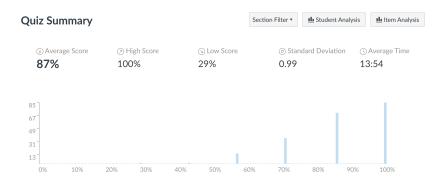
S&DS 265 / 565 Introductory Machine Learning

### **Trees and Forests**

October 3

### Reminders

- Assn 2 out; due Thursday at midnight
- Quiz 2 last week
- Midterm in class on Tuesday, October 17
- Questions?



### Communication

*Important*: For any communication regarding assignments, late work, etc. that requires email:

- Email the instructor John Lafferty (john.lafferty@yale.edu) and course manager Joanne Chen (haoting.chen@yale.edu)
- State that this is for Introductory Machine Learning (S&DS 265)
- Any email to me alone may go unanswered

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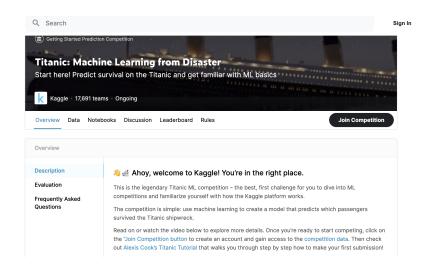
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- Response variables can be categorical or quantitative
- Yields a set of interpretable decision rules
- Predictive ability is mediocre, but can be improved by combining multiple trees (resampling, ensemble methods)

#### Titanic data



#### Titanic data

- Survived: Outcome of survival (0 = No; 1 = Yes)
- Pclass: Socio-economic class (1 = Upper class; 2 = Middle class; 3 = Lower class)
- · Name: Name of passenger
- · Sex: Sex of the passenger
- . Age: Age of the passenger (Some entries contain NaN)
- . SibSp: Number of siblings and spouses of the passenger aboard
- · Parch: Number of parents and children of the passenger aboard
- · Ticket: Ticket number of the passenger
- · Fare: Fare paid by the passenger
- . Cabin Cabin number of the passenger (Some entries contain NaN)
- Embarked: Port of embarkation of the passenger (C = Cherbourg; Q = Queenstown; S = Southampton)

### **Trees**



8

### **Trees**

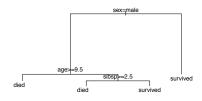


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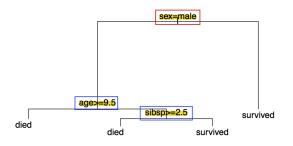
### **Trees**



### Modeling Titanic survival:



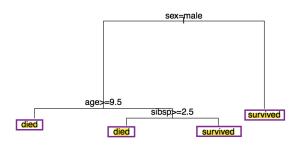
**Internal nodes** are points where the predictor space is split.



The internal node at the top is the **root** of the tree.

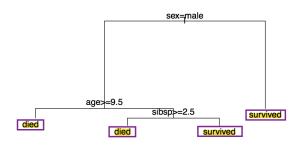
9

**Terminal nodes** (or **leaves**) are the ends of the tree where no further splitting occurs.



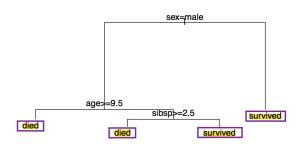
C

**Terminal nodes** (or **leaves**) are the ends of the tree where no further splitting occurs.



Denote these *J* regions as  $R_1, \ldots, R_J$ .

c



```
    R<sub>1</sub> = {i : sex<sub>i</sub> = male ∩ age<sub>i</sub> ≥ 9.5}
    R<sub>2</sub> = {i : sex<sub>i</sub> = male ∩ age<sub>i</sub> < 9.5 ∩ sibsp<sub>i</sub> ≥ 2.5}
    R<sub>3</sub> = {i : sex<sub>i</sub> = male ∩ age<sub>i</sub> < 9.5 ∩ sibsp<sub>i</sub> < 2.5}</li>
    R<sub>4</sub> = {i : sex<sub>i</sub> ≠ male}
```

9

# Let's go to the Titanic demo

#### **Bias-variance**

- Nodes are split by <u>greedily</u> choosing the best question (greatest reduction in error)
- As tree is grown deeper, bias decreases
- But the variance increases
- How to choose the right size of tree?

Once we stop, we relabel the terminal nodes to be  $R_1, \ldots, R_J$  and compute  $\bar{y}_{R_i}$  (means within each region) to serve as  $\hat{y}$  values.

But when do we stop?

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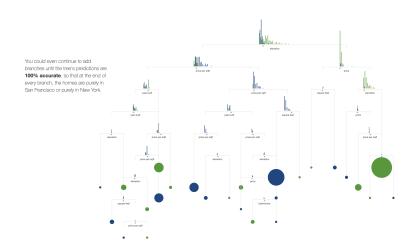
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Many options – resulting in tuning parameters that are hard to deal with.

