

# Multiclass Classification, Structured Prediction, & Decision Trees

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# Margin for Multiclass

- Binary
- Margin for  $(x^{(n)}, y^{(n)})$ :

$$y^{(n)} w^T x^{(n)} \quad (1)$$

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- Multiclass
- Class-specific margin for  $(x^{(n)}, y^{(n)})$ :

$$h(x^{(n)}, y^{(n)}) - h(x^{(n)}, y). \quad (2)$$

- Difference between scores of the correct class and each other class
- Want margin to be large and positive for all  $y \neq y^{(n)}$ .

# Multiclass SVM: separable case

## Binary

$$\min_w \quad \frac{1}{2} \|w\|^2 \quad (3)$$

$$\text{s.t.} \quad \underbrace{y^{(n)} w^T x^{(n)}}_{\text{margin}} \geq 1 \quad \forall (x^{(n)}, y^{(n)}) \in \mathcal{D} \quad (4)$$

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**Multiclass** As in the binary case, take 1 as our target margin.

$$m_{n,y}(w) \stackrel{\text{def}}{=} \underbrace{\langle w, \Psi(x^{(n)}, y^{(n)}) \rangle}_{\text{score of correct class}} - \underbrace{\langle w, \Psi(x^{(n)}, y) \rangle}_{\text{score of other class}} \quad (5)$$

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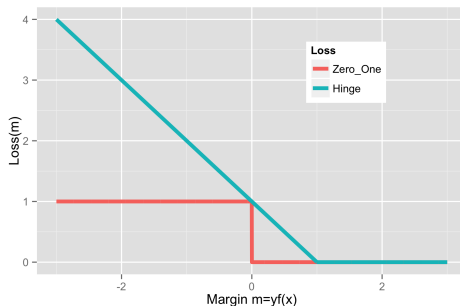
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**Exercise:** write the objective for the non-separable case

## Recap: hinge loss for binary classification

- Hinge loss: a convex upperbound on the 0-1 loss

$$\ell_{\text{hinge}}(y, \hat{y}) = \max(0, 1 - yh(x)) \quad (8)$$



## Generalized hinge loss

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- Upper bound on  $\Delta(y, y')$ .

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- Generalized hinge loss:

$$\ell_{\text{hinge}}(y, x, w) \stackrel{\text{def}}{=} \max_{y' \in \mathcal{Y}} (\Delta(y, y') - \langle w, (\Psi(x, y) - \Psi(x, y')) \rangle) \quad (13)$$

# Multiclass SVM with Hinge Loss

- Recall the hinge loss formulation for binary SVM (without the bias term):

$$\min_{w \in \mathbb{R}^d} \frac{1}{2} \|w\|^2 + C \sum_{n=1}^N \max \left( 0, 1 - \underbrace{y^{(n)} w^T x^{(n)}}_{\text{margin}} \right).$$

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- The multiclass objective:

$$\min_{w \in \mathbb{R}^d} \frac{1}{2} \|w\|^2 + C \sum_{n=1}^N \max_{y' \in \mathcal{Y}} \left( \Delta(y, y') - \underbrace{\langle w, (\Psi(x, y) - \Psi(x, y')) \rangle}_{\text{margin}} \right)$$

- $\Delta(y, y')$  as **target margin** for each class.
- If margin  $m_{n,y'}(w)$  meets or exceeds its target  $\Delta(y^{(n)}, y') \forall y \in \mathcal{Y}$ , then no loss on example  $n$ .



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  - Predict with  $\arg \max_{y \in \mathcal{Y}} h_y(x)$ .
  - Gave simple example where this fails for linear classifiers
- Solution 2: Multiclass loss
  - Train one model:  $h(x, y) : \mathcal{X} \times \mathcal{Y} \rightarrow \mathbb{R}$ .
  - Prediction involves solving  $\arg \max_{y \in \mathcal{Y}} h(x, y)$ .

# Does it work better in practice?

- Paper by Rifkin & Klautau: “In Defense of One-Vs-All Classification” (2004)
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    - (or at least, the advantages claimed by earlier papers for multiclass methods were not compelling)
- Compared
  - many multiclass frameworks (including the one we discuss)
  - one-vs-all for SVMs with RBF kernel
  - one-vs-all for square loss with RBF kernel (for classification!)
- All performed roughly the same

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- Generalizes to situations where  $k$  is very large and one-vs-all is intractable.
- Key idea is that we can generalize across outputs  $y$  by using features of  $y$ .



