

- Decision trees
- Ensemble methods
 - Bagging
 - Random forests
 - (Gradient) Boosting

Machine learning - Block 2

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Autumn 2023



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This week's lecture

- Trees
- Bagging
- Random Forest
- Boosting (Trees)



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Assignment 1

- 1. Research ammanuens positions (https://swerik-project.github.io/)
- 2. Master Thesis project proposals
- 3. Assignment 1: Overall satisfaction



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Section 1

Decision trees



- Decision trees
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 A popular method that can be used for both classification and regression is decision trees.



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- A popular method that can be used for both classification and regression is decision trees.
- Have you ever played the game "20 questions"?



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- Have you ever played the game "20 questions"?
- Decision trees is more or less that game!



Decision trees

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- A popular method that can be used for both classification and regression is decision trees.
- Have you ever played the game "20 questions"?
- Decision trees is more or less that game!
- In the case of classification, the idea is to classify the new observation by
 - 1. Asking a questions
 - 2. Based on the previous answer, ask new question
 - 3. Questions are asked until a conclusion is reached



- Decision trees
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Name	Body temp	Gives birth	Legs	Class
Human	warm-blooded	yes	yes	mammal
Whale	warm-blooded	yes	no	mammal
Cat	warm-blooded	yes	yes	mammal
Cow	warm-blooded	yes	yes	mammal
Python	cold-blooded	no	no	reptile
Komodo dragon	cold-blooded	no	yes	reptile
Turtle	cold-blooded	no	yes	reptile
Salmon	cold-blooded	no	no	fish
Eel	cold-blooded	no	no	fish
Pigeon	warm-blooded	no	yes	bird
Penguin	warm-blooded	no	yes	bird
	•		•	•



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Classify Komodo dragon with a decision tree:

1. Does it give live birth?



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Classify Komodo dragon with a decision tree:

- 1. Does it give live birth? (No!)
- 2. Is it warm-blooded?



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Classify Komodo dragon with a decision tree:

- 1. Does it give live birth? (No!)
- 2. Is it warm-blooded? (No!)
- 3. Does it have legs?



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Classify Komodo dragon with a decision tree:

- 1. Does it give live birth? (No!)
- 2. Is it warm-blooded? (No!)
- 3. Does it have legs? (Yes!) \rightarrow Reptile



Decision trees

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Regression trees: The regions of a tree

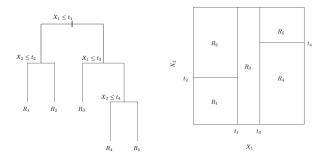


Figure: Regions of a tree (Garreth et al, 2013, Fig. 8.3)



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Regression Trees

$$T(x) = \sum_{m=1}^{M} \gamma_m I(x \in R_m),$$

where M is the total number of regions and $I(x \in R_m)$ is an indicator variable if x_i belongs to region R_m and γ_m is the prediction for region m.



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Regression Trees

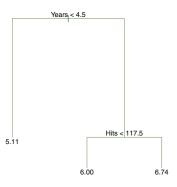


Figure: Regression Tree (Garreth et al, 2013, Fig. 8.1.)

- The Hitters dataset: log Salaries of Baseball players.
- The end of the tree contain the observations.



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Regression Trees

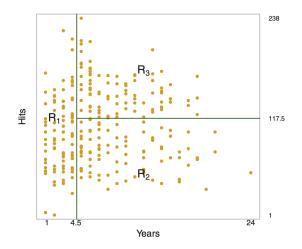


Figure: Hitters data and regression tree regions (Garreth et al, 2013, Fig. 8.2.)



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1. A tree has two groups of parameters $\Theta = (\gamma, R)$ that we need to learn.



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- 3. Usually we estimate γ_m as the mean of y_i in the region as:

$$\hat{\gamma}_m = \frac{1}{N_m} \sum_{\mathbf{x}:\in R_m}^{N_m} y_i \,,$$

where N_m is the number of observations in region R_m .



