

# Lecture 6 – Tree-based methods, Bagging and Boosting



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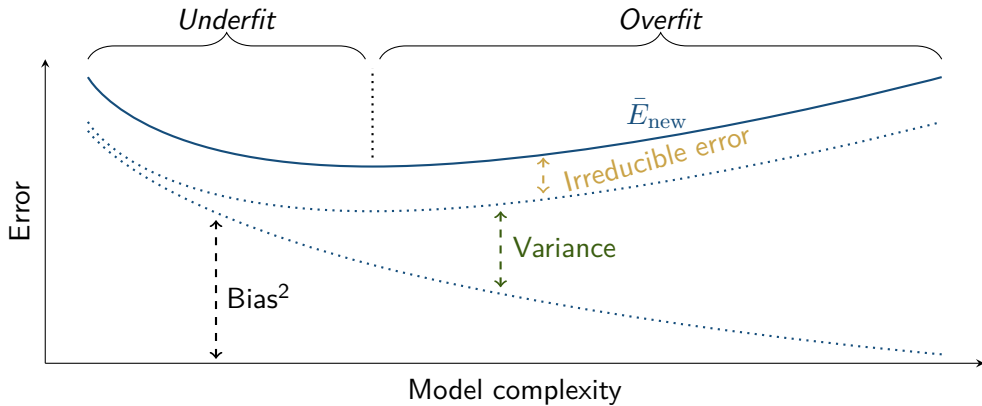
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# Summary of Lecture 5



Finding a balanced fit (neither over- nor underfit) is called the **the bias-variance tradeoff**.

# Contents – Lecture 6

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1. Classification and regression trees (CART)
2. Bagging – *a general variance reduction technique*
3. Random forests
4. Boosting

# The idea behind tree-based methods

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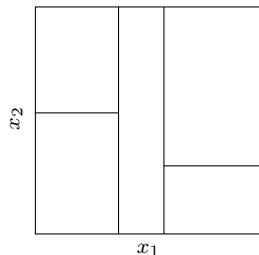
In both regression and classification settings we seek a function  $\hat{y}(\mathbf{x})$  which maps the input  $\mathbf{x}$  into a prediction.

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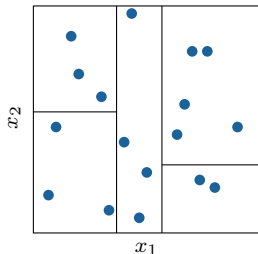
One **flexible** way of designing this function is to partition the input space into disjoint regions and fit a simple model in each region.



# The idea behind tree-based methods

In both regression and classification settings we seek a function  $\hat{y}(\mathbf{x})$  which maps the input  $\mathbf{x}$  into a prediction.

One **flexible** way of designing this function is to partition the input space into disjoint regions and fit a simple model in each region.



• = Training data

- **Classification:** Majority vote within the region.
- **Regression:** Mean of training data within the region.

# Finding the partition

---

The key challenge in using this strategy is to find a good partition.

Even if we restrict our attention to seemingly simple regions (e.g. “boxes”), finding an *optimal* partition w.r.t. minimizing the training error is *computationally infeasible!*

Instead, we use a “greedy” approach: **recursive binary splitting**.

1. Select one input variable  $x_j$  and a cut-point  $s$ . Partition the input space into two half-spaces,

$$\{\mathbf{x} : x_j < s\}$$

$$\{\mathbf{x} : x_j \geq s\}.$$

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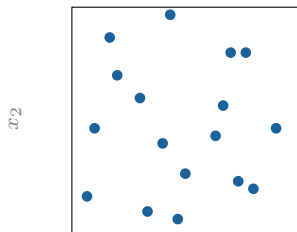
2. Repeat this splitting for each region until some stopping criterion is met (e.g., no region contains more than 5 training data points).

# Recursive binary splitting

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Partitioning of input space

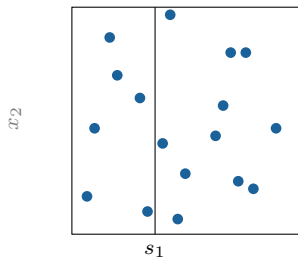
Tree representation



• = Training data

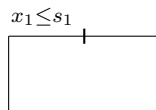
# Recursive binary splitting

Partitioning of input space



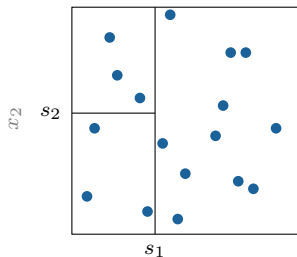
• = Training data

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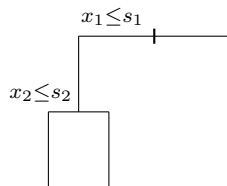
# Recursive binary splitting

Partitioning of input space



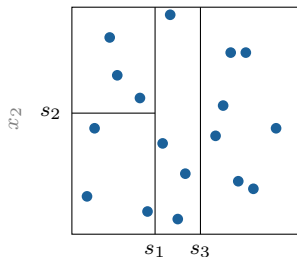
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Tree representation



