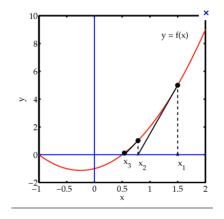
Committee Machines: Boosting Trees



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Outline

- Recap Adaboost
- Trees
- Boosting Trees
- Gradient Boosting motivation
- Steepest Descent
- Gradient Boosting algorithm
- Improving Gradient Boosting
- Interpretation (brief)
- Summary

AdaBoost.M1

- At each boosting step, weights are applied $w_1, w_2, ..., w_N$, to each of the training observations (x_i, y_i) , i = 1, 2, ..., N.
- At the initial step, $w_i = \frac{1}{N}$, $\forall i$.
- At each successive iteration, m = 2,3,...,M, the observation weights are individually modified and the classification algorithm is reapplied to the weighted observations.
- At step m, those observations that were misclassified by the classifier $G_{m-1}(x)$ induced at the previous step have their weights increased, whereas the weights are decreased for those that were classified correctly.

AdaBoost.M1

Algorithm 10.1 AdaBoost.M1.

- 1. Initialize the observation weights $w_i = 1/N, i = 1, 2, ..., N$.
- 2. For m=1 to M:
 - (a) Fit a classifier $G_m(x)$ to the training data using weights w_i .
 - (b) Compute

$$\operatorname{err}_{m} = \frac{\sum_{i=1}^{N} w_{i} I(y_{i} \neq G_{m}(x_{i}))}{\sum_{i=1}^{N} w_{i}}.$$

- (c) Compute $\alpha_m = \log((1 \operatorname{err}_m)/\operatorname{err}_m)$.
- (d) Set $w_i \leftarrow w_i \cdot \exp[\alpha_m \cdot I(y_i \neq G_m(x_i))], i = 1, 2, \dots, N$.
- 3. Output $G(x) = \operatorname{sign} \left[\sum_{m=1}^{M} \alpha_m G_m(x) \right]$.

AdaBoost.M1 aka "Discrete Adaboost", because returns a discrete class label. Real AdaBoost – specifically for real valued predictions.

Others include: gentle adaboost, modest adaboost.

Trees

- Recall: Regression and classification trees partition the model space into disjoint regions: $R_1, R_2, ..., R_J$, which are represented as the terminal nodes of the tree.
- A constant γ_i is assigned to each region, such that,

$$x \in R_j \Rightarrow f(x) = \gamma_j$$
.

Thus, a tree can be formally expressed as:

$$T(x;\Theta) = \sum_{j=1}^{J} \gamma_j I(x \in R_j),$$

where
$$\Theta = \left\{ R_j, \gamma_j \right\}_1^J$$
.

Solution requires numerical optimization:

$$\Theta = \arg\min_{\Theta} \sum_{j=1}^{J} \sum_{x_i \in R_i} L(y_i, \gamma_j).$$

Trees

2-stage solution:

- 1. Find γ_j given R_j : trivial, $\hat{\gamma}_j = \overline{y}_j$ by the modeling assumptions.
- 2. Find R_j : more complicated, use greedy, top-down recursive partitioning algorithm.

Recall: in two dimensions, splitting half planes and enumerating all pairs of split variables and split points.

Boosting Trees

The Boosted Tree:

$$f_M(x) = \sum_{m=1}^{N} T(x; \Theta_m),$$

Derived in a forward stagewise manner.

At each step, the minimization problem is solved:

$$\hat{\Theta}_m = \arg\min_{\Theta_m} \sum_{i=1}^N L(y_i, f_{m-1}(x_i) + T(x_i; \Theta_m)),$$

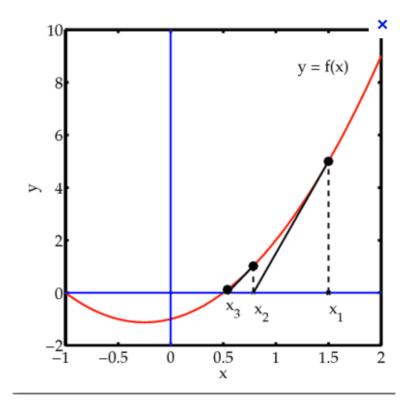
over regions and constants $\hat{\Theta}_m = \{R_{jm}, \gamma_{jm}\}_1^{J_m}$ of the next tree, given the current model $f_{m-1}(x)$.