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4. Learning R_m exact is generally computationally infeasable so we use a greedy heuristic.



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Growing a Decision Tree: Greedy Algorithm

Let S be the set of all observations $\{1, ..., N\}$ and S[[m]] be the set of observation indecies in R_m and 1 is the minimal number of leafs per node.

Input: S, X, y, 1

- 1. S[[1]] = S,M = 1, m = 1
- 2. while m <= M then do:
 - 2.1 if(size(S[[m]]) >= 2*1) 2.1.1 S[[M+1]], S[[M+2]], j[m], s[m] =
 - split_tree(X[S[[m]],], y[S[[m]],],1)
 212 M = M + 2
 - 2.2 else
 - 2.2.1 compute $\hat{\gamma}$ for S[[m]]
 - 2.3 m = m + 1

Output: j, s, γ

Example of a tree: j = {Years, Hits}, s = {4.5, 117.5}, $\hat{\gamma} = \{122, 317, 245\}$



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How to do a split?

Here we try to compute Eq. (9.12-9.14) in ESL:

$$R_1(j,s) = \{X | X_j \le s\} \text{ and } R_2(j,s) = \{X | X_j > s\}$$

$$\min_{j,s} \left(\min_{c_1} \sum_{x_i \in R_1(j,s)} (y_i - c_1)^2 + \min_{c_2} \sum_{x_i \in R_2(j,s)} (y_i - c_2)^2 \right)$$

Inner minimization is solved by:

$$\hat{c}_m = \frac{1}{N_m} \sum_{x_i \in R_-}^{N_m} y_i \,,$$



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How to do a split? Pseudo-code

Input: $\mathbf{X}, \mathbf{y}, I$

- 1. $SS = Inf \# Store sum of squares in matrix of dim <math>P \times N_s$
- 2. $S = Inf \# Store split point in matrix of dim <math>P \times N_s$
- 3. for $j \in \{1, ..., P\}$ # all features
 - 3.1 for $k \in \{1, ..., N_s\}$ # all observations in set s
 - 3.1.1 $s = x_{k,j} \# Split point (use the value of x)$
 - 3.1.2 if $(|R_1(s,j)| < l \text{ or } |R_2(s,j)| < l)$ next # Dont create too few leaves
 - 3.1.3 $\hat{c}_1 = \frac{1}{|R_1(s,j)|} \sum_{x_i \in R_1(s,j)} y_i$
 - 3.1.4 $\hat{c}_2 = \frac{1}{|R_2(s,j)|} \sum_{x_i \in R_2(s,j)} y_i$
 - 3.1.5 $SS_{k,j} = \sum_{x_i \in R_1(s,j)} (y_i c_1)^2 + \sum_{x_i \in R_2(s,j)} (y_i c_2)^2 \#$ Compute Sum of Squares
 - 3.1.6 $S_{k,j} = s$
- 4. $k_{final}, j_{final} = \min_{k,j} SS$
- 5. $s_{final} = S_{k_{final}, j_{final}}$
- 6. return $R_1(s_{final}, j_{final}), R_2(s_{final}, j_{final}), s_{final}, j_{final}$



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• How do we do if we have a classification tree?



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- We just change the loss function $L(\theta)$.



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- How do we do if we have a classification tree?
- We just change the loss function $L(\theta)$.
- Let p(j|t) be the fraction of observations in class j at the node t and let J be the number of classes.



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- How do we do if we have a classification tree?
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- Let p(j|t) be the fraction of observations in class j at the node t and let J be the number of classes.
- The Gini for node t is defined as

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• Gini is a measure of "impurity". If all observations belong to the same class, then

$$Gini(t) = 1 - 1^2 - 0 - \ldots - 0 = 0.$$



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• Gini is a measure of "impurity". If all observations belong to the same class, then

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- The Gini is maximized when all classes have the same number of observations at t
- One criterion for splitting could be to minimize the Gini in the next level of the tree. That way we will get "purer" nodes.



