CSCI567 Machine Learning (Spring 2025)

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Feb 28, 2025

Quiz 1 Logistics

Date: Friday, March 7th

Time: 1:00-3:20pm

Location: THH 101 (double seating), waiting for an overflow room (will announce on Piazza)

Individual effort, close-book (no cheat sheet), no calculators or any other electronics, *but need your phone to upload your solutions to Gradescope from 3:20-3:30pm*

Quiz 1 Coverage

Coverage: mostly Lec 1-6, some multiple-choice questions from Lec 7.

#Pages: $\sim$19 (but a lot of empty space!) + 5 for scratch work.

Five problems in total

- one problem of 15 multiple-choice *multiple-answer* questions
  - 0.5 point for selecting (not selecting) each correct (incorrect) answer
  - "which of the following is correct?" *does not imply one correct answer*
- four other homework-like problems, each has a couple sub-problems
  - linear regression, linear classifiers, backpropagation, kernel, SVM

Tips: expect to see variants of questions from discussion/homework

Quiz 1 Coverage

Formulas provided:

- chain rule, activation functions, SVM dual formulation, entropy definition

Basic stuff that will not be provided:

- SGD, softmax, CNN dimension/#parameters calculations, etc.

Most other formulas are not needed (*do not overthink!*)

- Taylor expansion, complete Backprop algorithm in matrix form, KKT conditions, etc.

Outline

Outline

Support Vector Machine

SVM: **max-margin linear classifier**

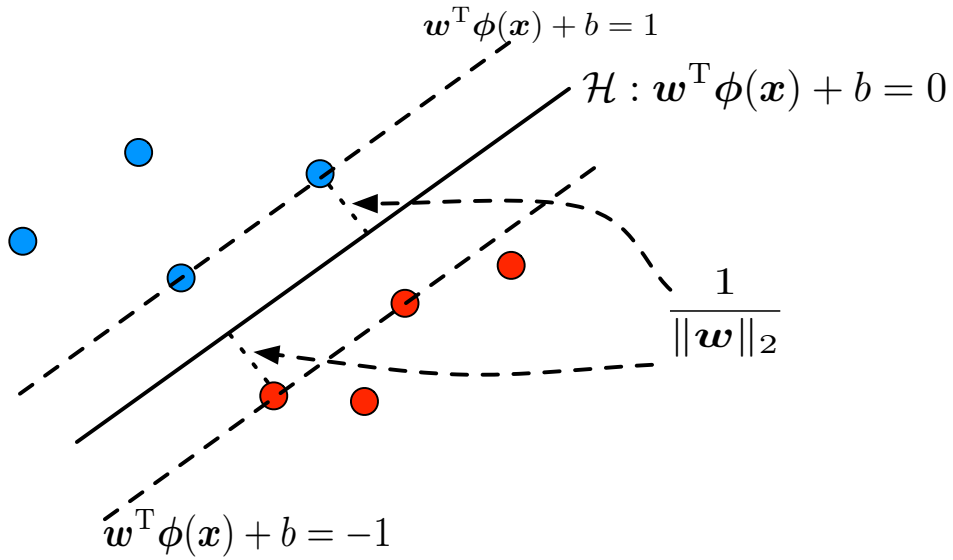
Primal (equivalent to minimizing L2 regularized hinge loss):

$$\begin{aligned} \min_{\boldsymbol{w},b,\{\xi_n\}} \quad & C\sum_n \xi_n + \frac{1}{2}\|\boldsymbol{w}\|_2^2 \\ \text{s.t.} \quad & 1 - y_n(\boldsymbol{w}^{\mathrm{T}}\boldsymbol{\phi}(\boldsymbol{x}_n) + b) \le \xi_n, \quad \forall\ n \\ & \xi_n \ge 0, \quad \forall\ n \end{aligned}$$

Dual (kernelizable, reveals what training points are support vectors):

$$\begin{aligned} \max_{\{\alpha_n\}} \quad & \sum_n \alpha_n - \frac{1}{2}\sum_{m,n} y_m y_n \alpha_m \alpha_n \boldsymbol{\phi}(\boldsymbol{x}_m)^{\mathrm{T}}\boldsymbol{\phi}(\boldsymbol{x}_n) \\ \text{s.t.} \quad & \sum_n \alpha_n y_n = 0 \quad \text{and} \quad 0 \le \alpha_n \le C, \quad \forall\ n \end{aligned}$$

Separable Case



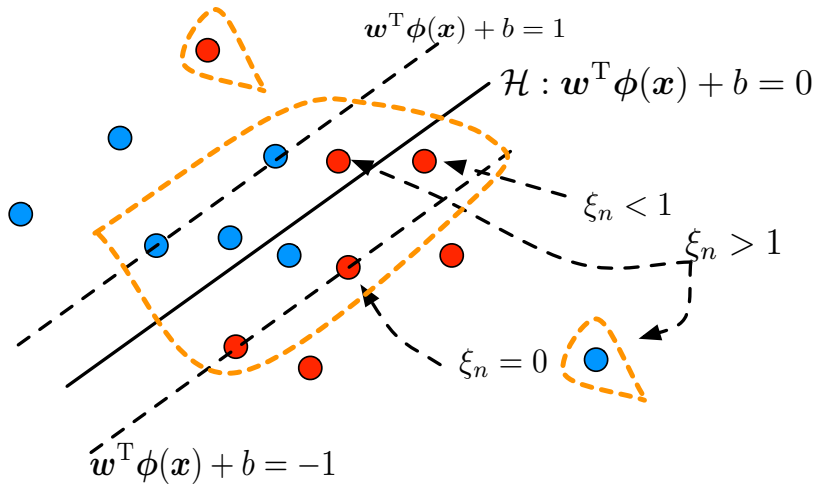
Geometric interpretation of support vectors

A support vector satisfies $\alpha_n^* \neq 0$ and

$$1 - \xi_n^* - y_n(\boldsymbol{w}^{*\mathrm{T}}\boldsymbol{\phi}(\boldsymbol{x}_n) + b^*) = 0$$

When

- $\xi_n^* = 0$, $y_n(\boldsymbol{w}^{*\mathrm{T}}\boldsymbol{\phi}(\boldsymbol{x}_n) + b^*) = 1$ and thus the point is $1/\|\boldsymbol{w}^*\|_2$ away from the hyperplane.
- $\xi_n^* < 1$, the point is classified correctly but does not satisfy the large margin constraint.
- $\xi_n^* > 1$, the point is misclassified.



