INFO 251: Applied Machine Learning

Trees and Forests



Thanks to machine-learning algorithms, the robot apocalypse was short-lived.

Key Concepts (last lecture)

- Churn prediction
- Decision boundaries
- Hyper-rectangles
- Splitting
- Information gain
- Recursive tree building

Intuition check

- True or false:
 - The recursive decision tree algorithm is deterministic (assuming we are not performing cross-validation and have fixed the hyperparameters in advance)

Course Outline

- Causal Inference and Research Design
 - Experimental methods
 - Non-experiment methods
- Machine Learning
 - Design of Machine Learning Experiments
 - Linear Models and Gradient Descent
 - Non-linear models
 - Neural models
 - Unsupervised Learning
 - Practicalities, Fairness, Bias
- Special topics

Outline

- Regression Trees
- Random Forests
- Feature Importance

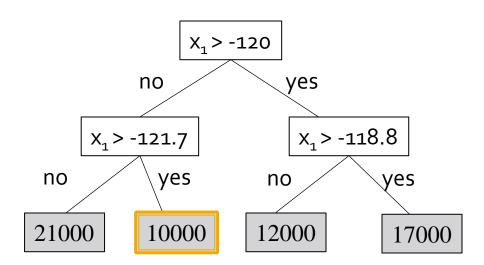
Regression Trees

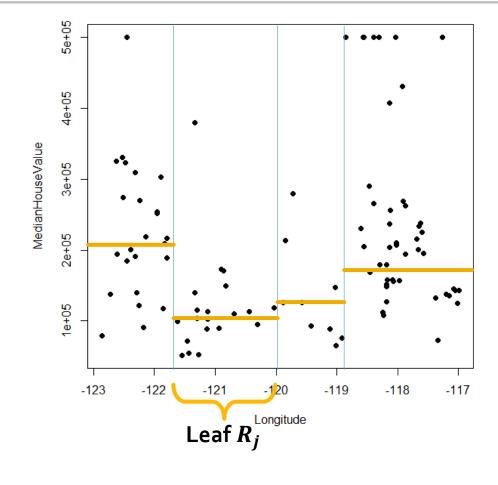
- What if output values are continuous or real-valued (i.e., not discrete)?
- Regression trees
 - Construct binary tree, minimize error in each leaf
 - Before, we counted # elements of each type in leaf
 - Now we choose predicted value that minimizes error
- Example: Predict median housing value based on a house's location (latitude, longitude)

Regression Trees

> head(calif[,c(1,8,9)])

	MedianHouseValue	Latitude	Longitude
1	452600	37.88	-122.23
2	358500	37.86	-122.22
3	352100	37.85	-122.24
4	341300	37.85	-122.25
5	342200	37.85	-122.25
6	269700	37.85	-122.25



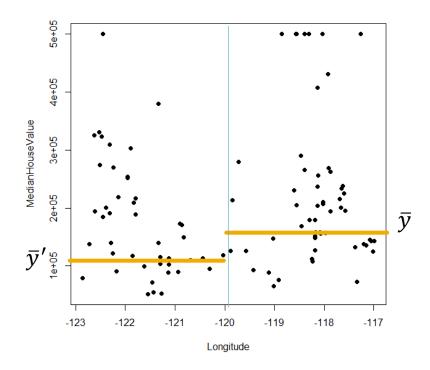


Regression Trees

- How to choose split point?
 - Idea: Minimize prediction error
- In 1-dimension: choose s to minimize

$$\min_{\bar{y}} \sum_{i:x_i > s} (\bar{y} - y_i)^2 + \min_{\bar{y}'} \sum_{i:x_i \le s} (\bar{y}' - y_i)^2$$

Consider finite splits (e.g. s between data)