# Introduction to Structured Prediction

## Example: Part-of-speech (POS) Tagging

- Given a sentence, give a part of speech tag for each word:

$x$	$\underbrace{[\text{START}]}_{x_0}$	$\underbrace{\text{He}}_{x_1}$	$\underbrace{\text{eats}}_{x_2}$	$\underbrace{\text{apples}}_{x_3}$
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- $\mathcal{P} = \{\text{START, Pronoun, Verb, Noun, Adjective}\}$
- $\mathcal{Y} = \mathcal{P}^n, n = 1, 2, 3, \dots$  [Part of speech sequence of any length]

# Multiclass Hypothesis Space

- Discrete output space:  $\mathcal{Y}(x)$ 
  - Very large but has structure, e.g., linear chain (sequence labeling), tree (parsing)
  - Size depends on input  $x$

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- Base Hypothesis Space:  $\mathcal{H} = \{h: \mathcal{X} \times \mathcal{Y} \rightarrow \mathbb{R}\}$ 
  - $h(x, y)$  gives **compatibility score** between input  $x$  and output  $y$

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  - $h(x, y)$  gives **compatibility score** between input  $x$  and output  $y$
- Multiclass hypothesis space

$$\mathcal{F} = \left\{ x \mapsto \arg \max_{y \in \mathcal{Y}} h(x, y) \mid h \in \mathcal{H} \right\}$$

- Final prediction function is an  $f \in \mathcal{F}$ .
- For each  $f \in \mathcal{F}$  there is an underlying compatibility score function  $h \in \mathcal{H}$ .

# Structured Prediction

- Part-of-speech tagging

$x$ :	he	eats	apples
$y$ :	pronoun	verb	noun

- Multiclass hypothesis space:

$$h(x, y) = w^T \Psi(x, y) \quad (14)$$

$$\mathcal{F} = \left\{ x \mapsto \arg \max_{y \in \mathcal{Y}} h(x, y) \mid h \in \mathcal{H} \right\} \quad (15)$$

- A special case of multiclass classification
- How to design the feature map  $\Psi$ ? What are the considerations?



# Unary features

- A **unary feature** only depends on
  - the label at a **single position**,  $y_i$ , and  $x$
- Example:

$$\phi_1(x, y_i) = \mathbb{1}[x_i = \text{runs}] \mathbb{1}[y_i = \text{Verb}]$$

$$\phi_2(x, y_i) = \mathbb{1}[x_i = \text{runs}] \mathbb{1}[y_i = \text{Noun}]$$

$$\phi_3(x, y_i) = \mathbb{1}[x_{i-1} = \text{He}] \mathbb{1}[x_i = \text{runs}] \mathbb{1}[y_i = \text{Verb}]$$

# Markov features

- A **markov feature** only depends on
  - two **adjacent** labels,  $y_{i-1}$  and  $y_i$ , and  $x$
- Example:

$$\theta_1(x, y_{i-1}, y_i) = \mathbb{1}[y_{i-1} = \text{Pronoun}] \mathbb{1}[y_i = \text{Verb}]$$

$$\theta_2(x, y_{i-1}, y_i) = \mathbb{1}[y_{i-1} = \text{Pronoun}] \mathbb{1}[y_i = \text{Noun}]$$

- Reminiscent of Markov models in the output space
- Possible to have higher-order features

# Local Feature Vector and Compatibility Score

- At each position  $i$  in sequence, define the **local feature vector** (unary and markov):

$$\Psi_i(x, y_{i-1}, y_i) = (\phi_1(x, y_i), \phi_2(x, y_i), \dots, \theta_1(x, y_{i-1}, y_i), \theta_2(x, y_{i-1}, y_i), \dots)$$

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- And **local compatibility score** at position  $i$ :  $\langle w, \Psi_i(x, y_{i-1}, y_i) \rangle$ .
- The compatibility score for  $(x, y)$  is the sum of local compatibility scores:

$$\sum_i \langle w, \Psi_i(x, y_{i-1}, y_i) \rangle = \left\langle w, \sum_i \Psi_i(x, y_{i-1}, y_i) \right\rangle = \langle w, \Psi(x, y) \rangle, \quad (16)$$

where we define the **sequence feature vector** by

$$\Psi(x, y) = \sum_i \Psi_i(x, y_{i-1}, y_i). \quad \text{decomposable}$$

# Structured perceptron

Given a dataset  $\mathcal{D} = \{(x, y)\}$ ;

Initialize  $w \leftarrow 0$ ;

**for**  $iter = 1, 2, \dots, T$  **do**

**for**  $(x, y) \in \mathcal{D}$  **do**

$\hat{y} = \arg \max_{y' \in \mathcal{Y}(x)} w^T \psi(x, y')$ ;

**if**  $\hat{y} \neq y$  **then** // We've made a mistake

$w \leftarrow w + \Psi(x, y)$  ; // Move the scorer towards  $\psi(x, y)$

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Identical to the multiclass perceptron algorithm except the  $\arg \max$  is now over the structured output space  $\mathcal{Y}(x)$ .