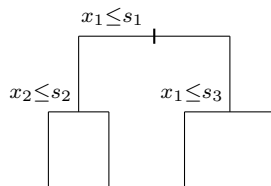
# Recursive binary splitting

Partitioning of input space



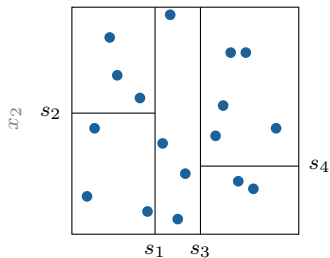
• = Training data

Tree representation



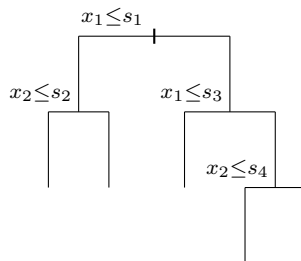
# Recursive binary splitting

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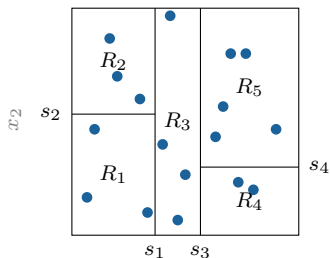
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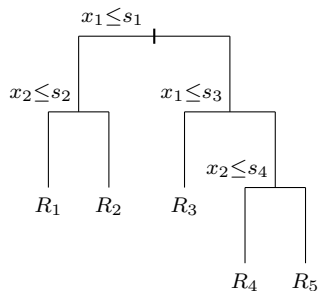
# Recursive binary splitting

Partitioning of input space



• = Training data

Tree representation



# Regression trees (I/II)

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Once the input space is partitioned into  $L$  regions,  $R_1, R_2, \dots, R_L$  the prediction model is

$$\hat{y}_\star = \sum_{\ell=1}^L \hat{y}_\ell \mathbb{I}\{\mathbf{x}_\star \in R_\ell\},$$

where  $\mathbb{I}\{\mathbf{x}_\star \in R_\ell\}$  is the indicator function

$$\mathbb{I}\{\mathbf{x}_\star \in R_\ell\} = \begin{cases} 1 & \text{if } \mathbf{x}_\star \in R_\ell \\ 0 & \text{if } \mathbf{x}_\star \notin R_\ell \end{cases}$$

and  $\hat{y}_\ell$  is a constant prediction within each region.

For regression trees we use

$$\hat{y}_\ell = \text{avarage}\{y_i : \mathbf{x}_i \in R_\ell\}$$



## Regression trees (II/II) - How do we find the partition?

Recursive binary splitting is **greedy** - each split is made in order to minimize the loss **without looking ahead** at future splits.

For any  $j$  and  $s$ , define

$$R_1(j, s) = \{\mathbf{x} \mid x_j < s\} \quad \text{and} \quad R_2(j, s) = \{\mathbf{x} \mid x_j \geq s\}.$$

We then seek  $(j, s)$  that minimize

$$\sum_{i: \mathbf{x}_i \in R_1(j, s)} (y_i - \hat{y}_1(j, s))^2 + \sum_{i: \mathbf{x}_i \in R_2(j, s)} (y_i - \hat{y}_2(j, s))^2$$

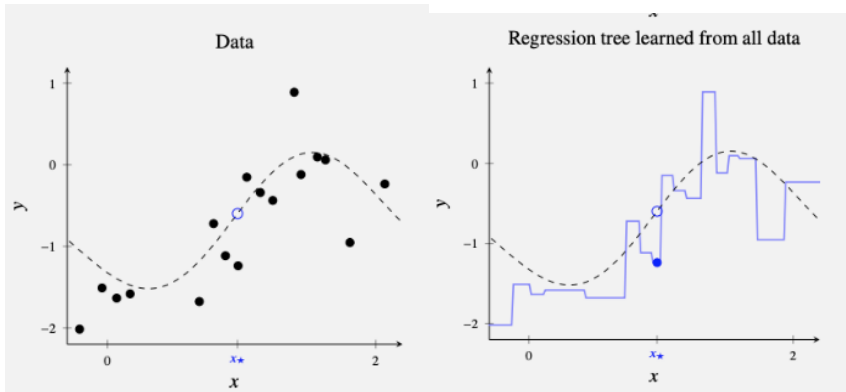
where

$$\hat{y}_1 = \text{average}\{y_i : \mathbf{x}_i \in R_1(j, s)\}$$

$$\hat{y}_2 = \text{average}\{y_i : \mathbf{x}_i \in R_2(j, s)\}$$

This optimization problem is easily solved by "brute force" by evaluating all possible splits.

# Example: Regression trees



# Classification trees

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Classification trees are constructed similarly to regression trees, but with *two differences*.

**Firstly**, the class prediction for each region is based on the proportion of data points from each class in that region. Let

$$\hat{\pi}_{\ell m} = \frac{1}{n_{\ell}} \sum_{i: \mathbf{x}_i \in R_{\ell}} \mathbb{I}\{y_i = m\}$$

be the proportion of training data points in the  $\ell$ th region that belong to the  $m$ th class.

$$p(y = m \mid \mathbf{x}_{\star}) \approx \sum_{\ell=1}^L \hat{\pi}_{\ell m} \mathbb{I}\{\mathbf{x}_{\star} \in R_{\ell}\}$$

# Classification trees

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**Secondly**, the squared loss used to construct the tree needs to be replaced by a measure suitable to categorical outputs.

