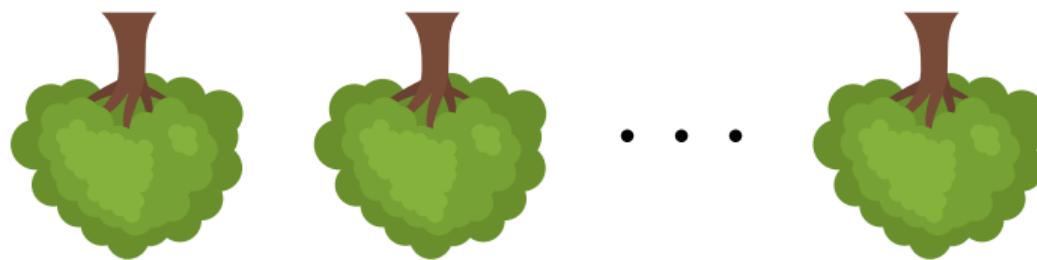
$$\hat{f}_{bag}(x) = \frac{1}{B} \sum_{b=1}^B \hat{f}^{*b}(x)$$

or majority vote (classification)

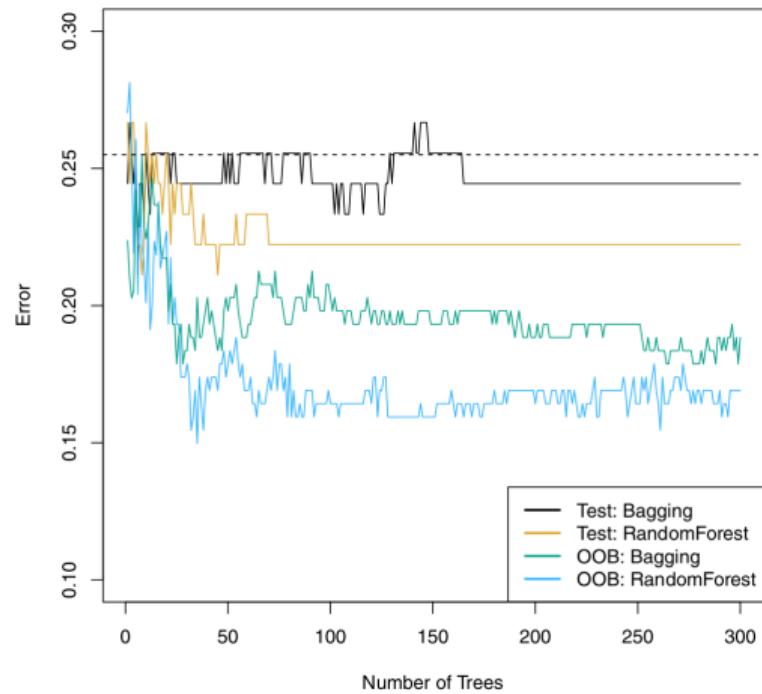
## Tree version



## Prediction on new data point

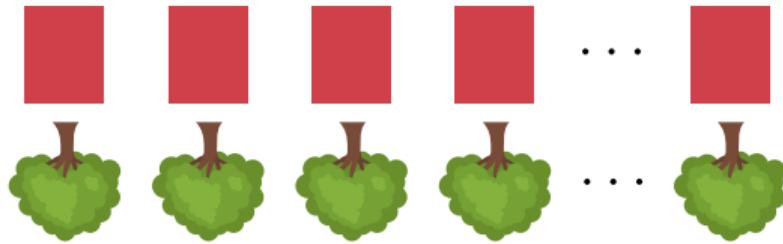


## Example: Heart classification data

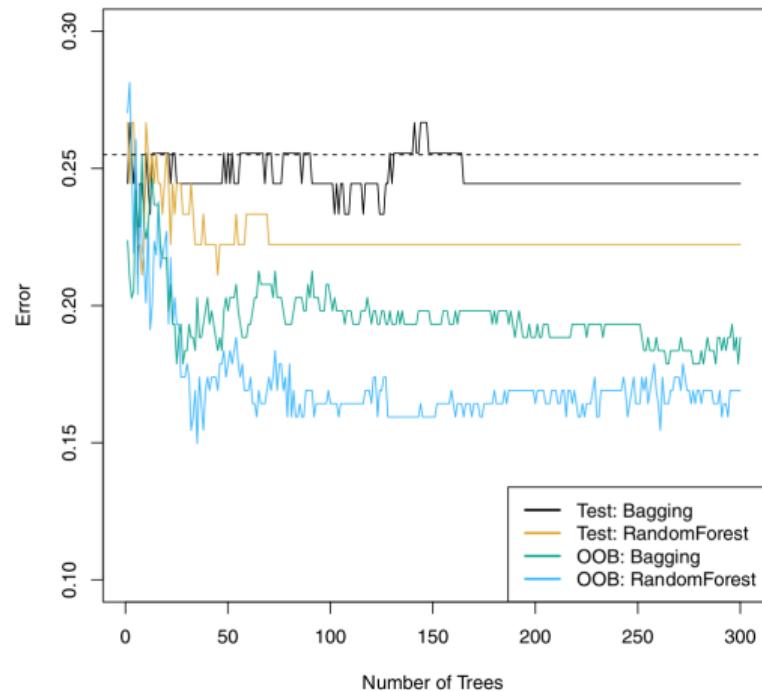


# Out of Bag Error Estimation

- On average, bootstrap sample uses about 2/3 of the data
- Remaining observations not used are called *out-of-bag* (OOB) observations
- For each observation, run through all the trees where it wasn't used for building
- Return the average (or majority vote) of those as test prediction! rather law of large number
- Bootstrapped version of LOOCV.



# Error using OOB



Test your understanding: [PolIEv](#)

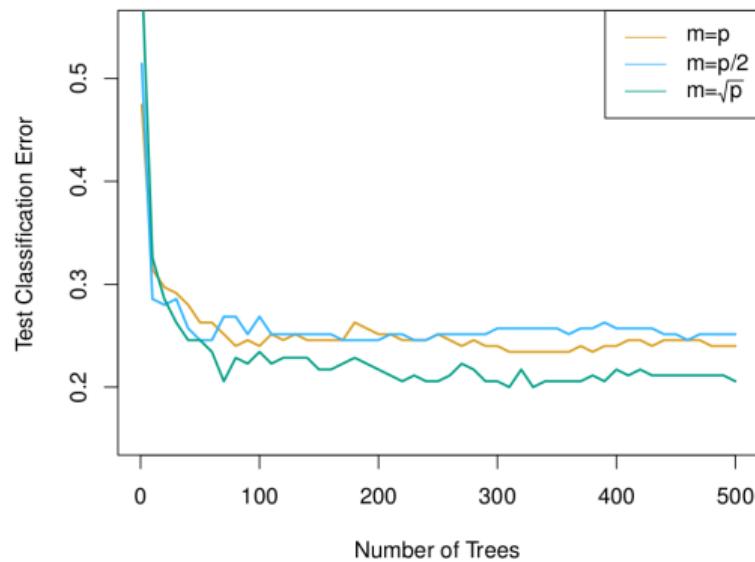
## Section 3

### Random Forests

# The idea

- Goal is to decorrelate the bagged trees:
  - ▶ If there is a strong predictor, the first split of most trees will be the same
  - ▶ Most or all trees will be highly correlated
  - ▶ Averaging highly correlated quantities doesn't decrease variance as much as uncorrelated
- The random forest fix:
  - ▶ Each time a split is considered, only use a random subset of  $m$  the predictors
  - ▶ Fresh sample taken every time
  - ▶ Typically  $m \approx \sqrt{p}$
  - ▶ On average,  $(p - m)/p$  of splits won't consider strong predictor
  - ▶  $m = p$  gives back bagging

## Example on gene expression



# Coding time!

- Bagging: trees grown independently on random samples. Trees tend to be similar to each other, can result in getting caught in local optima
- Random forest: trees independently on samples, but split is done using random subset of features

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