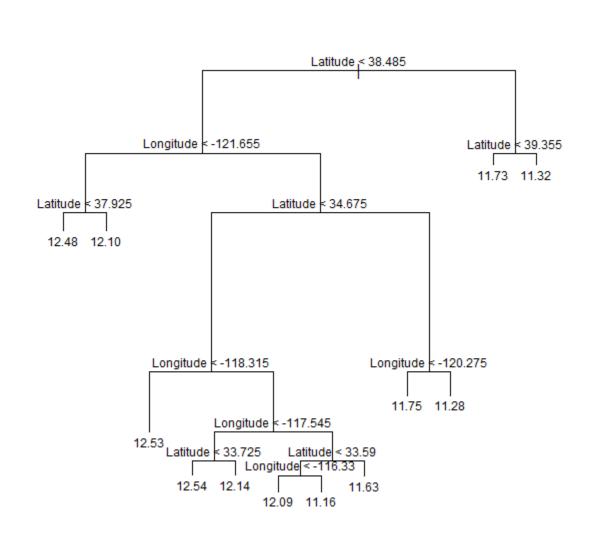


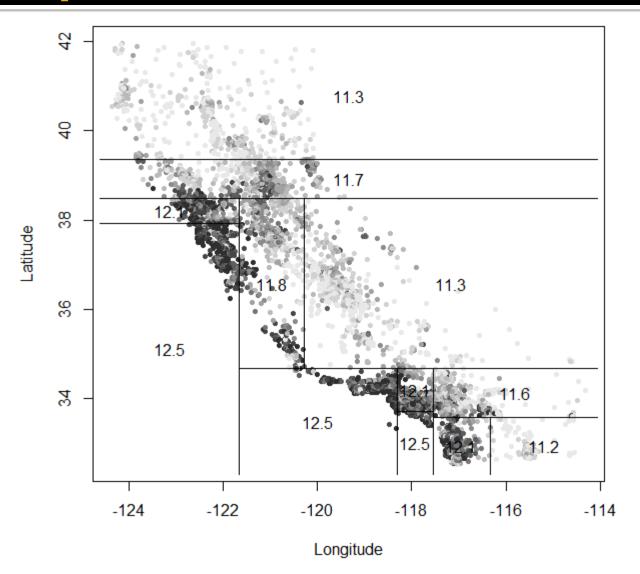
This intuition generalizes trivially to D dimensions

Regression Trees: Recursive Algorithm

- 1. Start with a single node (c_i) containing all points.
 - 1. Calculate predicted value $\bar{y}_{c_j} = \frac{1}{n} \sum_{i \in c_j} y_i$
 - 2. Calculate total error: $J = \sum_{c_j} \sum_{i \in c_j} \left(\overline{y}_{c_j} y_i \right)^2$
- If all points in the node have the identical features, stop.
 - Otherwise, search all binary splits of all variables for split that most reduces J
 - Stop if J decreases less than δ or if nodes are close to empty
 - Otherwise, make that split, creating two new nodes
- 3. Recurse on each new node

Regression Trees: Example



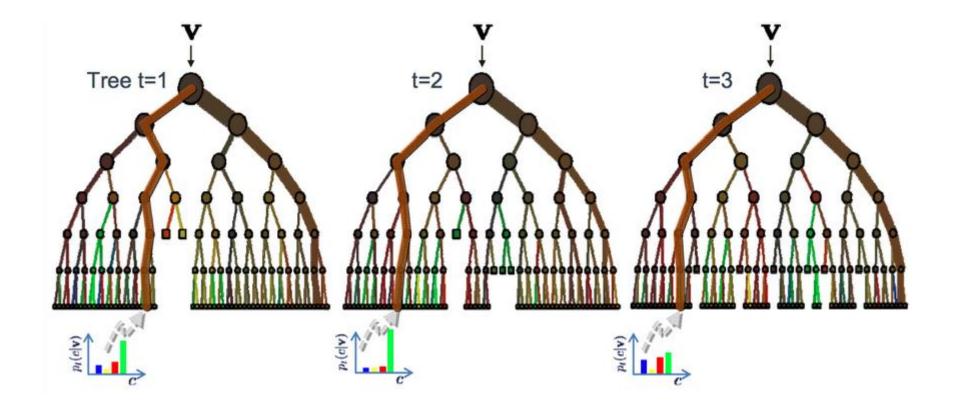


Outline

- Regression Trees
- Random Forests
- Feature Importance

Trees to forests

- Which classifier works best?
 - "Random forests" combine outputs of multiple classifiers



Building a forest

- Bootstrap sample a new training set
 - with replacement
- Build a decision tree
 - Random subset of splits/features
 - Forces differentiated trees
 - No pruning! "Regularization" through forest
- Repeat until you have lots of trees
- Predict by taking a vote among the trees

Example: The "CART" forest

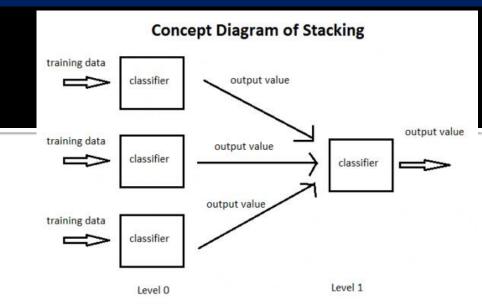
Formally:

$$\widehat{y_i} = \sum_{f_k \in \mathcal{F}} f_k(x_i)$$

- ullet ${\mathcal F}$ is the space of regression trees
 - Each f_k maps data examples x_i to tree leaves
 - With CART (a common decision tree algorithm), each leaf isn't a decision value, it's a "score"
 - Scores are summed across trees