# Structured hinge loss

- Recall the generalized hinge loss

$$\ell_{\text{hinge}}(y, \hat{y}) \stackrel{\text{def}}{=} \max_{y' \in \mathcal{Y}(x)} (\Delta(y, y') + \langle w, (\Psi(x, y') - \Psi(x, y)) \rangle) \quad (17)$$

- What is  $\Delta(y, y')$  for two sequences?



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- What is  $\Delta(y, y')$  for two sequences?
- Hamming loss** is common:

$$\Delta(y, y') = \frac{1}{L} \sum_{i=1}^L \mathbb{1}[y_i \neq y'_i]$$

where  $L$  is the sequence length.

## Exercise:

- Write down the objective of structured SVM using the structured hinge loss.
- Stochastic sub-gradient descent for structured SVM (similar to HW3 P3)
- Compare with the structured perceptron algorithm

# The argmax problem for sequences

**Problem** To compute predictions, we need to find  $\arg \max_{y \in \mathcal{Y}(x)} \langle w, \Psi(x, y) \rangle$ , and  $|\mathcal{Y}(x)|$  is exponentially large.

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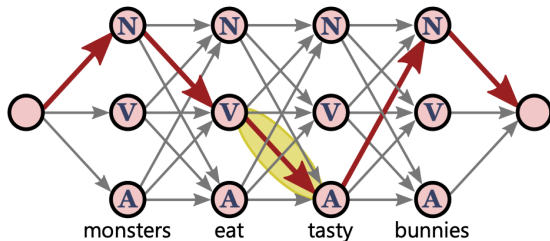
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**Solution** Dynamic programming (similar to the Viterbi algorithm)



What's the running time?

# Conditional random field (CRF)

- Recall that we can write logistic regression in a general form:

$$p(y|x) = \frac{1}{Z(x)} \exp(w^\top \psi(x, y)).$$

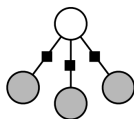
- $Z$  is normalization constant:  $Z(x) = \sum_{y \in Y} \exp(w^\top \psi(x, y))$ .

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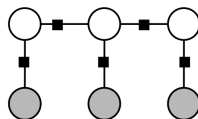
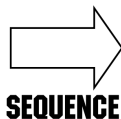
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- $Z$  is normalization constant:  $Z(x) = \sum_{y \in Y} \exp(w^\top \psi(x, y))$ .
- Example: linear chain  $\{y_t\}$
- We can incorporate unary and Markov features:  $p(y|x) = \frac{1}{Z(x)} \exp(\sum_t w^\top \psi(x, y_t, y_{t-1}))$



Logistic Regression



Linear-chain CRFs

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- We can draw samples in the output space.



# Conditional random field (CRF)

- Compared to Structured SVM, CRF has a probabilistic interpretation.
- We can draw samples in the output space.
- How do we learn  $w$ ? Maximum log likelihood, and regularization term:  $\lambda \|w\|^2$
- Loss function:

$$\begin{aligned} l(w) &= -\frac{1}{N} \sum_{i=1}^N \log p(y^{(i)} | x^{(i)}) + \frac{1}{2} \lambda \|w\|^2 \\ &= -\frac{1}{N} \sum_i \sum_t \sum_k w_k \psi_k(y_t^{(i)}, y_{t-1}^{(i)}) + \frac{1}{N} \sum_i \log Z(x^{(i)}) + \frac{1}{2} \sum_k \lambda w_k^2 \end{aligned}$$

# Conditional random field (CRF)

- Loss function:

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- Gradient:

$$\frac{\partial l(w)}{\partial w_k} = -\frac{1}{N} \sum_i \sum_t \sum_k \psi_k(x^{(i)}, y_t^{(i)}, y_{t-1}^{(i)}) \quad (18)$$

$$+ \frac{1}{N} \sum_i \frac{\partial}{\partial w_k} \log \sum_{y' \in Y} \exp\left(\sum_t \sum_{k'} w_{k'} \psi_{k'}(x^{(i)}, y'_t, y'_{t-1})\right) + \sum_k \lambda w_k \quad (19)$$

# Conditional random field (CRF)

- What is  $\frac{1}{N} \sum_i \sum_t \sum_k \psi_k(x^{(i)}, y_t^{(i)}, y_{t-1}^{(i)})$ ?

