

Bagging (Bootstrap Aggregation)

- Training data: $\mathbf{Z} = \{(x_i, y_i)_{i=1}^n\}$
- Bootstrap samples^a: $\mathbf{Z}^{*b} = \{(x_i^{*b}, y_i^{*b})_{i=1}^n\}$, where $b = 1 : B$
- \hat{f}^{*b} : classification/regression function trained by \mathbf{Z}^{*b}
- The bagging estimate is defined to be

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

- **Advantage: reduce variance.** So works well for high-variance, low-bias procedures, such as trees.

^asample with replacement from \mathbf{Z} .

Random Forest

1. For $b = 1 : B$:
 - (a) Draw a bs sample \mathbf{Z}^{*b} from the training data.
 - (b) Grow a **BIG** tree T_b (with some restriction).
2. Output the forest $\{T_b\}_{b=1}^B$.

To make a prediction at a new point x

Regression: $\frac{1}{B} \sum T_b(x)$.

Classification: majority voting among $T_1(x), \dots, T_B(x)$.

Restriction when growing a tree in the forest:

- At each split, randomly select m variables from the p variables, and then pick the best split among them.
- The recommended value for m is \sqrt{p} for classification and $p/3$ for regression.
- Purpose: reduce the correlation between trees in the forest.

Out-of-Bag (OOB) Samples

- OOB samples: sample points which are not included in \mathbf{Z}^{*b} , i.e., they are not used in building the tree T_b
- The OOB samples can be used to get a test error for T_b .
- The prediction and error rate returned by randomForest are calculated based on OOB. The error is usually close to a CV error.

Variable Importance

- Calculation of the m -th variable's importance based on Gini Index: the improvement in the split-criterion (Gini index) is the importance measure attributed to the splitting variable, and is accumulated over all the trees in the forest separately for each variable.
- The calculation can be easily extended to regression trees based on MSE.

- Another measure is computed from **permuting** OOB samples: For each tree T_b in the forest, calculate the prediction error (error rate for classification, MSE for regression). Then the same is done after permuting the j th predictor for the OOB samples. The difference between the two (**before and after** permutation) is then averaged over all trees, and normalized by the corresponding standard deviation^a.
- R returns both scaled and unscaled variable importance.

^aIf the standard deviation of the differences is equal to 0 for a variable, then the division is not applied.