Three common error measures are,

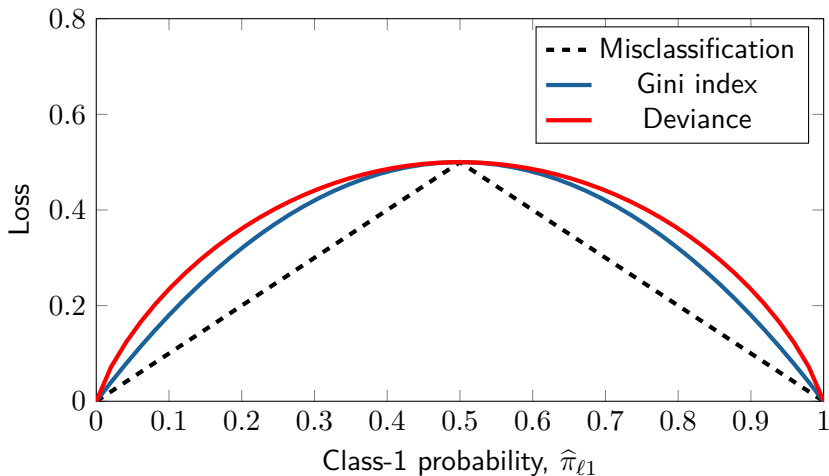
$$\text{Misclassification error: } 1 - \max_m \hat{\pi}_{\ell m}$$

$$\text{Entropy/deviance: } - \sum_{m=1}^M \hat{\pi}_{\ell m} \log \hat{\pi}_{\ell m}$$

$$\text{Gini index: } \sum_{m=1}^M \hat{\pi}_{\ell m} (1 - \hat{\pi}_{\ell m})$$

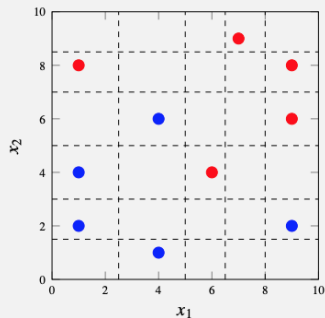
# Classification error measures

For a binary classification problem ( $M = 2$ )



# Example using entropy

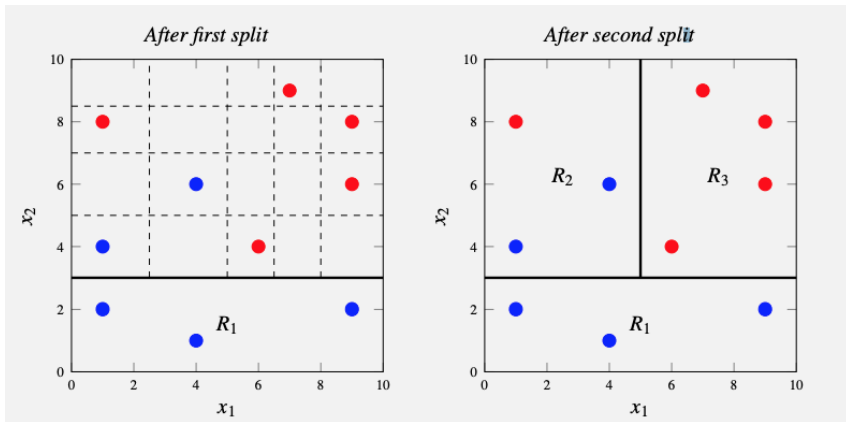
$x_1$	$x_2$	$y$
9.0	2.0	Blue
1.0	4.0	Blue
4.0	6.0	Blue
4.0	1.0	Blue
1.0	2.0	Blue
1.0	8.0	Red
6.0	4.0	Red
7.0	9.0	Red
9.0	8.0	Red
9.0	6.0	Red



# Example using entropy

Split ( $R_1$ )	$n_1$	$\hat{\pi}_{1B}$	$\hat{\pi}_{1R}$	$Q_1$	$n_2$	$\hat{\pi}_{2B}$	$\hat{\pi}_{2R}$	$Q_2$	$n_1Q_1 + n_2Q_2$
$x_1 < 2.5$	3	2/3	1/3	0.64	7	3/7	4/7	0.68	6.69
$x_1 < 5.0$	5	4/5	1/5	0.50	5	1/5	4/5	0.50	5.00
$x_1 < 6.5$	6	4/6	2/6	0.64	4	1/4	3/4	0.56	6.07
$x_1 < 8.0$	7	4/7	3/7	0.68	3	1/3	2/3	0.64	6.69
$x_2 < 1.5$	1	1/1	0/1	0.00	9	4/9	5/9	0.69	6.18
$x_2 < 3.0$	3	3/3	0/3	0.00	7	2/7	5/7	0.60	<b>4.18</b>
$x_2 < 5.0$	5	4/5	1/5	0.50	5	1/5	4/5	0.06	5.00
$x_2 < 7.0$	7	5/7	2/7	0.60	3	0/3	3/3	0.00	<b>4.18</b>
$x_2 < 8.5$	9	5/9	4/9	0.69	1	0/1	1/1	0.00	6.18

# Example using entropy





# Improving CART

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The flexibility/complexity of classification and regression trees (CART) is decided by the tree depth.

! To obtain a **small bias** the tree need to be grown deep,

! but this results in a **high variance!**

The performance of (simple) CARTs is often unsatisfactory!