Ensemble methods

- Bagging = bootstrap aggregating
 - Create artificial versions of data via bootstrap
 - 1 sample = bootstrap
 - M samples = bagging
- Stacking: train model (e.g. another tree, a logistic regression) on output of other models
- Boosting: Can a set of weak learners create a single strong learner?
 (Kearns, 1988)
 - Train a sequence of models, each emphasizes the examples misclassified by the previous model



Adaptive Boosting (AdaBoost)

- Adaboost:
- 1. Initially, set a weight for each training example = 1/n
- 2. Train a classifier where the objective respects the weights
- 3. Increase the weights for misclassified examples
- 4. Return to 2

Gradient Boosting



http://blog.kaggle.com/2017/01/23/a-kaggle-master-explains-gradient-boosting/

(Extreme) Gradient Boosting

Start with regression tree-based model:

$$\widehat{y}_i = \sum_{f_k \in \mathcal{F}} f_k(x_i)$$

Gradient boosting loss function "fits on residuals":

$$\mathcal{L}^{(t)} = \sum_{i=1}^{n} l(y_i, \hat{y}_i^{(t-1)} + f_t(x_i)) + \gamma T + \frac{1}{2} \lambda ||w||^2$$

$$f_t \text{ fits on residual of t-1} \qquad \text{Regularization penalty}$$

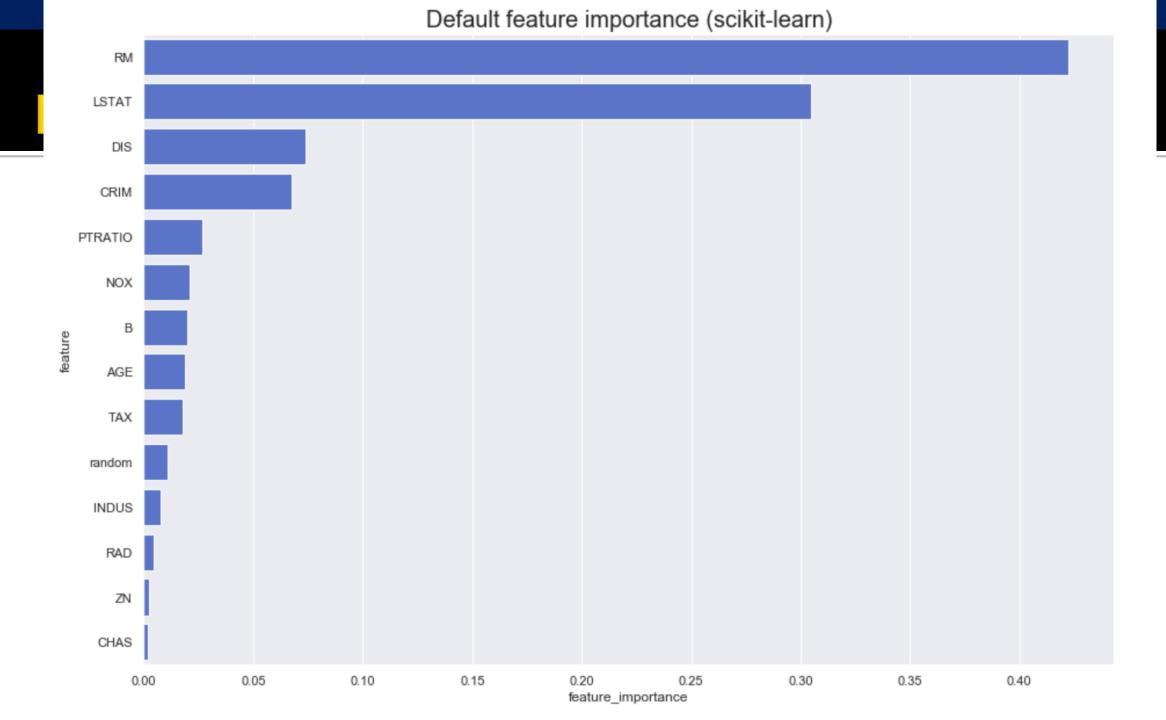
- T is the number of leaves
- t indexes training iterations
- w is vector of scores on each leaf (i.e., the leaf weights)
- Optimization is similar to gradient descent
 - Relies on being able to measure how good each tree is
 - Next tree solves for the loss of prior tree

