

Introduction to Machine Learning

Chapter 16: Random Forest cont.

Bernd Bischl, Christoph Molnar

Department of Statistics – LMU Munich Winter term 2017/18



- Single trees are highly interpretable
- Random Forests as an ensemble of many trees lose this feature
- Hence, contributions of a single covariate to the fit are difficult to evaluate
- Way out: variable importance measures

Measure based on permutations of OOB observations

- 1: After growing tree $\hat{b}^{[m]}(x)$, pass down OOB observations and record predictive accuracy.
- 2: Permute OOB observations of *j*th variable.
- 3: Pass down the permuted OOB observations and evaluate predictive accuracy again.
- 4: The loss of goodness induced by permutation is averaged over all trees and is used as a measure for the importance of the *j*th variable.

Measure based on improvement in split criterion

- 1: At each split in tree $\hat{b}^{[m]}(x)$ the improvement in the split criterion is attributed as variable importance measure for the splitting variable.
- 2: For each variable, this improvement is accumulated over all trees for the importance measure.

VARIABLE IMPORTANCE BASED ON PERMUTATIONS OF OOB OBSERVATIONS





Tree M

Inbag + oob of tree 1 Permuted oob obs. of x_1

| | X ₁ | | Xp | у | ŷ |
|---|----------------|-----|-----|---|---|
| 1 | 1.4 | | | 1 | |
| 2 | 2 | | | 0 | |
| 3 | 1.55 | 9 | 13 | 1 | 1 |
| 4 | 1.72 | 230 | 200 | 0 | 0 |
| 5 | 1.89 | | | 1 | |
| : | | | | | |
| n | 2.01 | 1 | 12 | 1 | 1 |



| | X ₁ | | Xp | y | ŷ |
|---|----------------|-----|----|---|---|
| 1 | 1.4 | | 1 | 1 | 1 |
| 2 | 2 | | | 0 | |
| 3 | 1.55 | 911 | 73 | 1 | 0 |
| 4 | 1.72 | | | 0 | |
| 5 | 1.89 | 23 | 6 | 1 | 1 |
| 1 | | | | | |
| n | 2.01 | | | 1 | |

| | X ₁ | | Xp | y | ŷ |
|---|----------------|----|-----|---|---|
| 1 | 1.89 | | 1 | 1 | 0 |
| 2 | 2 | | | 0 | |
| 3 | 1.4 | 38 | 13 | 1 | 0 |
| 4 | 1.72 | | | 0 | |
| 5 | 1.55 | 23 | 100 | 1 | 1 |
| i | | | | | |
| n | 2.01 | | | 1 | |

$$acc_{1, \text{ without permutation}} - acc_{1, \text{ with permutation}} = diff_1$$

$$acc_{1, \text{ without permutation}} - acc_{1, \text{ with permutation}} = diff_1$$
 $acc_{M, \text{ without permutation}} - acc_{M, \text{ with permutation}} = diff_M$

$$\frac{1}{M}\sum_{i=1}^{M} diff_i$$
 = variable importance for x_1



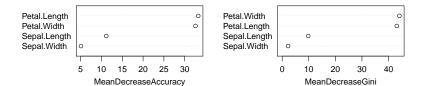


Figure: Two importance measures on iris.

iris_example (4 features)

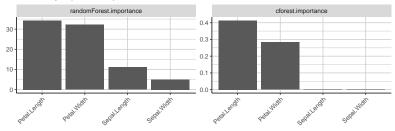


Figure: RF importance as filters in mlr.

RANDOM FOREST: PROXIMITIES

- the "closeness" or "nearness" between pairs of cases.
- Algorithm
 - After a tree is grown, put all of the data down the tree.
 - If cases x_1 and x_2 are in the same terminal node through one tree increase their proximity by one.
 - At the end of the run of all trees, normalize the proximities by dividing by the number of trees.
- The proximities originally form a NxN matrix.
- Proximities are used in replacing missing data, locating outliers, and producing illuminating low-dimensional views of the data.

We can visualize our Proximities $P(x_i, x_j)$ for each $i \in \{1, ..., n\}$ example by MDS.