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- What is  $\frac{1}{N} \sum_i \sum_t \sum_k \psi_k(x^{(i)}, y_t^{(i)}, y_{t-1}^{(i)})$ ?
- It is the expectation  $\psi_k(x^{(i)}, y_t, y_{t-1})$  under the empirical distribution  $\tilde{p}(x, y) = \frac{1}{N} \sum_i \mathbb{1}[x = x^{(i)}] \mathbb{1}[y = y^{(i)}]$ .

## Conditional random field (CRF)

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- It is the expectation of  $\psi_k(x^{(i)}, y'_t, y'_{t-1})$  under the model distribution  $p(y'_t, y'_{t-1} | x)$ .

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- Both problems are NP-hard for general graphs.

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- In general graphs, we rely on approximate inference (e.g. loopy belief propagation).

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An image may contain multiple class labels, e.g. a bus is likely to co-occur with a car.

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- Structured prediction: Structured SVM, CRF. Data containing structure. Extremely large output space. Text and image applications.



# Decision Trees

# Overview: Decision Trees

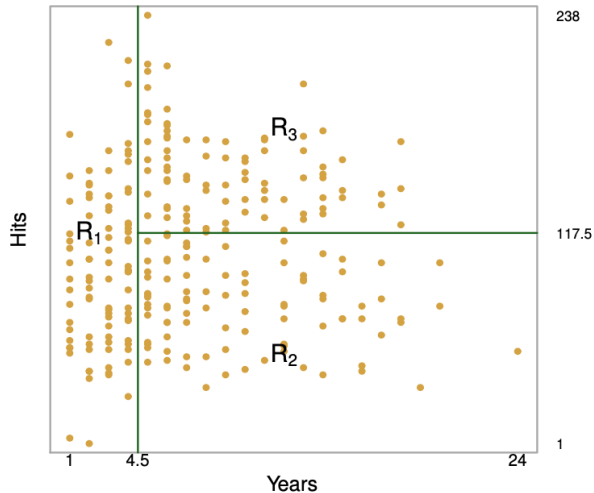
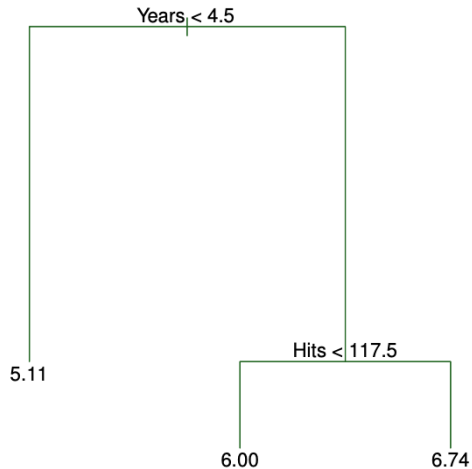
- Our first inherently non-linear classifier: decision trees.
- Ensemble methods: bagging and boosting.

# Decision Trees

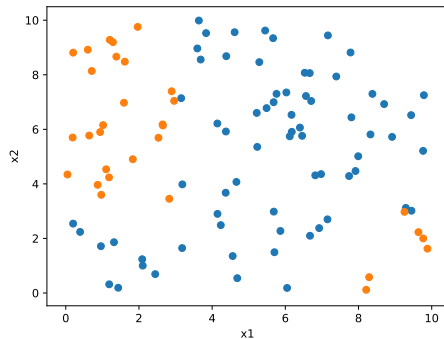
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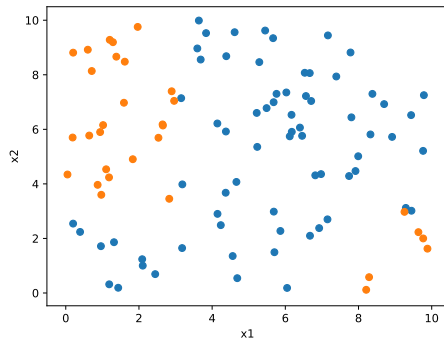