Outline

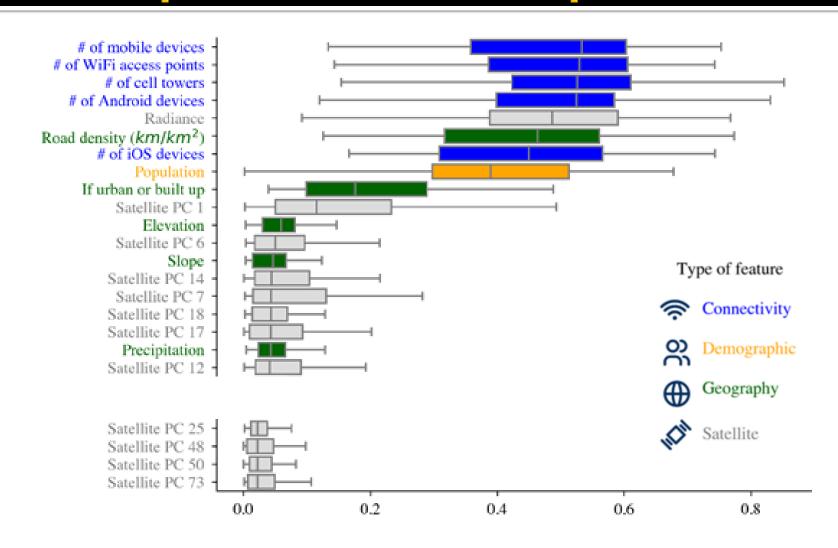
- Decision Trees: Loose ends
 - Overfitting and Pruning
 - Extensions
- Regression Trees
- Random Forests
- Feature Importance

- The primary focus of random forests is prediction, rather than inference
 - A single tree is fairly interpretable, but it's hard to interpret a forest
 - This is generally true for complex, non-parametric, and non-linear models
- Nonetheless, people frequently still want to do some ex-post interpretation
 - "What features are important to the classifier?"

- Intuitively, the features that "matter":
 - Occur high in the tree (high information gain for that tree)
 - Occur frequently in the tree (if feature is non-binary)
 - Occur in many trees (if it's a forest)



Feature Importance: Example



- Formally, two common approaches:
 - Mean decrease impurity (aka Gini importance): average (across trees) decrease in weighted impurity caused by that feature

```
from sklearn.datasets import load boston
from sklearn.ensemble import RandomForestRegressor
 3 import numpy as np
   #Load boston housing dataset as an example
 5 boston = load boston()
 6  X = boston["data"]
   Y = boston["target"]
8    names = boston["feature names"]
9    rf = RandomForestRegressor()
    rf.fit(X, Y)
     print "Features sorted by their score:"
     print sorted(zip(map(lambda x: round(x, 4), rf.feature importances ), names),
13
                   reverse=True)
 Features sorted by their score:
  [(0.5298, 'LSTAT'), (0.4116, 'RM'), (0.0252, 'DIS'), (0.0172, 'CRIM'), (0.0065, 'NOX'),
  (0.0035, 'PTRATIO'), (0.0021, 'TAX'), (0.0017, 'AGE'), (0.0012, 'B'), (0.0008, 'INDUS'),
  (0.0004, 'RAD'), (0.0001, 'CHAS'), (0.0, 'ZN')]
```

- Issues with impurity:
 - Biased towards features with multiple values
 - What happens when two features are closely correlated?

```
size = 10000
np.random.seed(seed=10)
X_seed = np.random.normal(0, 1, size)
X0 = X_seed + np.random.normal(0, .1, size)
X1 = X_seed + np.random.normal(0, .1, size)
X2 = X_seed + np.random.normal(0, .1, size)
X = np.array([X0, X1, X2]).T
Y = X0 + X1 + X2

rf = RandomForestRegressor(n_estimators=20, max_features=2)
rf.fit(X, Y);
print "Scores for X0, X1, X2:", map(lambda x:round (x,3), rf.feature_importances_)
```

Scores for X0, X1, X2: [0.278, 0.66, 0.062]

- Formally, two common approaches:
 - Mean decrease impurity: average (across trees) decrease in weighted impurity caused by that feature
 - 2. Mean decrease accuracy ("Permutation Importance"): average (across trees) decrease in performance when a given feature is randomized
 - Not implemented in sklearn (but very easy to do so by hand see ESLII reading)

- Issues
 - Interpret feature importances at your own risk!
 - They are informative, but rather atheoretical

Key Concepts (this lecture)

- Overfitting and pruning
- Regression trees
- Random forests
- AdaBoost
- Gradient boosting
- Feature Importance

Recap

- Regression
 - Parametric, fast training, linear
- Nearest Neighbors
 - Non-parametric, no training, complex decisions
- Naïve Bayes
 - Parametric, very fast training
- Decision Trees
 - Non-linear decisions, intuitive model

For Next Class:

- Read:
 - Daume, chapters 4 and 10
- Keep working on Problem Set 4!