Support vectors (circled with the orange line) are *the only points that matter!*

Outline

Decision tree

We have seen different ML models for classification/regression:

- linear models, neural nets and other nonlinear models induced by kernels

Decision tree is yet another one:

- nonlinear in general
- works for both classification and regression; we focus on classification
- one key advantage is good interpretability
- not to be confused with the "tree reduction" in Lec 4
- still very popular for small tabular data, especially when used in ensemble (i.e. "forest")

Tree-based models outperform neural nets sometimes

Why do tree-based models still outperform deep learning on tabular data?

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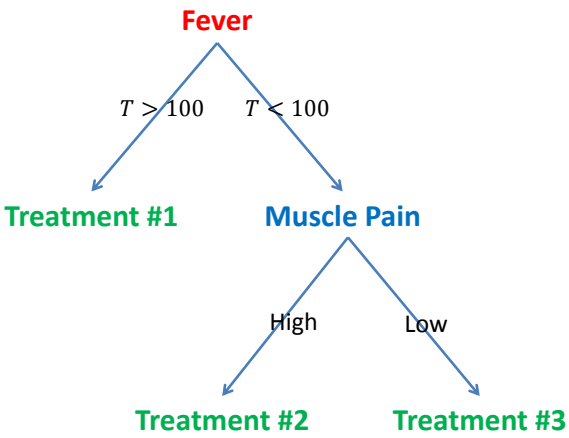
Abstract

While deep learning has enabled tremendous progress on text and image datasets, its superiority on tabular data is not clear. We contribute extensive benchmarks of standard and novel deep learning methods as well as tree-based models such as XGBoost and Random Forests, across a large number of datasets and hyperparameter combinations. We define a standard set of 45 datasets from varied domains with clear characteristics of tabular data and a benchmarking methodology accounting for both fitting models and finding good hyperparameters. Results show that tree-based models remain state-of-the-art on medium-sized data (∼10K samples) even without accounting for their superior speed. To understand this gap, we conduct an empirical investigation into the differing inductive biases of tree-based models and Neural Networks (NNs). This leads to a series of challenges which should guide researchers aiming to build tabular-specific NNs: **1.** be robust to uninformative features, **2.** preserve the orientation of the data, and **3.** be able to easily learn irregular functions. To stimulate research on tabular architectures, we contribute a standard benchmark and raw data for baselines: every point of a 20 000 compute hours hyperparameter search for each learner.