Another way to get around the overfitting problem is to grow a large tree and then **prune** it back.

Typically, pruning involves looking at subtrees of the fully-grown tree, and comparing how well the subtrees perform.



How do we prune?

- cross validation
- cost-complexity pruning

## **Cost-complexity pruning**

Minimize: Loss(
$$T$$
) +  $\lambda$  {# of nodes in T}
$$= \sum_{m=1}^{|T|} \sum_{i \in R_m} (y_i - \widehat{y}_{R_m})^2 + \lambda |T|$$

 $\lambda$  is a tuning parameter that controls for the complexity of the model.

# **Cost-complexity pruning**

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•  $\lambda = 0$  implies the full tree

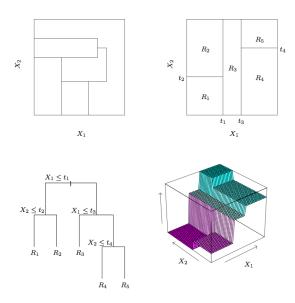
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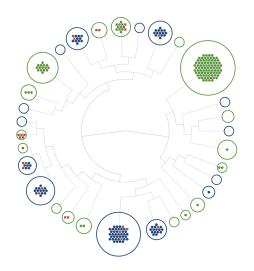
 $\lambda$  is a tuning parameter that controls for the complexity of the model.

- $\lambda = 0$  implies the full tree
- Larger  $\lambda$  implies higher penalty for complexity of model

- Grow a big tree on a training set.
- ② Obtain a nested set of subtrees T<sub>L</sub> ⊂ · · · ⊂ T<sub>2</sub> ⊂ T<sub>1</sub> ⊂ T<sub>0</sub> corresponding to a sequence of λ values.
- 3 Use (leave-one-out or k-fold) cross-validation to identify the subtree/ $\lambda$  that does best.



### Beautiful demo http://www.r2d3.us/



This demo gives a nice description of bias-variance tradeoff for trees. It's well worth a look!

### Trees vs. other methods

Decision trees are similar in spirit to *k*-nearest neighbors.

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 Both produce simple predictions (averages/maximally occurring) based on "neighborhoods" in the predictor space.

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- Both produce simple predictions (averages/maximally occurring) based on "neighborhoods" in the predictor space.
- Neighborhoods chosen very differently

Recall that linear regression fits models of the form

$$f(X) = \beta_0 + \sum_{j=1}^{p} X_j \beta_j$$

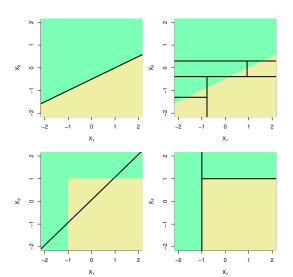
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$$f(X) = \beta_0 + \sum_{j=1}^{p} X_j \beta_j$$

Regression trees are like fitting linear regression models with a bunch of indicators

$$f(X) = \sum_{j=1}^{J} \beta_j \mathbb{1} \left\{ X \in R_j \right\}$$

Are trees always better than linear methods?



#### Summary so far

- Trees give interpretable, nonlinear prediction rules
- Deep trees have low bias, high variance
- Shallow trees have high bias, low variance
- Deep trees are pruned back using cross-validation to find best bias/variance tradeoff.

#### **Random Forests**

- Grow many trees and average their predictions
- Trees are grown deep, to have low bias, but high variance
- To "decorrelate" the trees and reduce variance, each tree is
  - grown on a bootstrap sample of the data
  - grown with random subsets of the predictors at each split
- Tree growing can be done in parallel

# Leo Breiman—"Keep it simple"



# **Random Forests Algorithm**

- **1** For b = 1 to B:
  - (a) Draw a bootstrap sample  $Z^*$  of size n from the training data
  - (b) Grow a random-forest tree T<sub>b</sub> to the bootstrapped data, recursively repeating following steps, until minimum node size reached:
    - i. Select *m* variables at random from the *p* variables
    - ii. Pick the best variable/split-point among the m
    - iii. Split the node into two children nodes
- ② Output the ensemble of trees  $\{T_b\}_{b=1}^B$ .

#### **Random Forests Algorithm**

To make a prediction at a new point *x*:

Regression: Average  $\hat{f}_{rf}^B(x) = \frac{1}{B} \sum_{b=1}^{B} T_b(x)$ 

Classification: Majority vote of the individual trees

# Out of bag (OOB) prediction

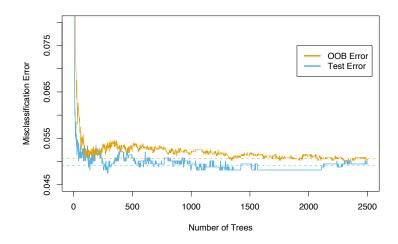
A bootstrap sample, or "bag" of data, is a set of n data points sampled with replacement from the original set of n data points. Will contain repetitions.