To improve the practical performance:

- Bagging and Random Forests
- Boosted trees

# Bagging (I/II)

---

For now, assume that we have access to  $B$  **independent** datasets  $\mathcal{T}^1, \dots, \mathcal{T}^B$ . We can then train a separate deep tree  $\hat{y}^b(\mathbf{x})$  for each dataset,  $1, \dots, B$ .

- Each  $\hat{y}^b(\mathbf{x})$  has a **low bias** but **high variance**

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- Each  $\hat{y}^b(\mathbf{x})$  has a **low bias** but **high variance**
- By averaging

$$\hat{y}_{\text{bag}}(\mathbf{x}) = \frac{1}{B} \sum_{b=1}^B \hat{y}^b(\mathbf{x})$$

the bias is kept small, but variance is reduced by a factor  $B$ !

# Bagging (II/II)

---

**Obvious problem** We only have access to one training dataset.

**Solution** Bootstrap the data!

- Sample  $n$  times with replacement from the original training data  $\mathcal{T} = \{\mathbf{x}_i, y_i\}_{i=1}^n$

For each bootstrapped dataset  $\tilde{\mathcal{T}}^b$  we train a tree  $\tilde{y}^b(\mathbf{x})$ .

$$\tilde{y}_{\text{bag}}^b(\mathbf{x}) = \frac{1}{B} \sum_{b=1}^B \tilde{y}^b(\mathbf{x})$$

is called "bootstrap aggregation", or bagging.

# Bagging (II/II)

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**Obvious problem** We only have access to one training dataset.

**Solution** Bootstrap the data!

- Sample  $n$  times with replacement from the original training data  $\mathcal{T} = \{\mathbf{x}_i, y_i\}_{i=1}^n$
- Repeat  $B$  times to generate  $B$  "bootstrapped" training datasets  $\tilde{\mathcal{T}}^1, \dots, \tilde{\mathcal{T}}^B$

For each bootstrapped dataset  $\tilde{\mathcal{T}}^b$  we train a tree  $\tilde{y}^b(\mathbf{x})$ .

$$\tilde{y}_{\text{bag}}^b(\mathbf{x}) = \frac{1}{B} \sum_{b=1}^B \tilde{y}^b(\mathbf{x})$$

is called "bootstrap aggregation", or bagging.

# Bagging - Toy example

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ex) Assume that we have a training set

$$\mathcal{T} = \{(\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), (\mathbf{x}_3, y_3), (\mathbf{x}_4, y_4)\}.$$

We generate, say,  $B = 3$  datasets by bootstrapping:

$$\tilde{\mathcal{T}}^1 = \{(\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), (\mathbf{x}_3, y_3), (\mathbf{x}_3, y_3)\}$$

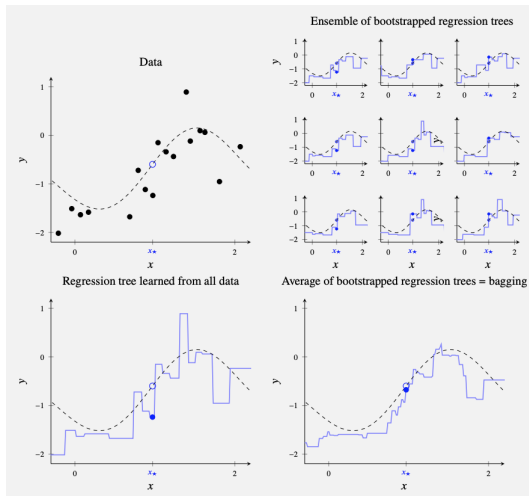
$$\tilde{\mathcal{T}}^2 = \{(\mathbf{x}_1, y_1), (\mathbf{x}_4, y_4), (\mathbf{x}_4, y_4), (\mathbf{x}_4, y_4)\}$$

$$\tilde{\mathcal{T}}^3 = \{(\mathbf{x}_1, y_1), (\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), (\mathbf{x}_2, y_2)\}$$

Compute  $B = 3$  (deep) regression trees  $\tilde{y}^1(\mathbf{x})$ ,  $\tilde{y}^2(\mathbf{x})$  and  $\tilde{y}^3(\mathbf{x})$ , one for each dataset  $\tilde{\mathcal{T}}^1$ ,  $\tilde{\mathcal{T}}^2$ , and  $\tilde{\mathcal{T}}^3$ , and average

$$\tilde{y}_{\text{bag}}(\mathbf{x}) = \frac{1}{3} \sum_{b=1}^3 \tilde{y}^b(\mathbf{x})$$

# Example: Regression trees



# Application) Predicting US Supreme Court behavior

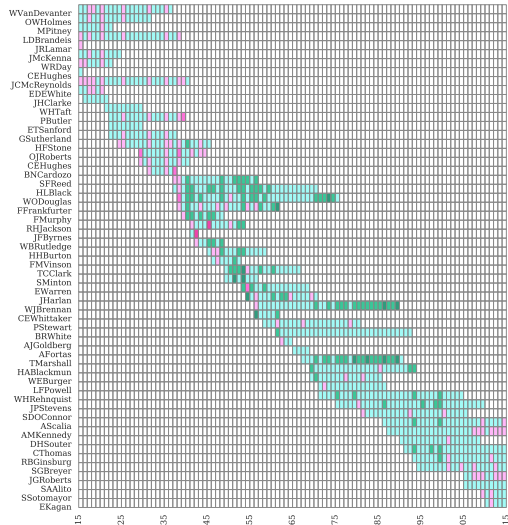
**Random forest** classifier built on SCDB data<sup>1</sup>  
to predict the votes of Supreme Court justices:

$$Y \in \{\text{affirm, reverse, other}\}$$

**Result:** 70% correct classifications

D. M. Katz, M. J. Bommarito II and J. Blackman. **A General Approach for Predicting the Behavior of the Supreme Court of the United States.**  
*arXiv.org*, arXiv:1612.03473v2, January 2017.