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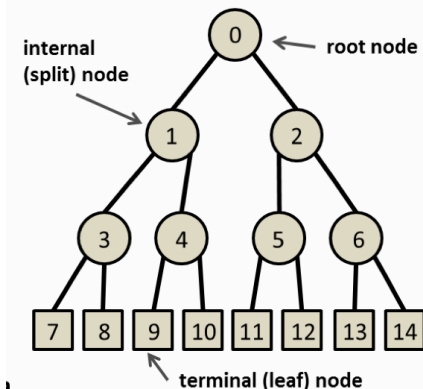
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- Can we classify these points using a linear classifier?
- Partition the data into axis-aligned regions **recursively** (on the board)

# Decision trees setup

## A general tree structure

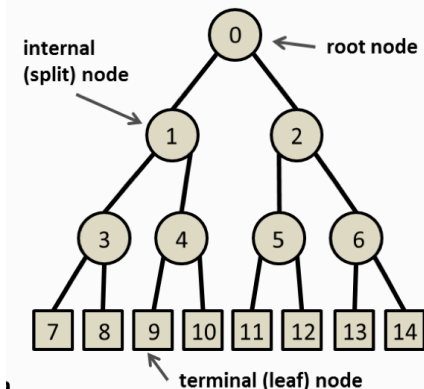


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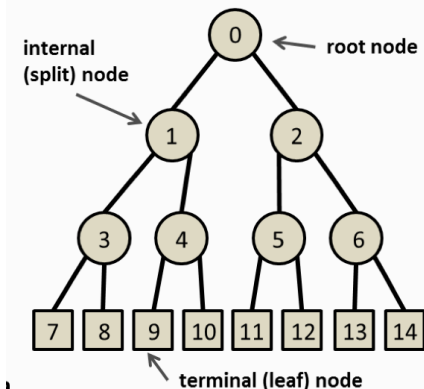
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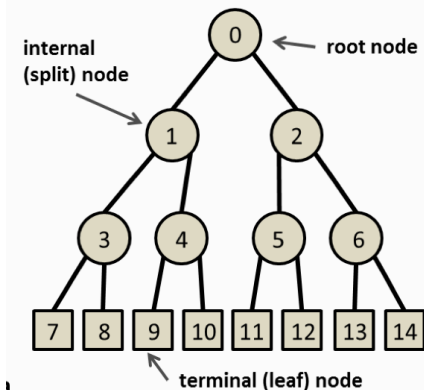
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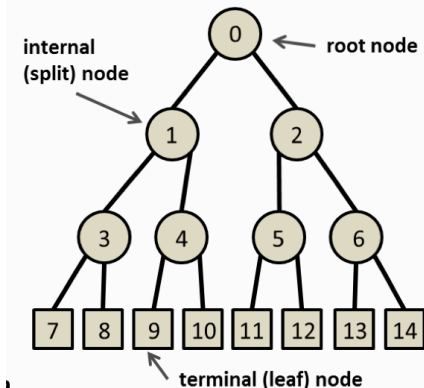
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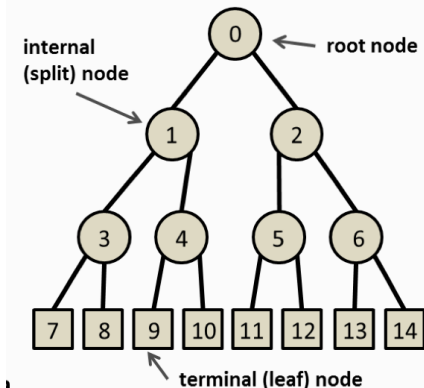
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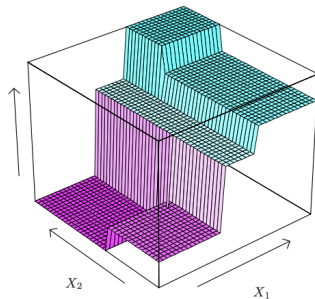
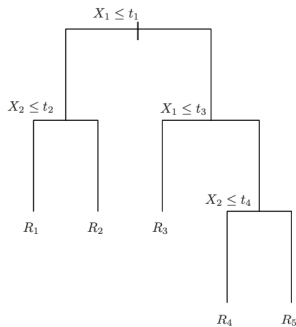
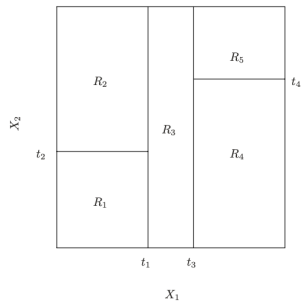
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# Prediction in a Regression Tree