Each tree is grown on such a sample, which contains about  $\frac{2}{3}$  of the original data (with repetitions). The remaining  $\frac{1}{3}$  can be used as validation data

- For each observation  $z_i = (x_i, y_i)$ , construct its random forest predictor by averaging only those trees corresponding to bootstrap samples in which  $z_i$  did not appear
- Thus, cross-validation can be performed "along the way"

Chance a sample  $x_i$  does not appear in a bootstrap sample is  $\left(1-\frac{1}{n}\right)^n \longrightarrow \frac{1}{e} \approx 0.37$ 

# Out of bag (OOB) prediction



# Performance on email spam task

Single tree: 8.7%

Random forest: 5.1%

(standard error of the estimates is  $\approx 0.6\%$ )

Random forests improve upon the predictive ability of trees, but this sacrifices the interpretability of the single tree model

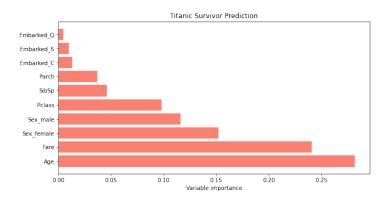
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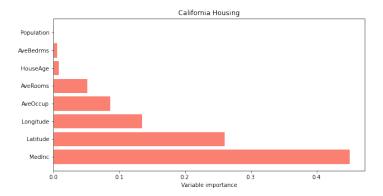
A good tool for interpreting a forest is **variable importance** 

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A good tool for interpreting a forest is variable importance

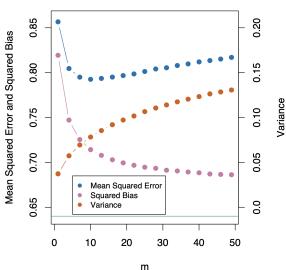
Variable importance is the amount that the RSS (or other loss) is reduced due to splits over a given predictor, averaged over all trees in the forest





#### **Random forest MSE**





# Understanding the bias-variance tradeoff for random forests

- Each tree has low (squared) bias, because it is deep
- The bootstrap sample and random subset of questions allowed at a given split result in "diversity" among of the trees
- This diversity translates to decorrelation, and tends to reduce the variance
- As we increase the number m in the random subset of questions allowed at each split, the complexity increases—variance goes up, squared bias goes down

#### Let's go to the notebook

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
```

In [2]: titanic\_train = pd.read\_csv('https://raw.githubusercontent.com/minsuk-heo/kaggle-titanic/master/input/train.csv')
titanic\_test = pd.read\_csv('https://raw.githubusercontent.com/minsuk-heo/kaggle-titanic/master/input/test.csv')
titanic\_train

#### Out[2]:

		Passengerld	Survived	Pclass	Name	Sex	Age	SibSp	Parch	Ticket	Fare	Cabin	Embarked
	0	1	0	3	Braund, Mr. Owen Harris	male	22.0	1	0	A/5 21171	7.2500	NaN	s
	1	2	1	1	Curnings, Mrs. John Bradley (Florence Briggs Th	female	38.0	1	0	PC 17599	71.2833	C85	С
	2	3	1	3	Heikkinen, Miss. Laina	female	26.0	0	0	STON/02. 3101282	7.9250	NaN	s
	3	4	1	1	Futrelle, Mrs. Jacques Heath (Lily May Peel)	female	35.0	1	0	113803	53.1000	C123	s
	4	5	0	3	Allen, Mr. William Henry	male	35.0	0	0	373450	8.0500	NaN	s
8	86	887	0	2	Montvila, Rev. Juozas	male	27.0	0	0	211536	13.0000	NaN	s
8	87	888	1	1	Graham, Miss. Margaret Edith	female	19.0	0	0	112053	30.0000	B42	s
8	88	889	0	3	Johnston, Miss. Catherine Helen "Carrie"	female	NaN	1	2	W./C. 6607	23.4500	NaN	s
8	89	890	1	1	Behr, Mr. Karl Howell	male	26.0	0	0	111369	30.0000	C148	С
8	90	891	0	3	Dooley, Mr. Patrick	male	32.0	0	0	370376	7.7500	NaN	Q

891 rows x 12 columns

#### **Summary**

- Trees give interpretable, nonlinear prediction rules
- Deep trees have low bias, high variance
- Random forests are a way of combining trees
- Want: Different trees should capture different aspects of the data
- How: Grow each tree on a random (bootstrap) sample of the data and choosing from a random set of questions at each split