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 Tree depth: the length of the longest path from the root to a leaf (i.e. greatest number of questions that the tree can ask).



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- Leaf size: the number of observations in a leaf.



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Decision trees can become quite large, which may lead to:

- Overfitting (high variance)
- Difficulties interpreting the tree



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The solution to this is

 Pruning: forcing the tree to be smaller by adding a stopping condition, e.g. a maximum depth or minimal leaf size.



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The solution to this is

- Pruning: forcing the tree to be smaller by adding a stopping condition, e.g. a maximum depth or minimal leaf size.
- But decision trees are quite bad predition models...



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Section 2

Ensemble methods



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Simulated example (Prize academy, see ESL):

 50 members vote in 10 categories, each with 4 nominations.



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Simulated example (Prize academy, see ESL):

- 50 members vote in 10 categories, each with 4 nominations.
- 2. For any category, only 15 voters have some knowledge (p > 0.25)



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Simulated example (Prize academy, see ESL):

- 50 members vote in 10 categories, each with 4 nominations.
- 2. For any category, only 15 voters have some knowledge (p > 0.25)
- 3. For each category, the 15 experts are chosen at random from the 50.



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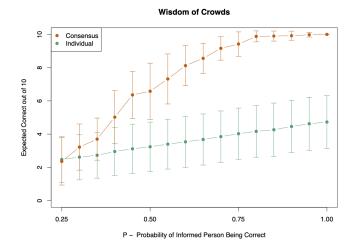


Figure: Simulated Award Voting, Fig. 8.11 (ESL)



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General idea of ensembles

The idea of an ensemble is simple: If it difficult to find one really good model perhaps we can find several weaker models and combine their predictions.



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General idea of ensembles

The idea of an ensemble is simple: If it difficult to find one really good model perhaps we can find several weaker models and combine their predictions.

A simple example: Say you have one outcome Y and 4 covariates X_1, X_2, X_3, X_4 . The goal is to predict Y. A possible ensemble would be to fit

$$y = \alpha_1 + \beta_1 X_1 + \epsilon_1$$
$$y = \alpha_2 + \beta_2 X_2 + \epsilon_2$$
$$y = \alpha_3 + \beta_3 X_3 + \epsilon_3$$
$$y = \alpha_4 + \beta_4 X_4 + \epsilon_4$$

and then use the mean of their predictions

$$\hat{y}_{ensemble} = \frac{1}{4} \sum_{i=1}^{4} \hat{y} = \frac{1}{4} \sum_{i=1}^{4} (\hat{\alpha}_i + \hat{\beta}_i X_i)$$
 (1)



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Two key parts of an ensemble

- 1. The prediction models (sometimes called 'learners')
 - A single model in an ensemble can be a simple or a complex model
 - Often the ensemble contains many simple models.



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Two key parts of an ensemble

- 1. The prediction models (sometimes called 'learners')
 - A single model in an ensemble can be a simple or a complex model
 - Often the ensemble contains many simple models.
- The weighting of each prediction in the final ensemble prediction
 - Many different algorithms for weighting together the predictions from many models
 - Models/learners with better predictive power can be given larger weights in the final prediction



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Ensambles of decision trees

- A common type of ensembles is ensembles of decision trees.
- We will focus on this case, but note that any type of model can be included in an ensemble in principle.