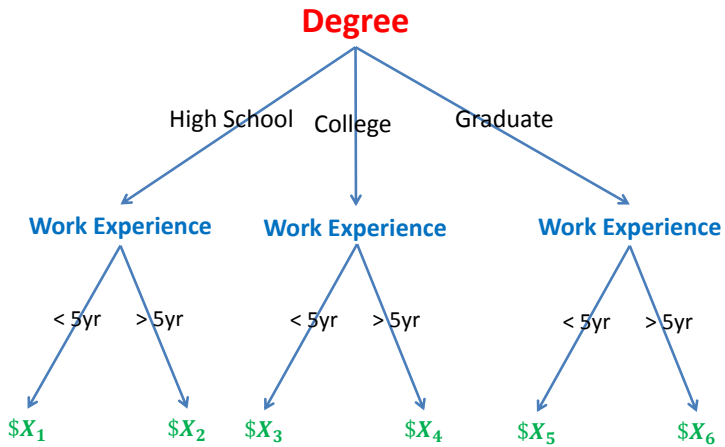
Example

Many decisions are made based on some tree structure

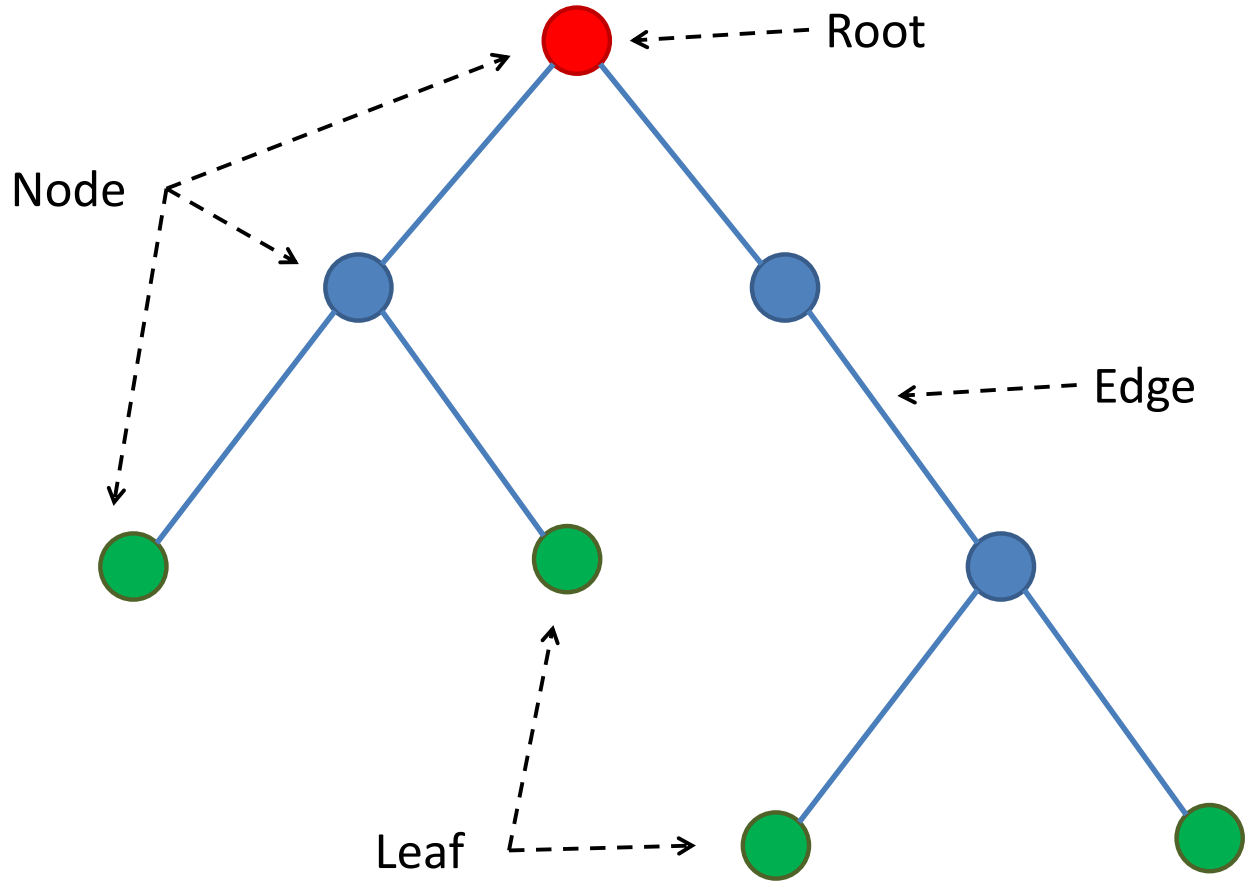
Medical treatment



Salary in a company



Tree terminology

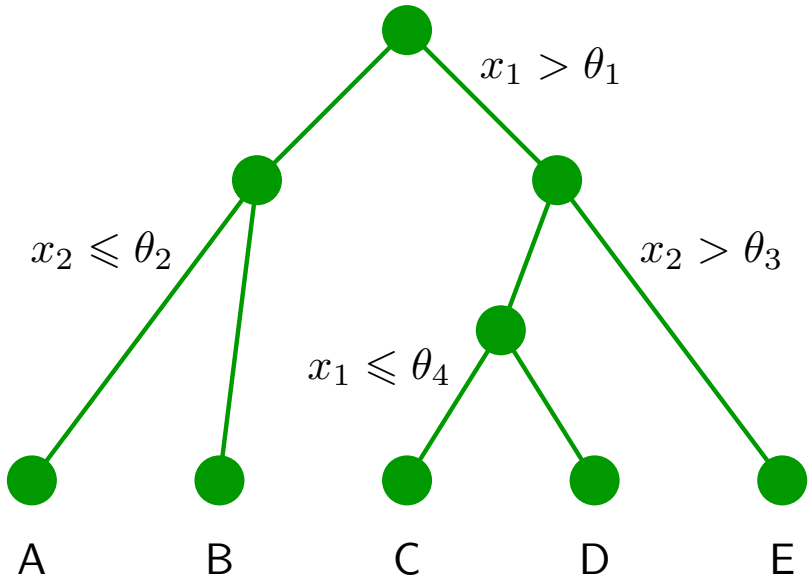


A more abstract example of decision trees

Input: $\boldsymbol{x} = (x_1, x_2)$

Output: $f(\boldsymbol{x})$ determined naturally by traversing the tree

- start from the root
- test at each node to decide which child to visit next
- finally the leaf gives the prediction $f(\boldsymbol{x})$

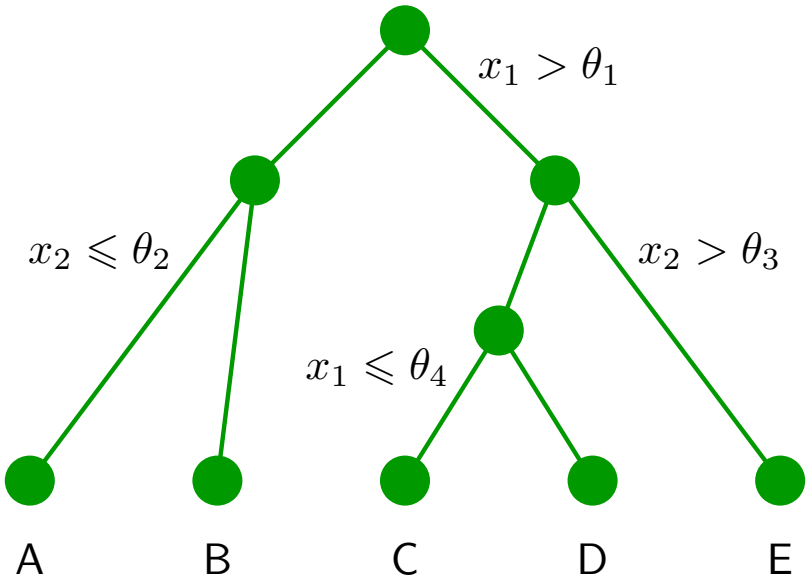


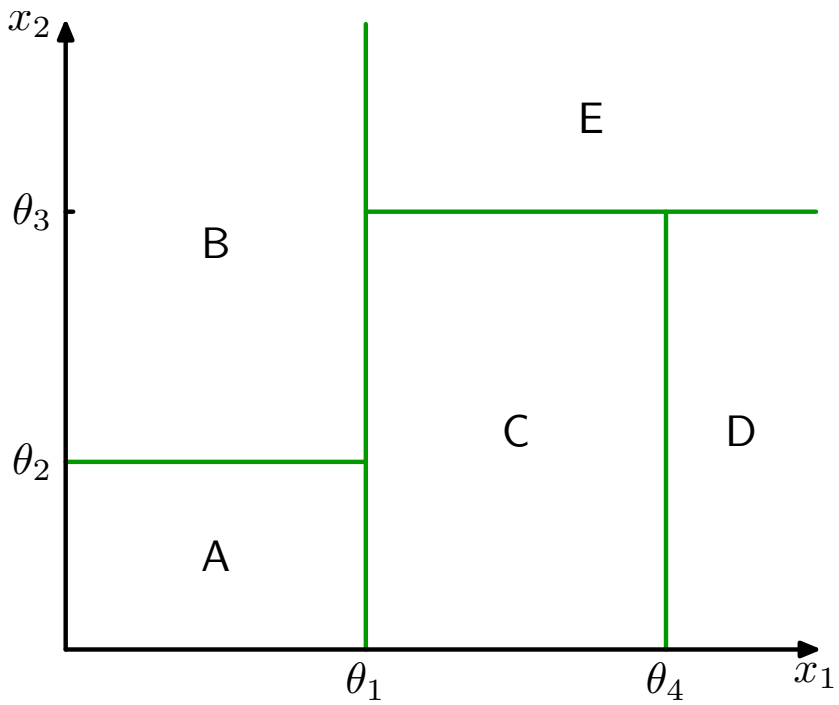
For example, $f((\theta_1 - 1, \theta_2 + 1)) = \text{B}$

Complex to formally write down, but easy to represent pictorially or as codes.

The decision boundary

Corresponds to a classifier with boundaries:

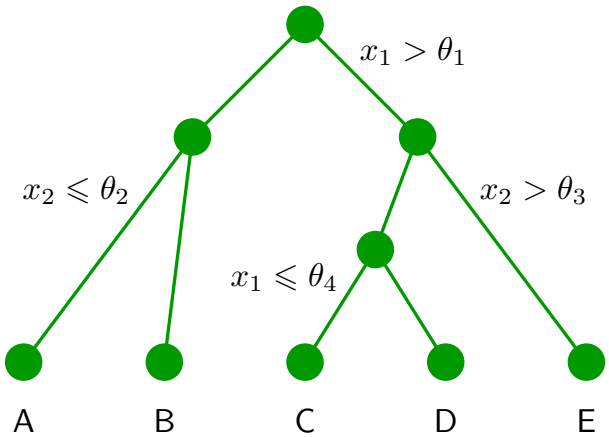




Parameters

Parameters to learn for a decision tree:

- the structure of the tree, such as the depth, #branches, #nodes, etc
  - some of them are sometimes considered as hyperparameters
  - unlike typical neural nets, the structure of a tree is *not fixed in advance, but learned from data*
- the test at each internal node
  - which feature(s) to test on?
  - if the feature is continuous, what threshold $(\theta_1, \theta_2, \ldots)$?