<sup>1</sup><http://supremecourtdatabase.org>





## *Application)* Predicting US Supreme Court behavior

---

*Not only have random forests proven to be “unreasonably effective” in a wide array of supervised learning contexts, but in our testing, random forests outperformed other common approaches including support vector machines [...] and feedforward artificial neural network models such as multi-layer perceptron*

— Katz, Bommarito II and Blackman (arXiv:1612.03473v2)

# Random forests

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Bagging can drastically improve the performance of CART!

However, the  $B$  bootstrapped dataset are ***correlated***  
 $\Rightarrow$  the variance reduction due to averaging is diminished.

Idea: De-correlate the  $B$  trees by randomly perturbing each tree.

A **random forest** is constructed by bagging, but for each split in each tree only a ***random subset*** of  $q \leq p$  inputs are considered as splitting variables.

Rule of thumb:  $q = \sqrt{p}$  for classification trees and  $q = p/3$  for regression trees.<sup>2</sup>

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<sup>2</sup>Proposed by Leo Breiman, inventor of random forests.

# Random forest pseudo-code

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## Algorithm Random forest for regression

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1. For  $b = 1$  to  $B$  (*can run in parallel*)
  - (a) Draw a bootstrap data set  $\tilde{\mathcal{T}}$  of size  $n$  from  $\mathcal{T}$ .
  - (b) Grow a regression tree by repeating the following steps until a minimum node size is reached:
    - i. Select  $q$  out of the  $p$  input variables uniformly at random.
    - ii. Find the variable  $x_j$  among the  $q$  selected, and the corresponding split point  $s$ , that minimizes the squared error.
    - iii. Split the node into two children with  $\{x_j \leq s\}$  and  $\{x_j > s\}$ .
2. Final model is the average the  $B$  ensemble members,

$$\hat{y}_{\star}^{\text{rf}} = \frac{1}{B} \sum_{b=1}^B \tilde{y}_{\star}^b.$$

# Random forests

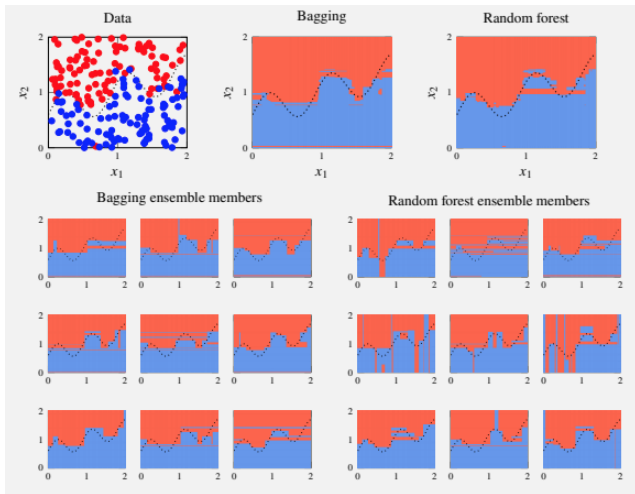
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The random input selection used in random forests:

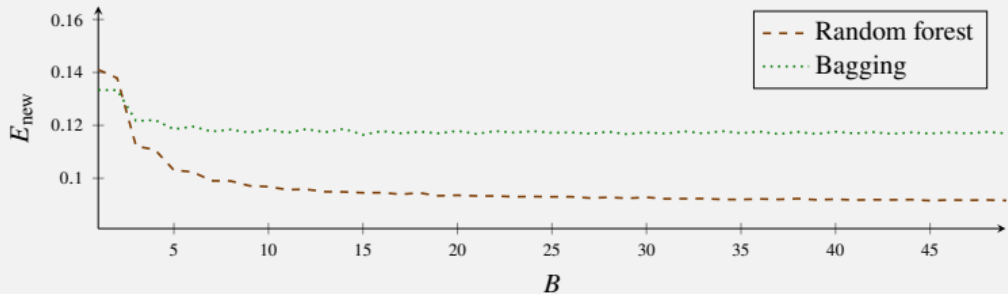
- ▼ increases the bias, but often very slowly
- ▼ adds to the variance ( $\sigma^2$ ) of each tree
- ▲ reduces the correlation between the trees

The reduction in correlation is typically the dominant effect  $\Rightarrow$  there is an overall reduction in MSE!

# ex)Random forest on regression data



## ex) Random forest on regression data



# Advantages of random forests

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Random forests have several **computational advantages**:

- ▲ Easy to do in parallel!
- ▲ Using  $q < p$  potential split variables reduces the computational cost of each split.
- ▲ We **could** bootstrap fewer than  $n$ , say  $\sqrt{n}$ , data points when creating  $\tilde{\mathcal{T}}^b$  — very useful for “big data” problems.

... and they also come with some other benefits:

- ▲ Often works well off-the-shelf – few tuning parameters
- ▲ Requires little or no input preparation
- ▲ Implicit input selection


## ex) Automatic music generation

**ALYSIA:** automated music generation using random forests.

- User specifies the lyrics
- ALYSIA generates accompanying music via
  - *rhythm model*
  - *melody model*
- Trained on a corpus of pop songs.