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Remember, the error of a prediction/classification can be decomposed as

$$error = bias^2 + variance + bayeserror.$$
 (2)



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 Complex models/strong learners (with many parameters) tend to have small bias and large variance (tend to be overfitted)



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- Complex models/strong learners (with many parameters) tend to have small bias and large variance (tend to be overfitted)
- Shallow models/weak learners (with few parameters) tend to have small variance and large bias



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- Complex models/strong learners (with many parameters) tend to have small bias and large variance (tend to be overfitted)
- Shallow models/weak learners (with few parameters) tend to have small variance and large bias

Bagging: Ensemble methods that aim to decrease the variance of complex/strong learners with low bias and large variance

Boosting: Ensemble methods that aim to decrease the bias of shallow/weak learners with low variance and large bias



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Subsection 1

Bagging



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Bagging (Bootstrap AGGregating)

- Train several deep trees and combine their results
- Use bootstrap to train different trees

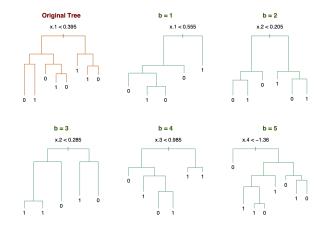


Figure: Bagging trees, Fig. 8.9 (ESL)



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Bagging (Bootstrap AGGregating)

- 1. Draw, *with replacement*, a random sample of *N* units from the original sample
- 2. Fit a prediction model (e.g., a deep decision tree)
- 3. Repeat steps 1-2 B times
- Weight together the predictions from the B models into a final ensemble prediction as

$$\hat{f}_{bag}(x_i) = \frac{1}{B} \sum_b \hat{f}^b(x_i) = \frac{1}{B} \sum_b \hat{T}(x_i | \Theta_b)$$



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Subsection 2

Random forests



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Random Forest

• A bagging ensemble method, but...



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Random Forest

- A bagging ensemble method, but...
- Sample of N observations and K covariates/features



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Random Forest

- A bagging ensemble method, but...
- Sample of *N* observations and *K* covariates/features
- It is common to use $k = \sqrt{K}$ (rounded down) for classification and k = K/3 for regression. But these are only rules of thumb: k is a *tuning parameter*.



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Random Forest: Algorithm

- 1. Draw, with replacement, a random sample of N units from the original sample
- 2. Draw, without replacement, a random subset of *k* covariates/features
- 3. Fit a prediction model (e.g., a decision tree)
- 4. Repeat step 1-3 B times
- 5. Weight together the predictions from the B models into a final ensemble prediction

$$\hat{f}_{\mathsf{rf}}(x_i) = \frac{1}{B} \sum_b \hat{T}(x_i | \Theta_b)$$



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- Consider each tree to be an i.i.d. random variable with variance σ^2 .
- The average of these trees then have variance

$$\mathbb{V}(\hat{f}(x) = \frac{1}{B}\sigma^2.$$



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- Consider each tree to be an i.i.d. random variable with variance σ^2 .
- The average of these trees then have variance

$$\mathbb{V}(\hat{f}(x) = \frac{1}{B}\sigma^2.$$

- Trees constructed from the same set of covariates will be correlated and therefore not independent.
- The variance of the average of these correlated trees then becomes

$$\mathbb{V}(\hat{f}(x)) = \rho \sigma^2 + \frac{1 - \rho}{B} \sigma^2.$$



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• The variance of the average of correlated trees:

$$\mathbb{V}(\hat{f}(x)) = \rho \sigma^2 + \frac{1 - \rho}{B} \sigma^2.$$

 The second term will vanish with increasing B leaving just the first term left: a function of the correlation between the trees



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• The variance of the average of correlated trees:

$$\mathbb{V}(\hat{f}(x)) = \rho \sigma^2 + \frac{1 - \rho}{B} \sigma^2.$$

- The second term will vanish with increasing B leaving just the first term left: a function of the correlation between the trees
- The remaining part of the variance is minimized by only consider a subset of the covariates when constructing trees - reducing the correlation between trees.



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Bagging vs. Random forest

- In bagging, the trees are often highly correlated
 - If some covariates are strong predictors of the outcome (in the training data), many trees in the 'bag' will us the same covariates in their decisions



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- Intuition: A random forest (with many trees) uses the predictive ability of all covariates rather than just a few → improved out of sample performance.