 As before, given the regions finding the optimal constants in each region is straightforward:

$$\hat{\gamma}_{jm} = \arg\min_{\gamma_{jm}} \sum_{x_i \in R_{im}} L(y_i, f_{m-1}(x_i) + \gamma_{jm}),$$

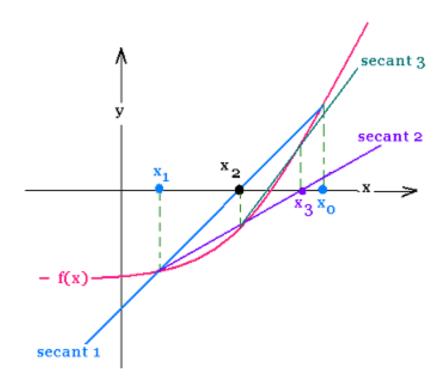
- General boosting algorithm that works well with a variety of different loss functions.
- Gradient boosting builds additive tree models.
- Tree size is the parameter that determines the order of interaction.
- Gradient boosting inherits all the good features of trees (variable selection, missing data, and mixed predictors), and improves on the weak features, such as prediction performance.

- What if we do not have a special case, and have no analytical solution - > optimization.
- Newton's method find a zero



What if derivatives are unavailable?

- In many practical application f'(x) is not given by a formula
- f'(x) may not exist over the domain.
- Quasi-Newton methods rely on f(x) only, to approximate the solution. In the simplest case, the secant method.



- Quasi-Newton's methods find a minimum, no requirement of hessian computation approximated by gradients.
- Consider a general loss:

$$L(f) = \sum_{i=1}^{N} L(y_i, f(x_i)).$$

the goal is to minimize the loss with respect to f(x).

Constrained to be the sum of trees

• Let the values of the approximating function f at each of the data points x_i be denoted as:

$$\mathbf{f} = \{f(x_1), f(x_2), \dots, f(x_N)\}.$$

• The numerical optimization problem:

$$\hat{\mathbf{f}} = \arg\min_{\mathbf{f}} L(\mathbf{f}).$$

where the "parameters" are $\hat{\mathbf{f}} \in \Re^N$.

• The solution can be expressed as a sum of component vectors:

• $\sum_{k=0}^{M} 1 = \sum_{k=0}^{M} N_k$

$$\mathbf{f}_{M} = \sum_{m=0}^{M} \mathbf{h}_{M}, \ \mathbf{h}_{M} \in \mathfrak{R}^{N},$$

where $\mathbf{f}_0 = \mathbf{h}_0$ is an initial guess and each successive \mathbf{f}_m is induced based on the current parameter vector \mathbf{f}_{m-1} .

Steepest Descent

 The solution is obtained via an iterative method, where the approximated function is updated repeatedly until convergence to the minimum. Updates are called "steps".

- How long should we step toward the minimum? is not a trivial question.
- What direction should we step? is also not a trivial question.
- The method of steepest descent steps as far as possible in the steepest direction.
- The negated gradient is the steepest downhill direction.

Steepest Descent

- The steepest descent chooses $\mathbf{h}_m = -\rho_m g_m$ where ρ_m is a scalar and $\mathbf{g}_m \in \mathfrak{R}^N$ is the gradient of $L(\mathbf{f})$ evaluated at $\mathbf{f} = \mathbf{f}_{m-1}$.
- The components of the gradient are:

$$g_{im} = \left[\frac{\partial L(y_i, f(x_i))}{\partial f(x_i)}\right]_{f(x_i) = f_{m-1}(x_i)}$$

and the step length ρ_m is the solution to:

$$\rho_m = \arg\min_{\rho} L(\mathbf{f}_{m-1} - \rho_m g_m).$$

• The update: $\mathbf{f}_m = \mathbf{f}_{m-1} - \rho_m g_m$.

 Steepest descent would be ideal if only concerned with minimizing the loss:

$$L(f) = \sum_{i=1}^{N} L(y_i, f(x_i)).$$

However, the gradient is only defined at training points x_i therefore, the approach may not be accurate to new data. The ultimate goal is to generalize $f_M(x)$ to new data.