- the value/prediction of the leaves (A, B, ...)

Learning the parameters

So how do we *learn all these parameters?*

Recall typical approach is to find the parameters that minimize some loss.

This is unfortunately *not feasible for trees*

- For Z nodes, there are roughly $\#\text{features}^Z$ different ways to decide "which feature to test on each node", which is *a lot*.
- enumerating all these configurations to find the one that minimizes some loss is too computationally expensive.

Instead, we turn to some **greedy top-down approach**.

A running example

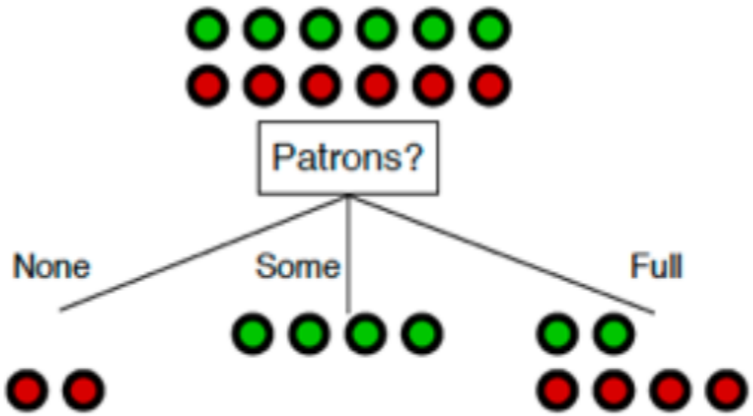
[Russell & Norvig, AIMA]

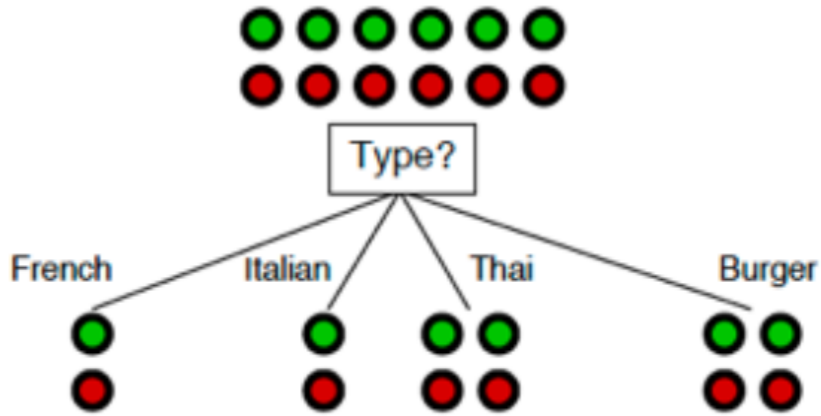
- predict whether a customer will wait for a table at a restaurant
- 12 training examples
- 10 features (all discrete)

Example	Attributes										Target
	Alt	*Bar*	*Fri*	*Hun*	*Pat*	*Price*	*Rain*	*Res*	*Type*	*Est*	*WillWait*
X_1	T	F	F	T	Some	$$$	F	T	French	0–10	T
X_2	T	F	F	T	Full	$	F	F	Thai	30–60	F
X_3	F	T	F	F	Some	$	F	F	Burger	0–10	T
X_4	T	F	T	T	Full	$	F	F	Thai	10–30	T
X_5	T	F	T	F	Full	$$$	F	T	French	>60	F
X_6	F	T	F	T	Some	$$	T	T	Italian	0–10	T
X_7	F	T	F	F	None	$	T	F	Burger	0–10	F
X_8	F	F	F	T	Some	$$	T	T	Thai	0–10	T
X_9	F	T	T	F	Full	$	T	F	Burger	>60	F
X_{10}	T	T	T	T	Full	$$$	F	T	Italian	10–30	F
X_{11}	F	F	F	F	None	$	F	F	Thai	0–10	F
X_{12}	T	T	T	T	Full	$	F	F	Burger	30–60	T

First step: how to build the root?

I.e., which feature should we test at the root? Examples:





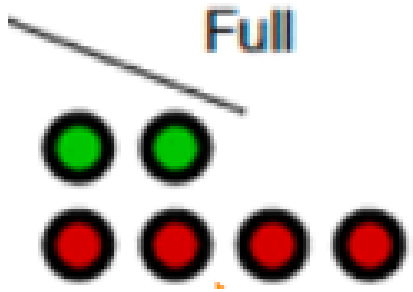
Which split is better?

- intuitively "patrons" is a better feature since it leads to "more pure" or "more certain" children
- how to quantify this intuition?

Measure of uncertainty of a node

It should be **a function of the distribution of classes**

- e.g. a node with 2 positive and 4 negative examples can be summarized by a distribution P with $P(Y = +1) = 1/3$ and $P(Y = -1) = 2/3$



One classic uncertainty measure of a distribution is its *(Shannon) entropy*:

$$H(P) = -\sum_{k=1}^{C} P(Y = k) \log P(Y = k)$$

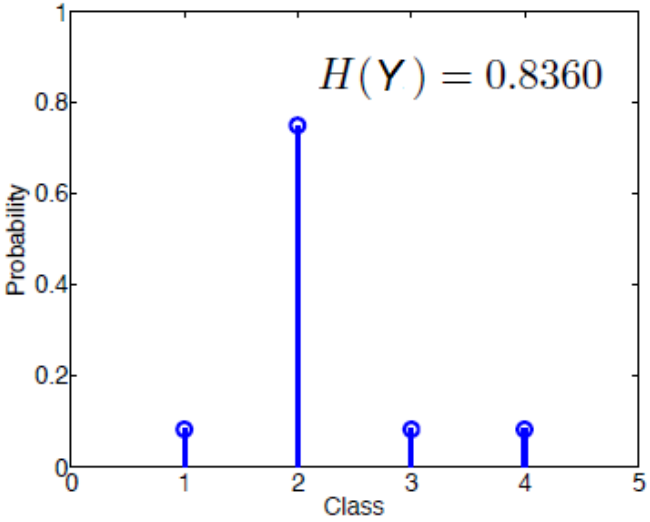
Properties of entropy

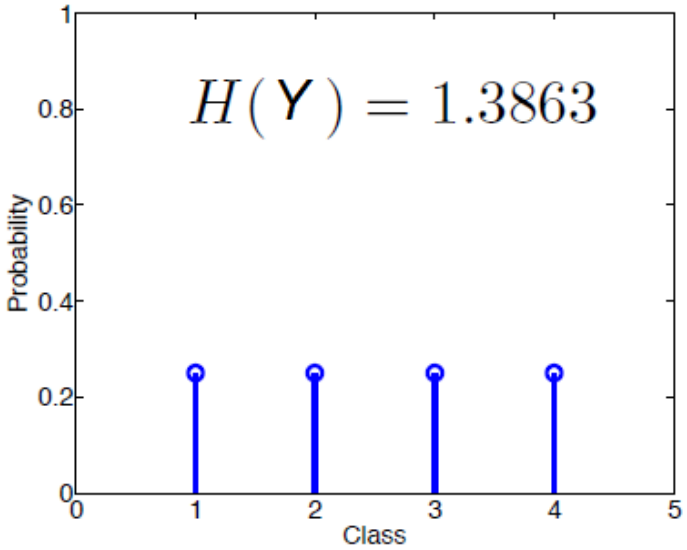
$$H(P) = -\sum_{k=1}^{\mathsf{C}} P(Y = k) \log P(Y = k)$$

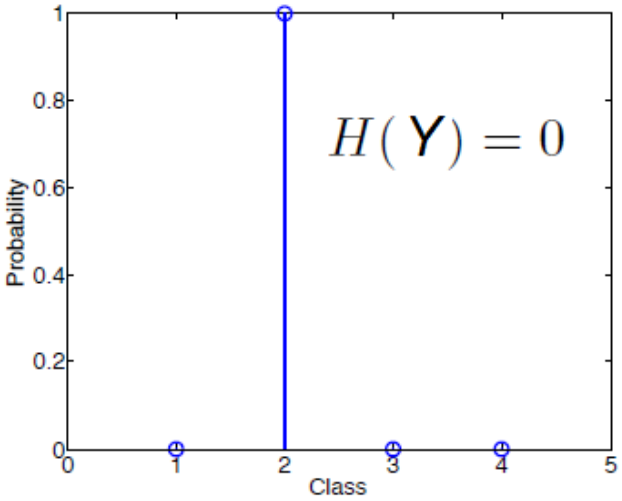
- the base of log can be 2, e or 10
- always non-negative
- it's the *smallest codeword length to encode symbols drawn from* P
- maximized if P is uniform (max $= \log \mathsf{C}$): most uncertain case
- minimized if P focuses on one class (min $= 0$): most certain case
  - e.g. $P = (1, 0, \ldots, 0)$
  - $0 \log 0$ is defined naturally as $\lim_{z \to 0+} z \log z = 0$

Examples of computing entropy

With base e and 4 classes:







Another example

Entropy in each child if root tests on "patrons"

For "None" branch

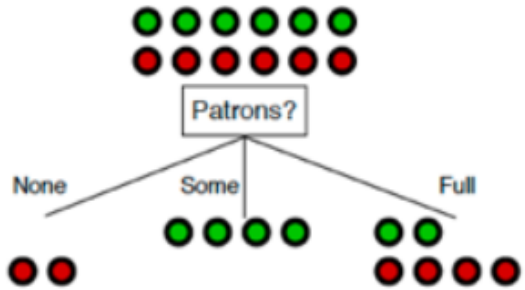
$$-\left(\frac{0}{0+2}\log\frac{0}{0+2}+\frac{2}{0+2}\log\frac{2}{0+2}\right)=0$$

For "Some" branch

$$-\left(\frac{4}{4+0}\log\frac{4}{4+0}+\frac{0}{4+0}\log\frac{0}{4+0}\right)=0$$

For "Full" branch

$$-\left(\frac{2}{2+4}\log\frac{2}{2+4}+\frac{4}{2+4}\log\frac{4}{2+4}\right)\approx 0.9$$



So how good is choosing "patrons" overall?

Very naturally, we take the weighted average of entropy:

$$\frac{2}{12}\times 0+\frac{4}{12}\times 0+\frac{6}{12}\times 0.9=0.45$$

Measure of uncertainty of a split

Suppose we split based on a discrete feature A, the uncertainty can be measured by the **conditional entropy**:

$$
\begin{aligned}
& H(Y \mid A) \\
&= \sum_a P(A=a) H(Y \mid A=a) \\
&= \sum_a P(A=a) \left(-\sum_{k=1}^{\mathsf{C}} P(Y \mid A=a) \log P(Y \mid A=a) \right) \\
&= \sum_a \text{“fraction of example at node } A=a\text{”} \times \text{“entropy at node } A=a\text{”}
\end{aligned}
$$

Pick the feature that leads to the smallest conditional entropy.

Deciding the root

For "French" branch

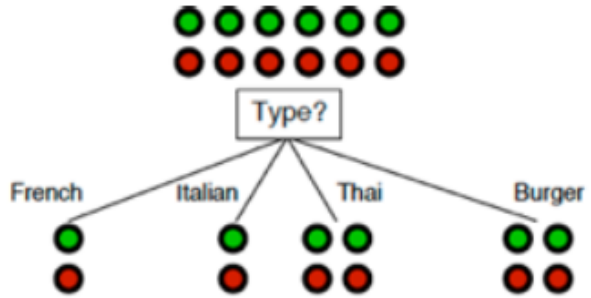
$$-\left(\frac{1}{1+1}\log\frac{1}{1+1}+\frac{1}{1+1}\log\frac{1}{1+1}\right)=1$$

For "Italian" branch

$$-\left(\frac{1}{1+1}\log\frac{1}{1+1}+\frac{1}{1+1}\log\frac{1}{1+1}\right)=1$$

For "Thai" and "Burger" branches

$$-\left(\frac{2}{2+2}\log\frac{2}{2+2}+\frac{2}{2+2}\log\frac{2}{2+2}\right)=1$$



The conditional entropy is $\frac{2}{12}\times 1+\frac{2}{12}\times 1+\frac{4}{12}\times 1+\frac{4}{12}\times 1=1>0.45$

So splitting with "patrons" is better than splitting with "type".