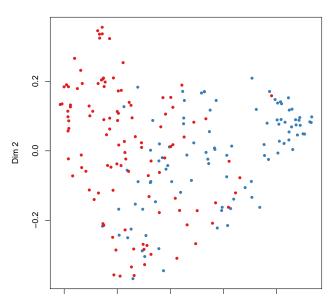
Our data contains 111 patterns obtained by bouncing sonar signals off a metal cylinder at various angles and under various conditions and 97 patterns obtained from rocks under similar conditions.

| ## | | V53 | V54 | V55 | V56 | V57 | V58 | V59 | V60 | Class |
|----|----|--------|--------|--------|--------|--------|--------|--------|--------|-------|
| ## | 1 | 0.0065 | 0.0159 | 0.0072 | 0.0167 | 0.0180 | 0.0084 | 0.0090 | 0.0032 | R |
| ## | 2 | 0.0089 | 0.0048 | 0.0094 | 0.0191 | 0.0140 | 0.0049 | 0.0052 | 0.0044 | R |
| ## | 3 | 0.0166 | 0.0095 | 0.0180 | 0.0244 | 0.0316 | 0.0164 | 0.0095 | 0.0078 | R |
| ## | 4 | 0.0036 | 0.0150 | 0.0085 | 0.0073 | 0.0050 | 0.0044 | 0.0040 | 0.0117 | R |
| ## | 5 | 0.0054 | 0.0105 | 0.0110 | 0.0015 | 0.0072 | 0.0048 | 0.0107 | 0.0094 | R |
| ## | 6 | 0.0014 | 0.0038 | 0.0013 | 0.0089 | 0.0057 | 0.0027 | 0.0051 | 0.0062 | R |
| ## | 7 | 0.0248 | 0.0131 | 0.0070 | 0.0138 | 0.0092 | 0.0143 | 0.0036 | 0.0103 | R |
| ## | 8 | 0.0120 | 0.0045 | 0.0121 | 0.0097 | 0.0085 | 0.0047 | 0.0048 | 0.0053 | R |
| ## | 9 | 0.0128 | 0.0145 | 0.0058 | 0.0049 | 0.0065 | 0.0093 | 0.0059 | 0.0022 | R |
| ## | 10 | 0.0223 | 0.0179 | 0.0084 | 0.0068 | 0.0032 | 0.0035 | 0.0056 | 0.0040 | R |

We try to predict the type, based on signals obtained from a variety of different aspect angles, spanning 90 degrees for the cylinder and 180 degrees for the rock

We calculate the proximities $P(x_i, x_j)$ based on out-of-bag observations.

Now we can visualize proximities $P(x_i, x_j)$ by MDS, using as distance matrix D= 1 - P



ADAPTIVE NEAREST NEIGHBORS

Let $P(x, x_i) \in [0, 1]$ be the Proximity between the observation x and our original point x_i .

- For classification, the prediction will be the weighted greatest number of hits, which is proportional to the proximities $P(x, x_i)$.
- For a regression we can calculate the prediction of Random Forests at x as:
 - if every leaf node contains the same number of observations.

$$\hat{Y}_{RF}(x) = \frac{\sum_{n=0}^{n} P(x, x_i) Y_i}{\sum_{n=0}^{n} P(x, x_i)}$$

• if some leaf node contains the different number of observations, $P(x, x_i)$ is the percentage of trees where x and x_i fall into the same leaf node, and weights are inversely proportional for each tree to the number of samples in the leaf node where x_i falls into.

RANDOM FOREST: ADVANTAGES

- Easy to implement
- Can be applied to basically any model
- Easy to parallelize
- Often works well (enough)
- Enables variance analysis
- Integrated estimation of OOB error
- Can work on high-dimensional data
- Often not much tuning necessary

RANDOM FOREST: DISADVANTAGES

- Often suboptimal for regression
- Hard to interpret, especially interactions
- Does not really optimize loss aggressively
- No real way to adapt to problem (see e.g. loss in GBM, kernel in SVM)
- Implementations sometimes memory-hungry
- Prediction can be slow