Random Forests

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Outline

- Growing Random Forests
- Parameters
- Results
- (Neural Networks)

Random Forests

- Breiman (2001), Breiman & Cutler (2004)
- Tree Ensemble built by randomly sampling cases and variables
- Each case classified once for each tree in the ensemble

How do Random Forests work

- Large number (at least 500) of 'different' trees is grown
- Each tree gives a classification for each record, i.e. the tree "votes" for that class.
- The forest determines the overall classification for each record by a majority vote.

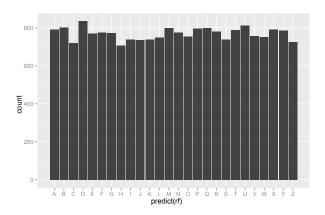
Growing a Random Forest

for sample size N and M explanatory variables $X_1, ..., X_M$

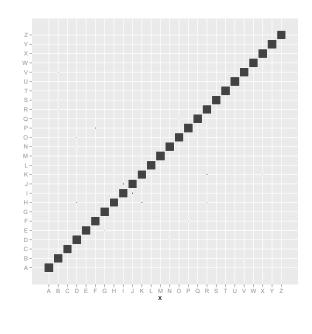
- draw bootstrap sample of data (i.e. draw sample of size N with replacement)
- at each node, select m << M variables at random and find best split.
- each tree is grown to the largest extent possible, i.e. no pruning!

randomForest package

Results



Misclassification

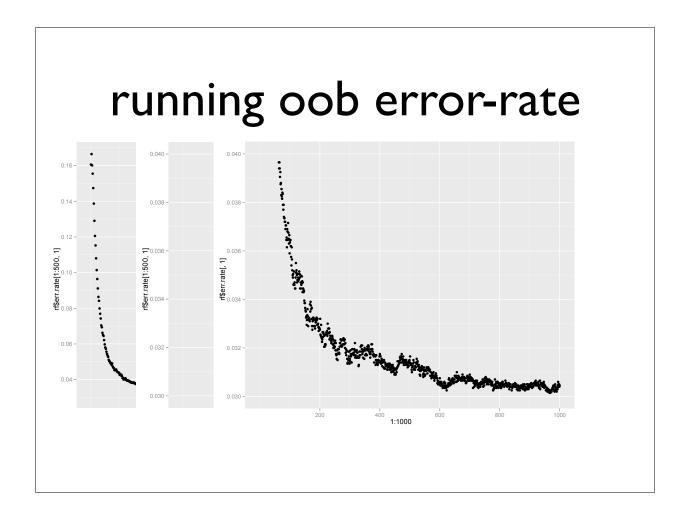


Forest Error

- Increasing correlation between any two trees increases the forest error rate.
- Trees with low individual error rates are stronger classifiers. Increasing strength of individual trees decreases the overall forest error rate.
- decreased m reduces both correlation and strength. "optimal" range of m usually quite wide.

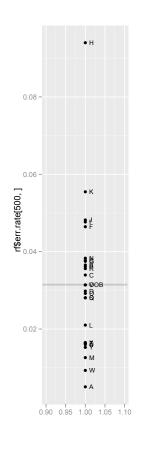
Out of bag (oob) error

- Slight modification to bootstrap samples:
- for each tree, leave about 1/3 of data out of sample, then draw bootstrap sample of size N.
- use out-of bag data to get (running)
 unbiased estimate of classification error as
 each tree is added to forest.



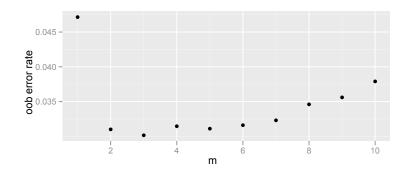


 oob classification allows to assess error rates for each class



optimal choice of m

based on oob error:



sqrt(M) works well in most cases

Variable Importance

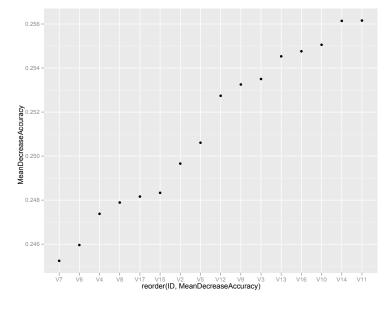
Permutation Criterion:

- based on out-of bag data
- for each tree, count # of correctly classified oob records
- permute values of variable m, re-count correctly classified oob records, subtract from first count
- for each variable, average over all trees





Mean Decrease Accuracy



Proximity

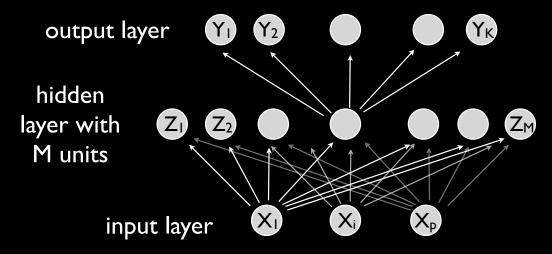
- N x N matrix of proximity values
- for each tree: if two records k and l are in the same leaf, increase proximity by one
- normalize proximity by dividing by number of trees
- size problematic

Neural Networks

- Historically used to model (biological) networks of neurons:
 - nodes represent neurons
 - edges represent nerves
 - network illustrates activity and flow of signals

Setup

- Response Y has K categories
- Network:



Formula Setup

• Relationship between layers:

$$Z_m = \sigma(\alpha_{0m} + \alpha'_m X)$$
 $m = 1, ..., M$
 $T_k = \beta_{0k} + \beta'_k Z$ $k = 1, ..., K$
 $f_k(X) = g_k(T)$ $k = 1, ..., K$

• where sigma is the activation function, e.g.

$$\sigma(\nu) = \frac{1}{1 + e^{-\nu}}$$

• gk is final transformation between T and Y

Formula Setup

 g_k with continuous response usually chosen as identity, with categorical response usually softmax:

$$g_k(T) = \frac{e^{T_k}}{\sum_{\ell=1}^K e^{T_\ell}}$$

• i.e. estimates are positive and sum to 1

Issues with Neural Nets

Model generally highly over-parametrized:

weights:
$$\begin{cases} \{\alpha_{0m}, \alpha_m : m = 1, ..., M\} & M(p+1) \\ \{\beta_{0k}, \beta_k : k = 1, ..., K\} & K(M+1) \end{cases}$$

- Optimization problem convex & unstable -> convergence is tricky
- Over-parametrization leads to overfit at minimum

Fitting Strategies

- Standardize input variables X
- Pick starting values for alpha, beta close to zero (i.e. close to linear fit)
- Stop run before convergence (to avoid overfitting)
- Alternatively: use penalty on size of weights (decay)

$$\lambda \cdot \left(\sum \beta^2 + \sum \alpha^2\right)$$

Fitting Strategies

- Pick large number of hidden units OR do cross-validation to figure out good size
- # parameters (and with it #units) bounded by sample size
- average results from set of networks (bagging)

Neural Networks are fickle

Choice of M is important

starting parameters
are important some models do not
even come close to
a good solution in
100 iterations