Why Do I Still Miss You?

Maya Ackerman ALYSIA



Vo. Now that you're gone, I just rea-lized I'm all a - lone.

Vo. For - give me if I, throw a - way the phone, stop won -

Vo. Chorus: dring where we when wrong. Tell me a - fter all you've

Vo. done, why do I still miss you? Why, do I still miss you.

[https://www.youtube.com/watch?v=whgudcj82\\_I](https://www.youtube.com/watch?v=whgudcj82_I)  
<https://www.withalysia.com/>

M. Ackerman and D. Loker. **Algorithmic Songwriting with ALYSIA.** In: *Correia J., Ciesielski V., Liapis A. (eds) Computational Intelligence in Music, Sound, Art and Design. EvoMUSART, 2017.*



# Boosting

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Even a simple (classification or regression) model can typically capture some aspects of the input-output relationship.

Can we then learn an **ensemble** of “weak models”, each describing some part of this relationship, and combine these into one “strong model”?

## Boosting:

- **Sequentially** learns an ensemble of **weak models**.
- Combine these into one **strong model**.
- General strategy – can in principle be used to improve any supervised learning algorithm.
- One of the most successful machine learning ideas!



# Boosting

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# Binary classification

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We will restrict our attention to binary classification.

- Class labels are  $-1$  and  $1$ , i.e.  $y \in \{-1, 1\}$ .
  - We have access to some (weak) base classifier, e.g. a classification tree.
- 

*Note.* Using labels  $-1$  and  $1$  is mathematically convenient as it allows us to express a majority vote between  $B$  classifiers  $\hat{G}^1(\mathbf{x}), \dots, \hat{G}^B(\mathbf{x})$  as

$$\text{sign} \left( \sum_{b=1}^B \hat{G}^b(\mathbf{x}) \right) = \begin{cases} +1 & \text{if more plus-votes than minus-votes,} \\ -1 & \text{if more minus-votes than plus-votes.} \end{cases}$$

# Boosting procedure (for classification)

---

## Boosting procedure:

1. Assign weights  $w_i^1 = 1/n$  to all data points.

# Boosting procedure (for classification)

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2. For  $b = 1$  to  $B$ 
  - (a) Train a weak classifier  $\hat{G}^{(b)}(\mathbf{x})$  on the **weighted training data**  $\{(\mathbf{x}_i, y_i, w_i^b)\}_{i=1}^n$ .

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  - (b) *Update the weights*  $\{w_i^{b+1}\}_{i=1}^n$  *from*  $\{w_i^b\}_{i=1}^n$ :
    - i. Increase weights for all points misclassified by  $\hat{G}^{(b)}(\mathbf{x})$ .
    - ii. Decrease weights for all points correctly classified by  $\hat{G}^{(b)}(\mathbf{x})$ .

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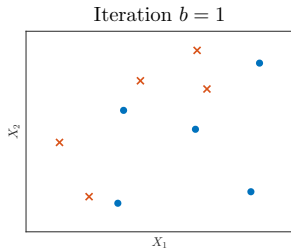
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The predictions of the  $B$  classifiers,  $\hat{G}^{(1)}(\mathbf{x}), \dots, \hat{G}^{(B)}(\mathbf{x})$ , are combined using a **weighted** majority vote:

$$\hat{G}_{\text{boost}}^B(\mathbf{x}) = \text{sign} \left( \sum_{b=1}^B \alpha^{(b)} \hat{G}^{(b)}(\mathbf{x}) \right).$$

## ex) Boosting illustration

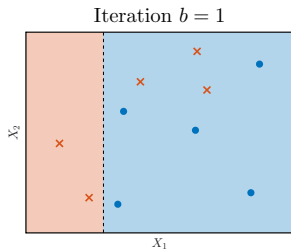
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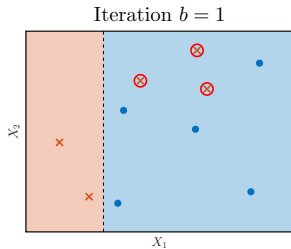
## ex) Boosting illustration

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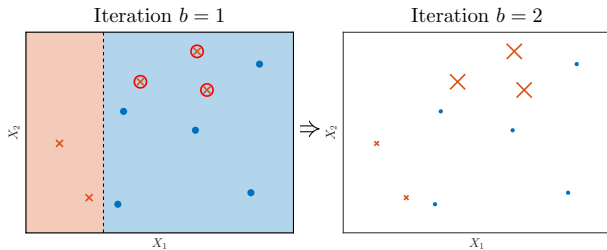


## ex) Boosting illustration

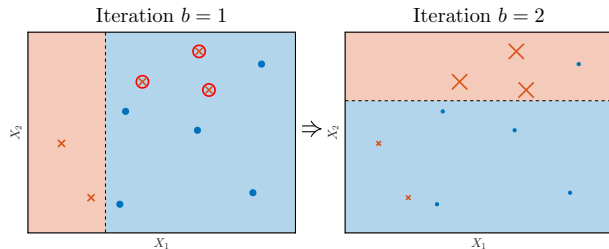
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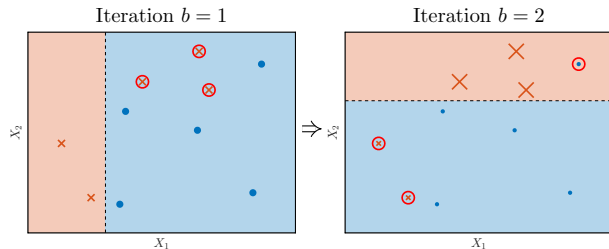
# ex) Boosting illustration