• A resolution is to induce a tree $\{T(x_i;\Theta_m)\}$ as close as possible to the negative gradient,

$$\Theta_{m} = \arg\min_{\Theta} \sum_{i=1}^{N} \left(-g_{im} - T(x_{i}; \Theta) \right)^{2}$$

Note that the resulting regions will not be the same, but they will be close. The corresponding $\hat{\gamma}_{jm} = \arg\min_{\gamma_{jm}} \sum_{x \in \mathcal{R}} L(y_i, f_{m-1}(x_i) + \gamma_{jm})$,

TABLE 10.2. Gradients for commonly used loss functions.

Setting	Loss Function	$-\partial L(y_i, f(x_i))/\partial f(x_i)$
Regression	$\frac{1}{2}[y_i - f(x_i)]^2$	$y_i - f(x_i)$
Regression	$ y_i - f(x_i) $	$sign[y_i - f(x_i)]$
Regression	Huber	$y_i - f(x_i)$ for $ y_i - f(x_i) \le \delta_m$ $\delta_m \text{sign}[y_i - f(x_i)]$ for $ y_i - f(x_i) > \delta_m$ where $\delta_m = \alpha \text{th-quantile}\{ y_i - f(x_i) \}$
Classification	Deviance	kth component: $I(y_i = \mathcal{G}_k) - p_k(x_i)$

K least squares trees are constructed at each iteration. Each tree is fit to its corresponding gradient vector.

*Only one tree is needed for binary classification.

Gradient Boosting (MART)

Algorithm 10.3 Gradient Tree Boosting Algorithm.

- 1. Initialize $f_0(x) = \arg\min_{\gamma} \sum_{i=1}^{N} L(y_i, \gamma)$.
- 2. For m=1 to M:
- (a) For $i = 1, 2, \dots, N$ compute

$$r_{im} = -\left[\frac{\partial L(y_i, f(x_i))}{\partial f(x_i)}\right]_{f=f_{m-1}}.$$

- (b) Fit a regression tree to the targets r_{im} giving terminal regions $R_{jm}, j = 1, 2, ..., J_m$.
- (c) For $j = 1, 2, \ldots, J_m$ compute

$$\gamma_{jm} = \arg\min_{\gamma} \sum_{x_i \in R_{jm}} L(y_i, f_{m-1}(x_i) + \gamma).$$

- (d) Update $f_m(x) = f_{m-1}(x) + \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$.
 - 3. Output $\hat{f}(x) = f_M(x)$.

Tree Size

- Tree size one approach, decide independently. Does not work well in practice for the boosting.
- Alternative strategy: require all of the trees to be the same size $J_m = J$, $\forall m$, where J_m is a j-terminal node tree.
- *J* is a meta-parameter.

What is the effect of different size J?

Tree Size

 Interaction level is limited to tree size and this constraint carries over to boosted tree.

Example:

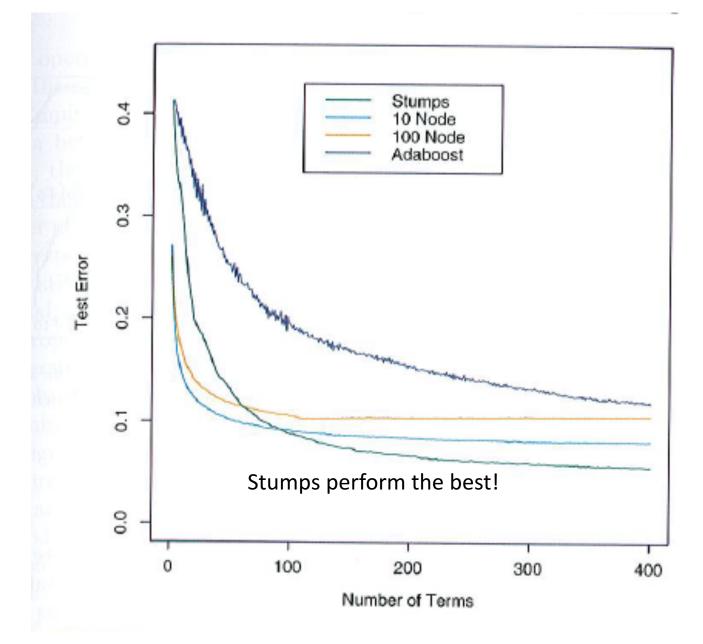
 $J = 2 \sim \text{produces a boosted model with only main effects.}$

 $J = 3 \sim 2$ variable interactions permitted.