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Subsection 3

(Gradient) Boosting



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Boosting

1. In boosting, models/trees are trained sequentially



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Boosting

- 1. In boosting, models/trees are trained sequentially
- 2. Each new model tries to target weak spots of the previous models



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Boosting (AdaBoost)

- 1. Initialize weight for all observations as $w_i = n^{-1}$
- 2. Repeat B times
 - 2.1 Fit a simple classifier $h_b(x)$ using w_i s
 - 2.2 Compute the weighted error (between 0-1) as

$$e_b = \frac{\sum w_i I(y_i \neq h_b(x_i))}{\sum w_i}$$

2.3 Compute the importance of the classifier as

$$\alpha_b = \log((1 - e_b)/e_b) = \log it(1 - e_b)$$

2.4 Add the classifier to the ensemble

$$\hat{f}_b(x) = \hat{f}_{b-1}(x) + \alpha_b h_b(x)$$

2.5 Update weights so that badly classified observations is weighted more

$$w_i \leftarrow w_i \exp(\alpha_b I(y_i \neq h_b(x_i)))$$

3. Output the final ensemble $\hat{f}_B(x)$



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Boosting example

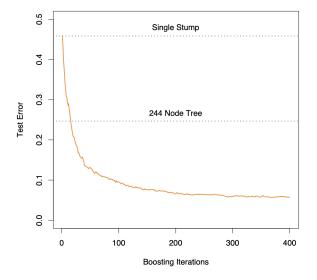


Figure: Boosting example (using stumps as h_b), Fig. 10.2 (ESL)



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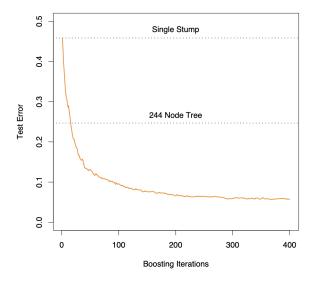


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• A more general approach is (gradient) boosting trees



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- Let

$$f_{\rm gb}(x) = \sum_{b}^{B} \hat{T}(x|\Theta_b)$$



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We want to minimize the loss

$$L(y, f_{\mathsf{gb}}(x))$$



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We want to minimize the loss

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• This means finding

$$\Theta_b = \arg\min \sum_{i}^{N} L(y_i, f_{b-1}(x_i) + \hat{T}(x|\Theta_b))$$



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Gradient Boosting Trees

- 1. Initialize $f_0(x)$
- 2. Repeat B times
 - **2.1** For i = 1, 2, ..., N compute

$$r_{ib} = -\frac{\partial L(y_i, f(x_i))}{\partial f(x_i)}$$

- 2.2 Compute a regression tree $\hat{T}(x|\Theta_b)$ on r_b
- 2.3 Add the classifier to the ensemble

$$\hat{f}_b(x) = \hat{f}_{b-1}(x) + \hat{T}(x|\Theta_b)$$

3. Output the final ensemble $\hat{f}_B(x)$



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Comparisons

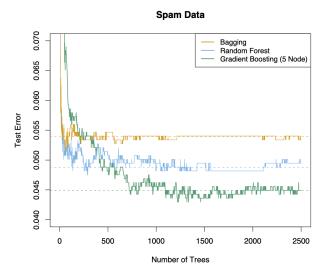


Figure: Comparing bagging, random forests and boosting, Fig. 15.1 (ESL)



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• State of the Art method



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- State of the Art method
- Use gradient boosting trees with regularization

$$L(y, f_{\mathsf{boost}}(x)) + \sum_{b} \Omega(\hat{T}(x_i|\Theta_b))$$



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$$L(y, f_{\mathsf{boost}}(x)) + \sum_{b} \Omega(\hat{T}(x_i|\Theta_b))$$

where

$$\Omega(\hat{T}(x_i|\Theta_b)) = \lambda_1 Q_b + \lambda_2 ||\gamma_b||_2^2$$

where Q_b is the number of leafs in tree b and $||\cdot||_2$ is the Euclidian/ L^2 norm.



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• Is scalable to very large data