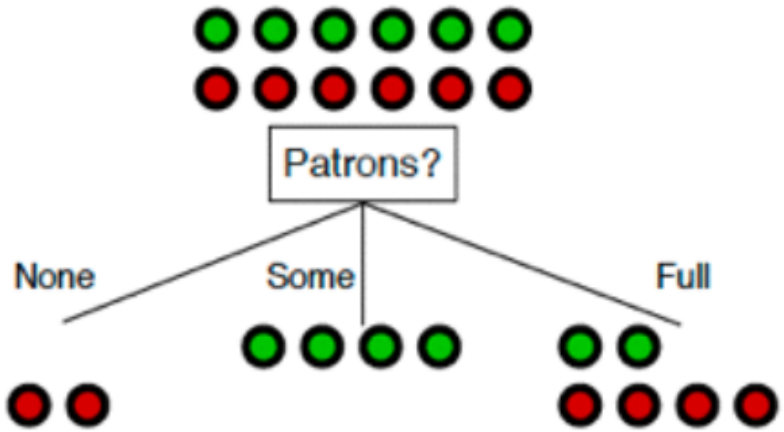
In fact by similar calculation "patrons" is the best split among all features.

We are now done with building the root (this is also called a **stump**).

Repeat recursively

Split each child in the same way.

- but no need to split children "none" and "some": they are pure already and become leaves
- for "full", repeat, focusing on those 6 examples:



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X_8	F	F	F	T	Some	$$	T	T	Thai	0–10	T
X_9	F	T	T	F	Full	$	T	F	Burger	>60	F
X_{10}	T	T	T	T	Full	$$$	F	T	Italian	10–30	F
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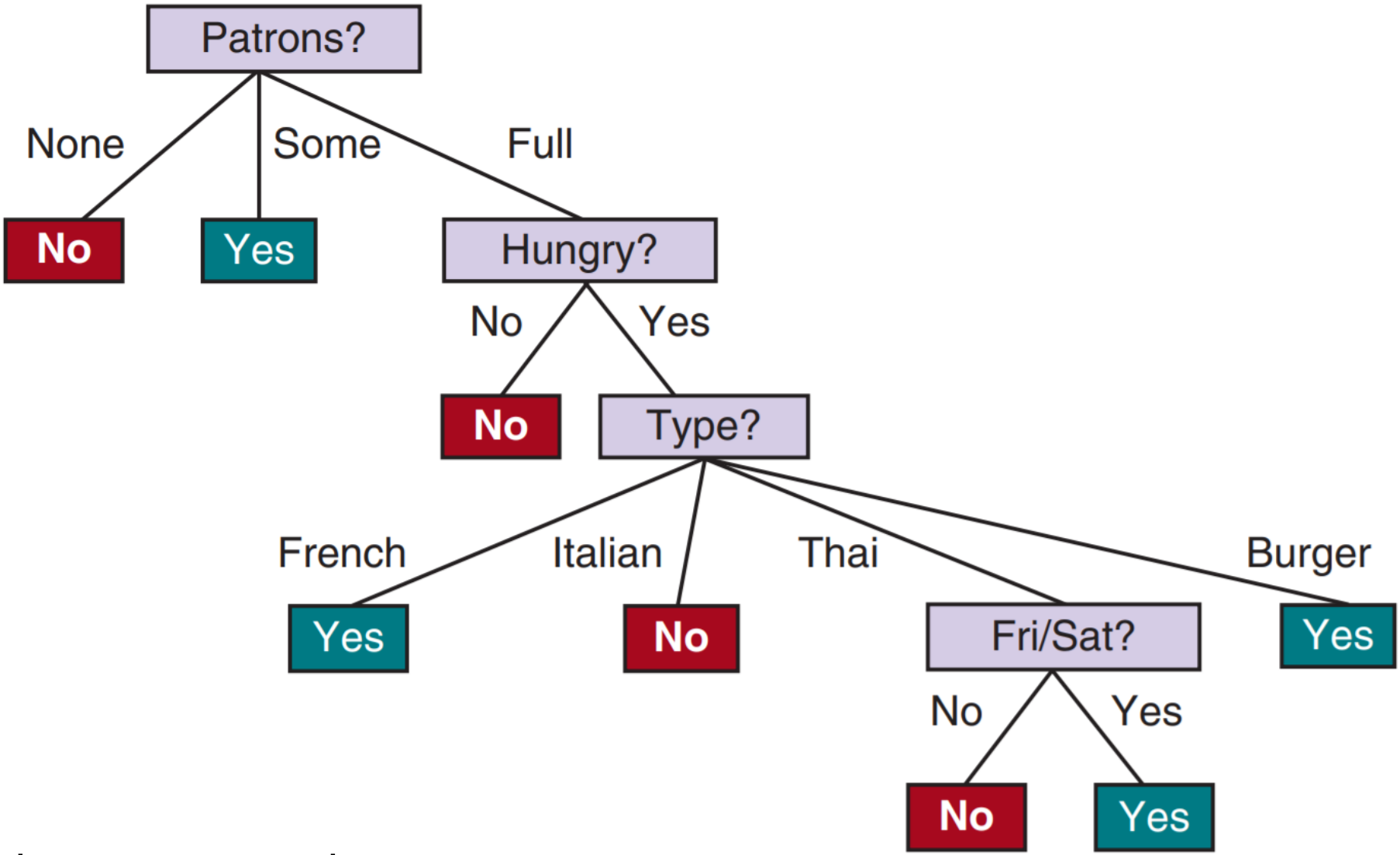


Again, very easy to interpret.

Putting it together

DecisionTreeLearning(Examples, Features)

- if Examples have the same class, return a leaf with this class
- else if Features is empty, return a leaf with the majority class
- else if Examples is empty, return a leaf with majority class of parent
- else

  find the best feature A to split (e.g. based on conditional entropy)

  Tree $\leftarrow$ a root with test on A

  For each value a of A:

  Child $\leftarrow$ **DecisionTreeLearning**(Examples with $A = a$, Features$\backslash\{A\}$)
  add **Child** to **Tree** as a new branch
- return **Tree**

Variants

Popular decision tree algorithms (e.g. C4.5, CART, etc) are all based on this framework.

Variants:

- replace entropy by **Gini impurity**:

$$G(P) = \sum_{k=1}^{C} P(Y = k)(1 - P(Y = k))$$

  meaning: *probability of two randomly drawn classes being different*

- if a feature is continuous, we need to find a threshold that leads to minimum conditional entropy or Gini impurity. *Think about how to do it efficiently.*

Regularization

If the dataset has no contradiction (i.e. same $\boldsymbol{x}$ but different y), the training error of a tree is always zero, which might indicate overfitting.

Pruning is a typical way to prevent overfitting for a tree:

- restrict the depth or #nodes
- other more principled approaches
- all make use of a validation set

Random Forests

Random forest is an **ensemble** of trees:

- each tree is built using *a bootstrapped dataset* (that is, a set of points randomly sampled from the training set with replacement)
- each split of each tree is selected from *a random subset of features*
- final prediction is the *majority vote* of all tress (for classification tasks) or the *averaged prediction* of all trees (for regression tasks)
- much better performance than a single tree, trivially parallelizable!