• The value of J should reflect knowledge about the number of interactions (usually unknown, but most often low).

Rule of thumb: J=2 too small, J>10 too big, between 4 and 10 is good.

Tree Size



Shrinkage

- The second consideration, how many trees to include in the boosting.
- Same issues wrt overfitting.
- Strategy: model prediction risk (error) on a validation sample. Find the M^* that minimizes the predicted risk.

Shrinkage

Algorithm 10.3 Gradient Tree Boosting Algorithm.

- 1. Initialize $f_0(x) = \arg\min_{\gamma} \sum_{i=1}^{N} L(y_i, \gamma)$.
- 2. For m=1 to M:
 - (a) For $i = 1, 2, \dots, N$ compute

$$r_{im} = -\left[\frac{\partial L(y_i, f(x_i))}{\partial f(x_i)}\right]_{f=f_{m-1}}.$$

- (b) Fit a regression tree to the targets r_{im} giving terminal regions $R_{jm}, j = 1, 2, ..., J_m$.
- (c) For $j = 1, 2, \ldots, J_m$ compute

$$\gamma_{jm} = \arg\min_{\gamma} \sum_{x_i \in R_{jm}} L(y_i, f_{m-1}(x_i) + \gamma).$$

- (d) Update $f_m(x) = f_{m-1}(x) + \sum_{j=1}^{J_m} \gamma_{jm} I(x \in R_{jm})$.
- 3. Output $\hat{f}(x) = f_M(x)$.

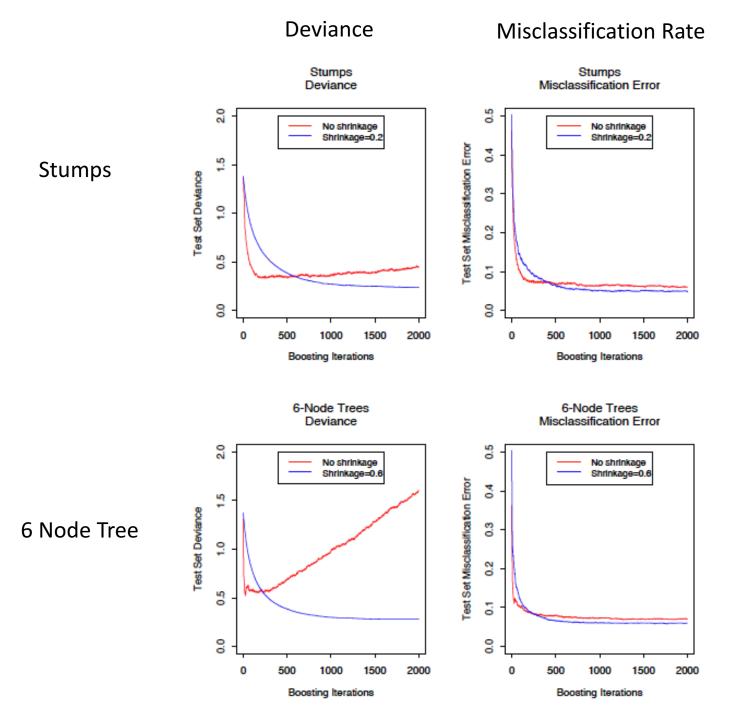
$$f_m(x) = f_{m-1}(x) + v \cdot \sum_{j=1}^{J} \gamma_{jm} I\left(x \in R_{jm}\right)$$

Shrinkage

$$f_m(x) = f_{m-1}(x) + v \cdot \sum_{j=1}^{J} \gamma_{jm} I(x \in R_{jm})$$

- Now we have another complexity parameter that also controls prediction risk.
- Inverse relationship between v and M.
- Empirically, it has been shown that smaller values of $\ ^{
 u}$ favor better test error and require corresponding larger values of M .

Another possibility.... Subsampling.