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- It is common to split half way between two adjacent values:

$$s_j \in \left\{ \frac{1}{2} (x_{j(r)} + x_{j(r+1)}) \mid r = 1, \dots, n-1 \right\}. \quad n-1 \text{ splits} \quad (24)$$

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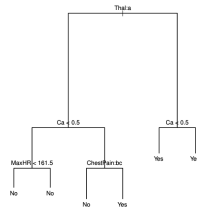
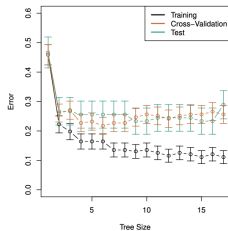
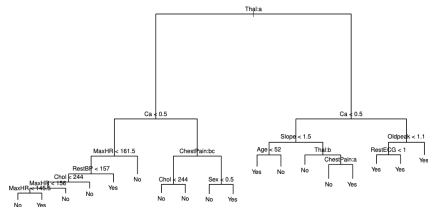
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    - 2 *Prune* the tree back greedily, potentially all the way to the root, until validation performance starts decreasing.

# Pruning: Example



# What Makes a Good Split for Classification?

Our plan is to predict the **majority label** in each region.

Which of the following splits is better?

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Intuition: we want to produce *pure* nodes, i.e. nodes where most instances have the same class.



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- The Gini index encourages  $\hat{p}_{mk}$  to be close to 0 or 1

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- Entropy / Information gain

$$-\sum_{k=1}^K \hat{p}_{mk} \log \hat{p}_{mk}.$$

# Node Impurity Measures

- Three measures of **node impurity** for leaf node  $m$ :

- Misclassification error

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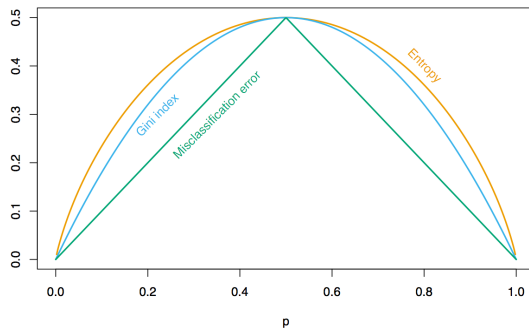
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- The Gini index and entropy are numerically similar to each other, and both work better in practice than the misclassification error.



# Impurity Measures for Binary Classification

( $p$  is the relative frequency of class 1)



HTF Figure 9.3

# Quantifying the Impurity of a Split

Scoring a potential split that produces the nodes  $R_L$  and  $R_R$ :

- Suppose we have  $N_L$  points in  $R_L$  and  $N_R$  points in  $R_R$ .

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- Let  $Q(R_L)$  and  $Q(R_R)$  be the node impurity measures for each node.
- We aim to find a split that minimizes the *weighted average of node impurities*:

$$\frac{N_L Q(R_L) + N_R Q(R_R)}{N_L + N_R}$$

## Discussion: Interpretability of Decision Trees



- Trees are easier to visualize and explain than other classifiers (even linear regression)

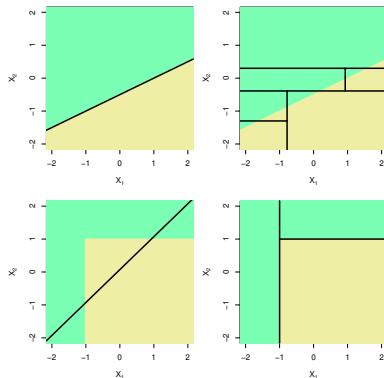
## Discussion: Interpretability of Decision Trees



- Trees are easier to visualize and explain than other classifiers (even linear regression)
- Small trees are interpretable – large trees, maybe not so much

## Discussion: Trees vs. Linear Models

Trees may have to work hard to capture linear decision boundaries, but can easily capture certain nonlinear ones:



## Discussion: Review

Decision trees are:

- Non-linear: the decision boundary that results from splitting may end up being quite complicated
- Non-metric: they do not rely on the geometry of the space (inner products or distances)
- Non-parametric: they make no assumptions about the distribution of the data



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Additional pros:

- Interpretable and simple to understand

Cons:

- Struggle to capture linear decision boundaries
- They have high variance and tend to **overfit**: they are sensitive to small changes in the training data (The ensemble techniques we discuss next can mitigate these issues)