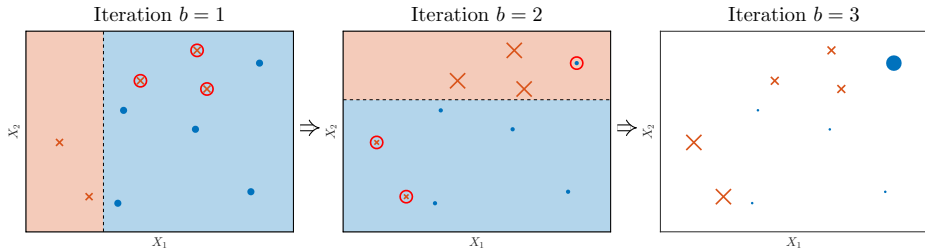
# ex) Boosting illustration



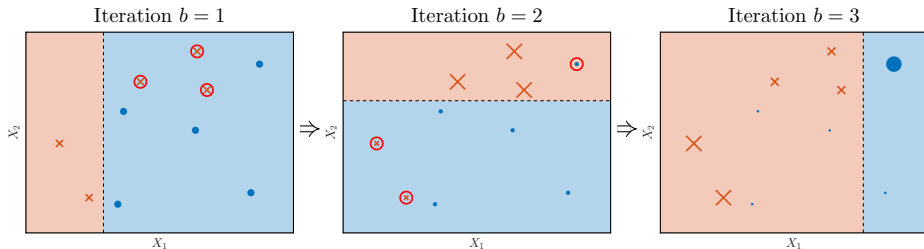
# ex) Boosting illustration



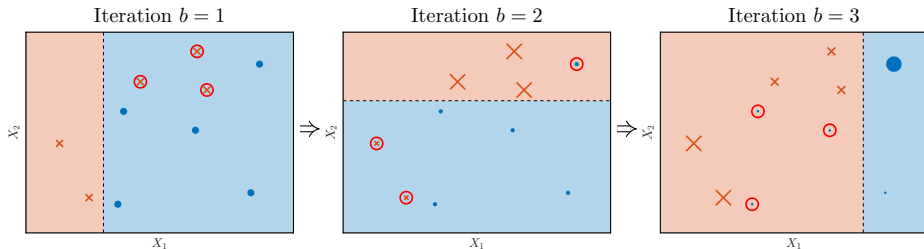
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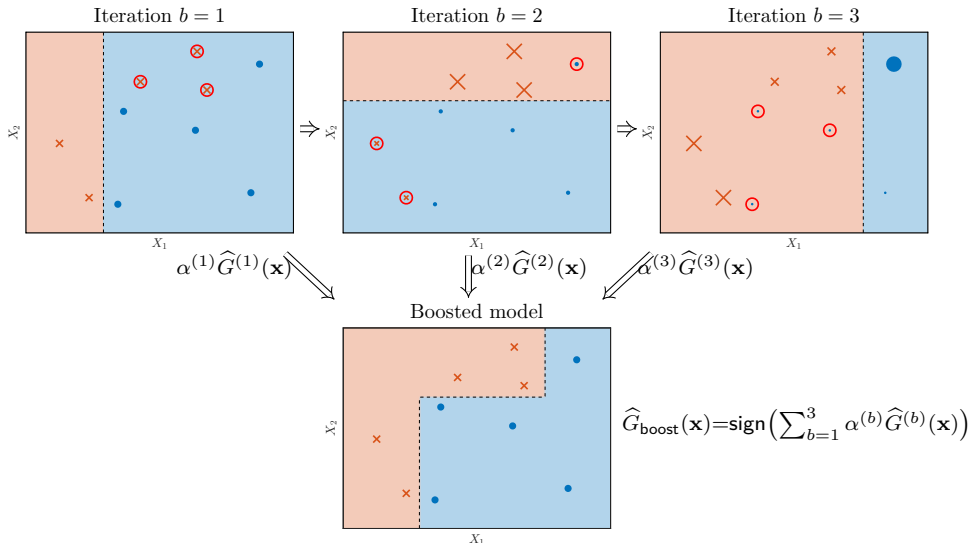


# ex) Boosting illustration





# ex) Boosting illustration



# The technical details. . .

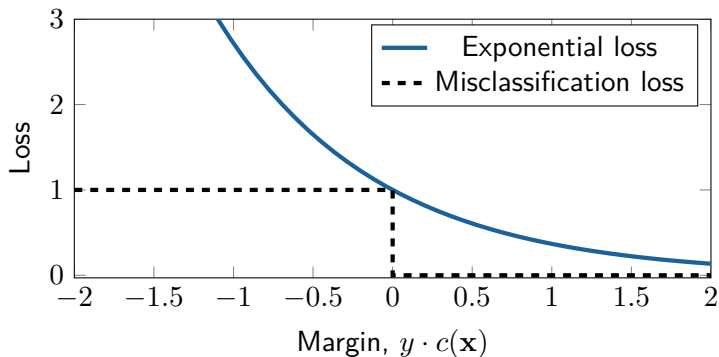
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**Q1:** How do we reweight the data?