Blue: No Shrinkage

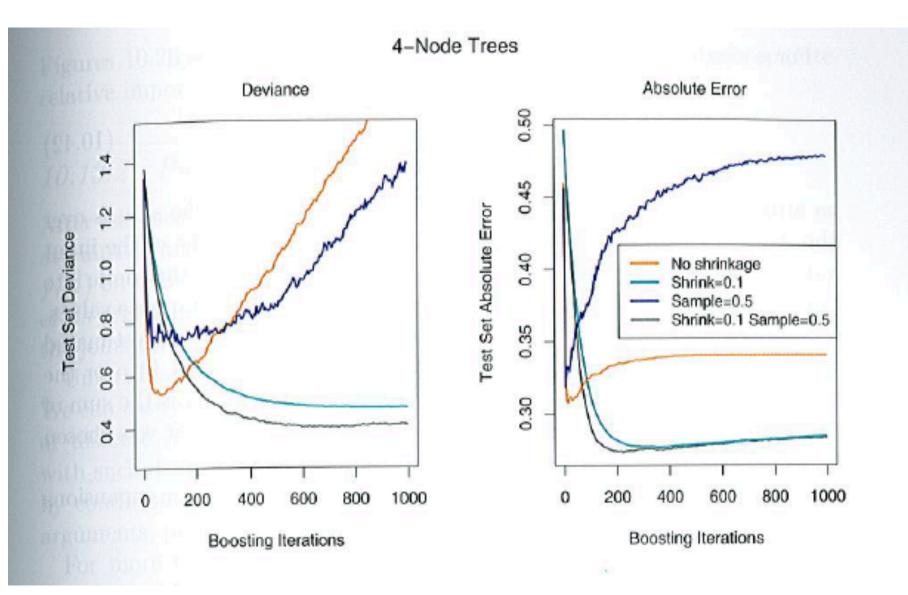
Red: Shrinkage

Number of Trees

Stochastic Gradient Boosting \sim at each iteration, we sample a fraction of the training data η without replacement. Grow the next tree using the subsample.

- Reduction in computation
- Often produces a more accurate model when used with shrinkage.
- We are up to four complexity parameters: J,M,v, and η .

Number of Trees



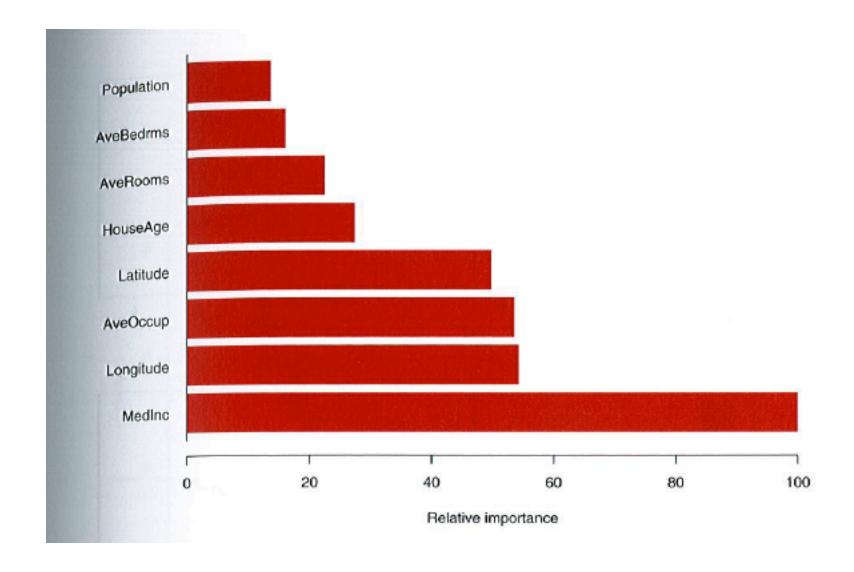
Interpretation

- Interpretation is lost with boosting.
- We need to rely on alternative metrics to reveal the model structure.

Relative Importance of a Predictor variable.

- Often many nuisance predictors which cloud model interpretation.
- Important to understand how important a variable is in predicting response, and dividing classes.

Relative Importance



Summary

- Boosting trees is complicated.
- Steepest descent a good idea... but prone to being too "greedy" by nature of the step.
- Gradient boosting a "less greedy" alternative.
- Main idea: fit trees to "residuals" defined as gradient function evaluations. Update functional approximation in an iterative manner. (Quasi-Newton like).
- Several improvements can be made to gradient boosting, including: subsampling, and shrinkage.
- Lots of parameters have to be set: number of trees, depth, subsampling, and shrinkage!

But..... It really works!