# Bagging and Random Forests

## Recap: Statistics and Point Estimators

- We observe data  $\mathcal{D} = (x_1, x_2, \dots, x_n)$  sampled i.i.d. from a parametric distribution  $p(\cdot \mid \theta)$

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  - E.g., sample mean, sample variance, histogram, empirical data distribution
- A statistic  $\hat{\theta} = \hat{\theta}(\mathcal{D})$  is a **point estimator** of  $\theta$  if  $\hat{\theta} \approx \theta$

## Recap: Bias and Variance of an Estimator

- Statistics are random, so they have probability distributions.
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- Why does variance matter if an estimator is unbiased?
  - $\hat{\theta}(\mathcal{D}) = x_1$  is an unbiased estimator of the mean of a Gaussian, but would be farther away from  $\theta$  than the sample mean.

## Variance of a Mean

- Let  $\hat{\theta}(\mathcal{D})$  be an unbiased estimator with variance  $\sigma^2$ :  $\mathbb{E}[\hat{\theta}] = \theta$ ,  $\text{Var}(\hat{\theta}) = \sigma^2$ .
- So far we have used a single statistic  $\hat{\theta} = \hat{\theta}(\mathcal{D})$  to estimate  $\theta$ .

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- The average has the same expected value but smaller standard error (recall that  $\text{Var}(cX) = c^2 \text{Var}(X)$ , and that the  $\hat{\theta}_i$ -s are uncorrelated):

$$\mathbb{E}\left[\frac{1}{n} \sum_{i=1}^n \hat{\theta}_i\right] = \theta \quad \text{Var}\left[\frac{1}{n} \sum_{i=1}^n \hat{\theta}_i\right] = \frac{\sigma^2}{n} \quad (25)$$

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- We will define the average prediction function as:

$$\hat{f}_{\text{avg}} \stackrel{\text{def}}{=} \frac{1}{B} \sum_{b=1}^B \hat{f}_b \quad (26)$$

# Averaging Reduces Variance of Predictions

- The average prediction for  $x_0$  is

$$\hat{f}_{\text{avg}}(x_0) = \frac{1}{B} \sum_{b=1}^B \hat{f}_b(x_0).$$

- $\hat{f}_{\text{avg}}(x_0)$  and  $\hat{f}_b(x_0)$  have the same expected value, but
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- **Problem:** in practice we don't have  $B$  independent training sets!

# The Bootstrap Sample

How do we simulate multiple samples when we only have one?

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- For large  $n$ ,

$$\left(1 - \frac{1}{n}\right)^n \approx \frac{1}{e} \approx .368. \quad (27)$$

- So we expect ~63.2% of elements of  $\mathcal{D}_n$  will show up at least once.

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- Use these values as though  $D_n^1, \dots, D_n^B$  were i.i.d. samples from  $P$ .
- This often ends up being very close to what we'd get with independent samples from  $P$ !

# Independent Samples vs. Bootstrap Samples

- Point estimator  $\hat{\alpha} = \hat{\alpha}(\mathcal{D}_{100})$  for samples of size 100, for a synthetic case where the data generating distribution is known
- Histograms of  $\hat{\alpha}$  based on
  - 1000 independent samples of size 100 (left), vs.
  - 1000 bootstrap samples of size 100 (right)

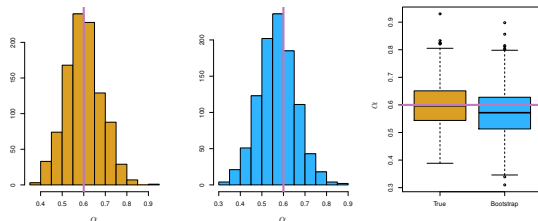


Figure 5.10 from *ISLR* (Springer, 2013) with permission from the authors: G. James, D. Witten, T. Hastie and R. Tibshirani.