**Q2:** How are the coefficients  $\alpha^{(1)}, \dots, \alpha^{(B)}$  computed?

# Exponential loss

Loss functions for binary classifier  $\hat{G}(\mathbf{x}) = \text{sign}(c(\mathbf{x}))$ .



Exponential loss function  $L(y, c(\mathbf{x})) = \exp(-y \cdot c(\mathbf{x}))$  plotted vs. **margin**  $y \cdot c(\mathbf{x})$ . The misclassification loss  $\mathbb{I}\{y \neq \hat{G}(\mathbf{x})\} = \mathbb{I}\{y \cdot c(\mathbf{x}) < 0\}$  is plotted as comparison.

# AdaBoost pseudo-code

---

1. Assign weights  $w_i^1 = 1/n$  to all data points.
2. For  $b = 1$  to  $B$ 
  - (a) Train a weak classifier  $\hat{G}^{(b)}(\mathbf{x})$  on the **weighted training data**  $\{(\mathbf{x}_i, y_i, w_i^b)\}_{i=1}^n$ .
  - (b) *Update the weights*  $\{w_i^{b+1}\}_{i=1}^n$  *from*  $\{w_i^b\}_{i=1}^n$ :
    - i. Increase weights for all points misclassified by  $\hat{G}^{(b)}(\mathbf{x})$ .
    - ii. Decrease weights for all points correctly classified by  $\hat{G}^{(b)}(\mathbf{x})$ .

# AdaBoost pseudo-code

---

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  - (b) *Update the weights*  $\{w_i^{b+1}\}_{i=1}^n$  *from*  $\{w_i^b\}_{i=1}^n$ :
    - i. Compute weighted classification error
    - ii. Compute classifier “confidence”
    - iii. Compute new weights

# AdaBoost pseudo-code

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  - (b) *Update the weights*  $\{w_i^{b+1}\}_{i=1}^n$  *from*  $\{w_i^b\}_{i=1}^n$ :
    - i. Compute  $E_{\text{train}}^b = \sum_{i=1}^n w_i^b \mathbb{I}\{y_i \neq \hat{G}^{(b)}(\mathbf{x}_i)\}$
    - ii. Compute classifier “confidence”
    - iii. Compute new weights

# AdaBoost pseudo-code

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    - i. Compute  $E_{\text{train}}^b = \sum_{i=1}^n w_i^b \mathbb{I}\{y_i \neq \hat{G}^{(b)}(\mathbf{x}_i)\}$
    - ii. Compute  $\alpha^b = 0.5 \log((1 - E_{\text{train}}^b)/E_{\text{train}}^b)$ .
    - iii. Compute new weights

# AdaBoost pseudo-code

---

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    - iii. Compute  $w_i^{b+1} = w_i^b \exp(-\alpha^{(b)} y_i \hat{G}^{(b)}(\mathbf{x}_i))$ ,  $i = 1, \dots, n$



# AdaBoost pseudo-code

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    - ii. Compute  $\alpha^b = 0.5 \log((1 - E_{\text{train}}^b)/E_{\text{train}}^b)$ .
    - iii. Compute  $w_i^{b+1} = w_i^b \exp(-\alpha^{(b)} y_i \hat{G}^{(b)}(\mathbf{x}_i))$ ,  $i = 1, \dots, n$
    - iv. *Normalize*. Set  $w_i^{b+1} \leftarrow w_i^{b+1} / \sum_{j=1}^n w_j^{b+1}$ , for  $i = 1, \dots, n$ .

# AdaBoost pseudo-code

---

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3. Output  $\hat{G}_{\text{boost}}^{(B)}(\mathbf{x}) = \text{sign}\left(\sum_{b=1}^B \alpha^{(b)} \hat{G}^{(b)}(\mathbf{x})\right)$ .

# AdaBoost pseudo-code

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Y. Freund and R. E. Schapire. **Experiments with a New Boosting Algorithm**. *Proceedings of the 13th International Conference on Machine Learning (ICML)*. Bari, Italy, 1996.



2003 Gödel Prize

# Boosting vs. bagging

Bagging	Boosting
Learns base models in parallel	Learns base models sequentially
Uses bootstrapped datasets	Uses reweighted datasets
Does not overfit as $B$ becomes large	Can overfit as $B$ becomes large
Reduces variance but not bias (requires deep trees as base models)	Also reduces bias! (works well with shallow trees)

**N.B.** Boosting does *not* require each base model to have low bias. Thus, a shallow classification tree (say, 4-8 terminal nodes) or even a tree with a single split (2 terminal nodes, a “stump”) is often sufficient.

# A few concepts to summarize lecture 6

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**CART:** Classification and regression trees. A class of nonparametric methods based on partitioning the input space into regions and fitting a simple model for each region.

**Recursive binary splitting:** A greedy method for partitioning the input space into “boxes” aligned with the coordinate axes.

**Gini index and deviance:** Commonly used error measures for constructing classification trees.

**Ensemble methods:** Umbrella term for methods that average or combine multiple models.

**Bagging:** Bootstrap aggregating. An ensemble method based on the statistical bootstrap.

**Random forests:** Bagging of trees, combined with random feature selection for further variance reduction (and computational gains).