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  - We try to find new learners that do well where previous learners fall short

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- The **bagged prediction function** is a *combination* of these:

$$\hat{f}_{\text{avg}}(x) = \text{Combine} \left( \hat{f}_1(x), \hat{f}_2(x), \dots, \hat{f}_B(x) \right)$$



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- Is there a downside, compared to having a single decision tree?
- Yes: if we have many trees, the bagged predictor is much less interpretable

## Aside: Out-of-Bag Error Estimation

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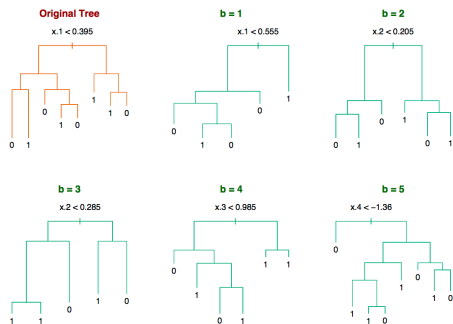
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- The OOB error is a good estimate of the test error
- Similar to cross validation error: both are computed on the training set



# Applying Bagging to Classification Trees

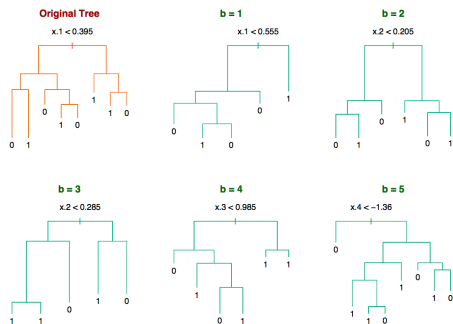
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From HTF Figure 8.9

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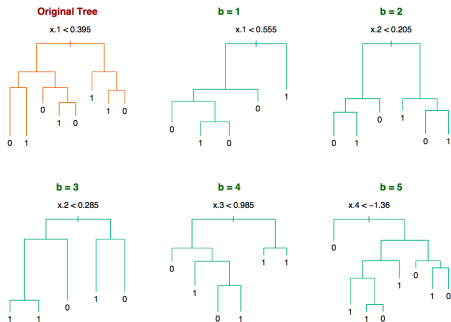


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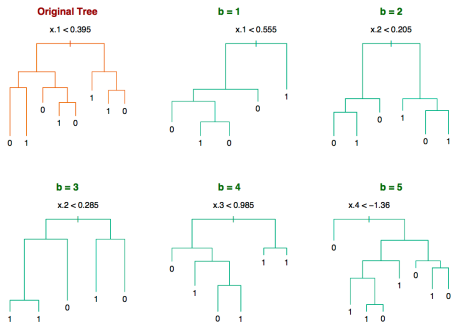


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- Each bootstrap tree is quite different: different splitting variable at the root!
- High variance:** small perturbations of the training data lead to a high degree of model variability
- Bagging helps most when the base learners are relatively unbiased but have high variance (exactly the case for decision trees)

From HTF Figure 8.9

# Motivating Random Forests: Correlated Prediction Functions

Recall the motivating principle of bagging:

- For  $\hat{\theta}_1, \dots, \hat{\theta}_n$  *i.i.d.* with  $\mathbb{E}[\hat{\theta}] = \theta$  and  $\text{Var}[\hat{\theta}] = \sigma^2$ ,

$$\mathbb{E}\left[\frac{1}{n}\sum_{i=1}^n \hat{\theta}_i\right] = \mu \quad \text{Var}\left[\frac{1}{n}\sum_{i=1}^n \hat{\theta}_i\right] = \frac{\sigma^2}{n}.$$

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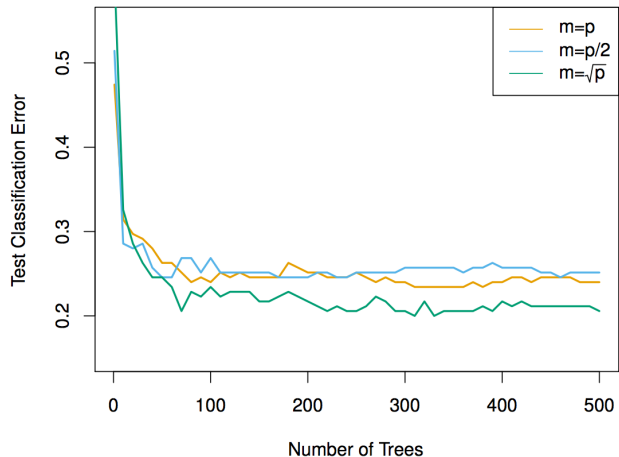
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# Random Forests: Effect of $m$



From *An Introduction to Statistical Learning, with applications in R* (Springer, 2013) with permission from the authors: G. James, D. Witten, T. Hastie and R. Tibshirani.



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- But bootstrap samples (and the induced models) are correlated
- Ensembling works better when we combine a diverse set of prediction functions
  - $\implies$  Random forests: select a random subset of features for each decision tree