Outline

Introduction

Boosting (an even more powerful/general ensemble method):

- is a meta-algorithm, which takes a base algorithm (classification, regression, ranking, etc) as input and boosts its accuracy
- main idea: combine weak "rules of thumb" (e.g. 51% accuracy) to form a highly accurate predictor (e.g. 99% accuracy)
- works very well in practice (especially in combination with trees)
- often is *resistant to overfitting*
- has strong theoretical guarantees

We again focus on binary classification.

A simple example

Email spam detection:

- given a training set like:
  - ("Want to make money fast? ...", **spam**)
  - ("Viterbi Research Gist ...", **not spam**)
- first obtain a classifier by applying a base algorithm, which can be a rather simple/weak one, like decision stumps:
  - e.g. contains the word "money" $\Rightarrow$ spam
- reweight the examples so that "difficult" ones get more attention
  - e.g. spam that doesn't contain the word "money"
- obtain another classifier by applying the same base algorithm:
  - e.g. empty "to address" $\Rightarrow$ spam
- repeat ...
- final classifier is the (weighted) majority vote of all weak classifiers

The base algorithm

A **base algorithm** $\mathcal{A}$ (also called weak learning algorithm/oracle) takes a training set S weighted by D as input, and outputs classifier $h \leftarrow \mathcal{A}(S, D)$

- this can be any off-the-shelf classification algorithm (e.g. decision trees, logistic regression, neural nets, etc)
- many algorithms can deal with a weighted training set (e.g. for algorithm that minimizes some loss, we can simply replace "total loss" by "weighted total loss")
- even if it's not obvious how to deal with weight directly, we can always resample according to D to create a new unweighted dataset

Boosting Algorithms

Given:

- a training set S
- a base algorithm $\mathcal{A}$

Two things to specify a boosting algorithm:

- how to **reweight** the examples?
- how to **combine** all the weak classifiers?

Focus on AdaBoost, one of the most successful boosting algorithms.

The AdaBoost Algorithm

Given a training set S and a base algorithm $\mathcal{A}$, initialize D_1 to be uniform

For $t = 1, \ldots, T$

- obtain a weak classifier $h_t \leftarrow \mathcal{A}(S, D_t)$
- calculate the importance of h_t as

$$\beta_t = \frac{1}{2} \ln \left(\frac{1 - \epsilon_t}{\epsilon_t} \right) \qquad (\beta_t > 0 \Leftrightarrow \epsilon_t < 0.5)$$

  where $\epsilon_t = \sum_{n: h_t(\boldsymbol{x}_n) \neq y_n} D_t(n)$ is the weighted error of h_t.

- update distributions

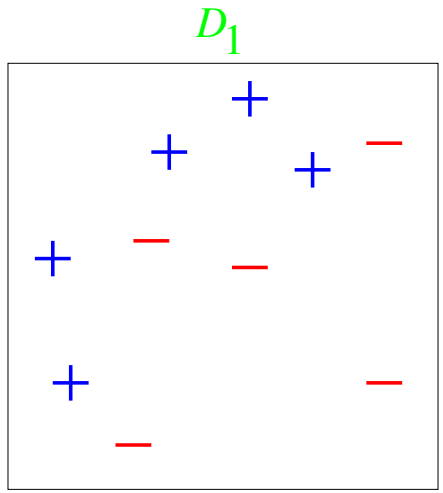
$$D_{t+1}(n) \propto D_t(n) e^{-\beta_t y_n h_t(\boldsymbol{x}_n)} = \begin{cases} D_t(n) e^{-\beta_t} & \text{if } h_t(x_n) = y_n \\ D_t(n) e^{\beta_t} & \text{else} \end{cases}$$

Output the final classifier $H(\boldsymbol{x}) = \text{sgn} \left(\sum_{t=1}^{T} \beta_t h_t(\boldsymbol{x}) \right)$

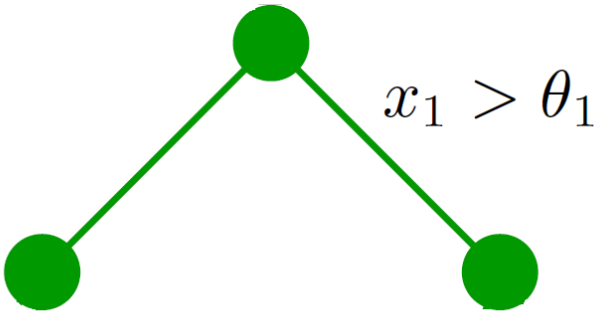
Example

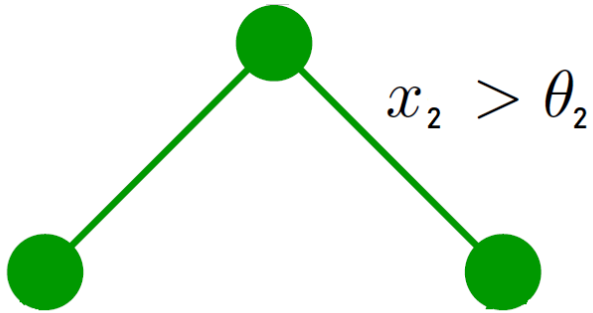
10 data points in $\mathbb{R}^2$

The size of $+$ or - indicates the weight, which starts from uniform D_1



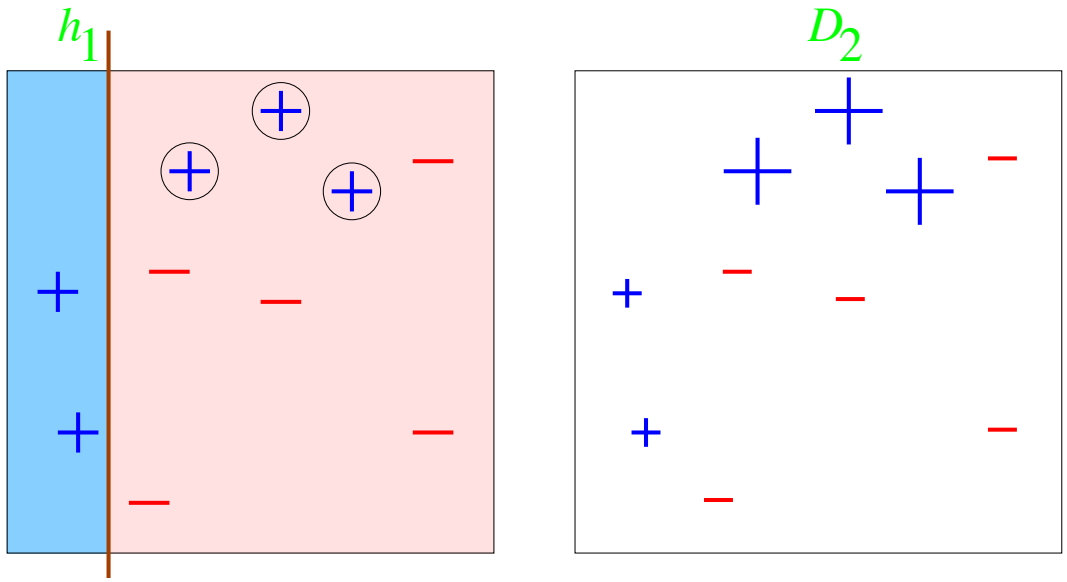
Base algorithm is decision stump:





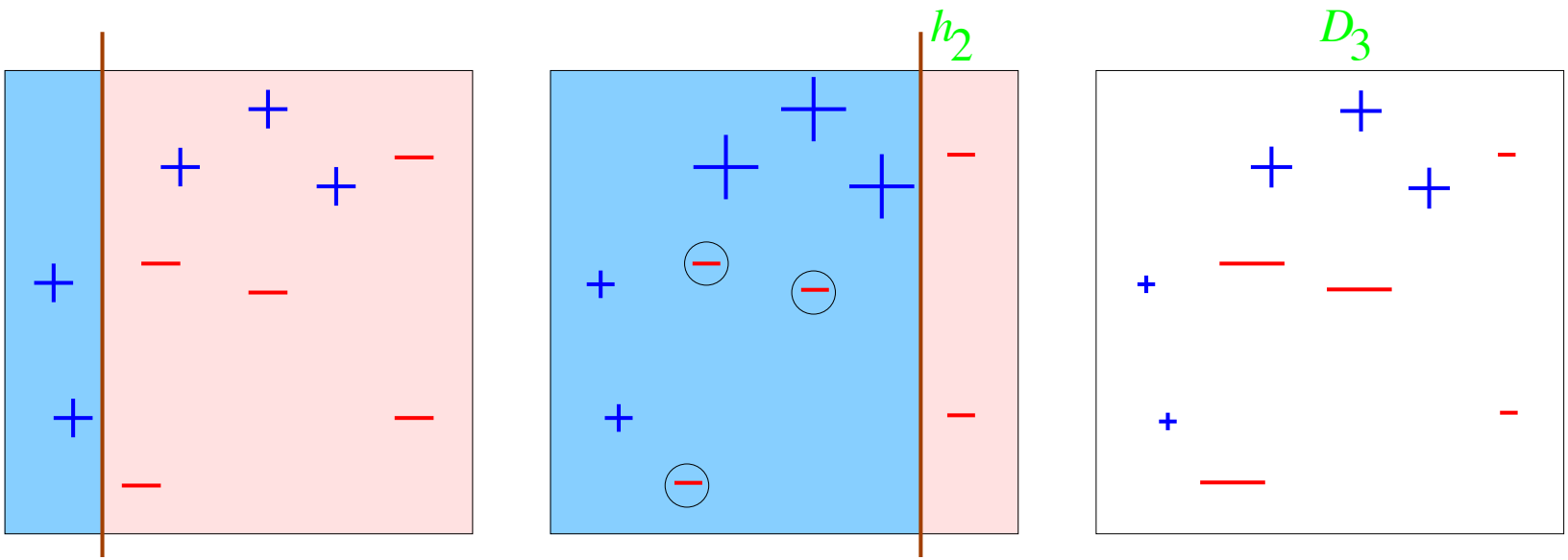
Observe that *no stump can predict very accurately for this dataset*

Round 1: $t = 1$



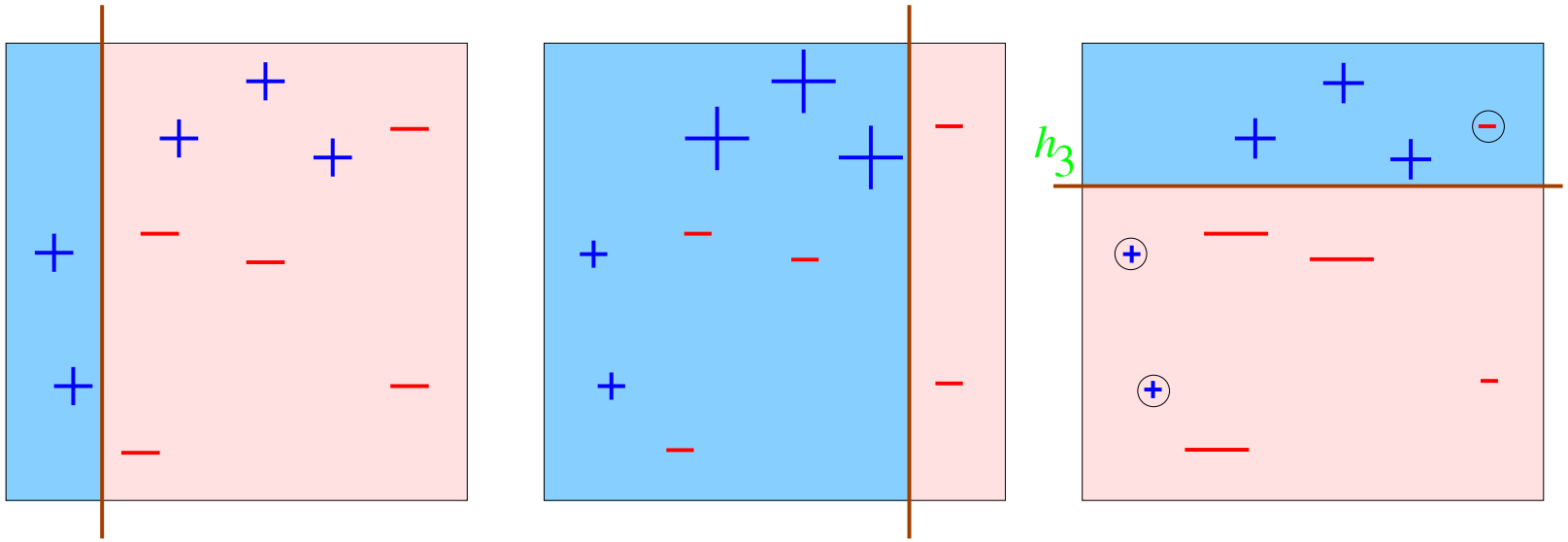
- 3 misclassified (circled): $\epsilon_1 = 0.3 \rightarrow \beta_1 = \frac{1}{2} \ln \left(\frac{1-\epsilon_t}{\epsilon_t} \right) \approx 0.42.$
- D_2 puts more weights on those examples

Round 2: $t = 2$



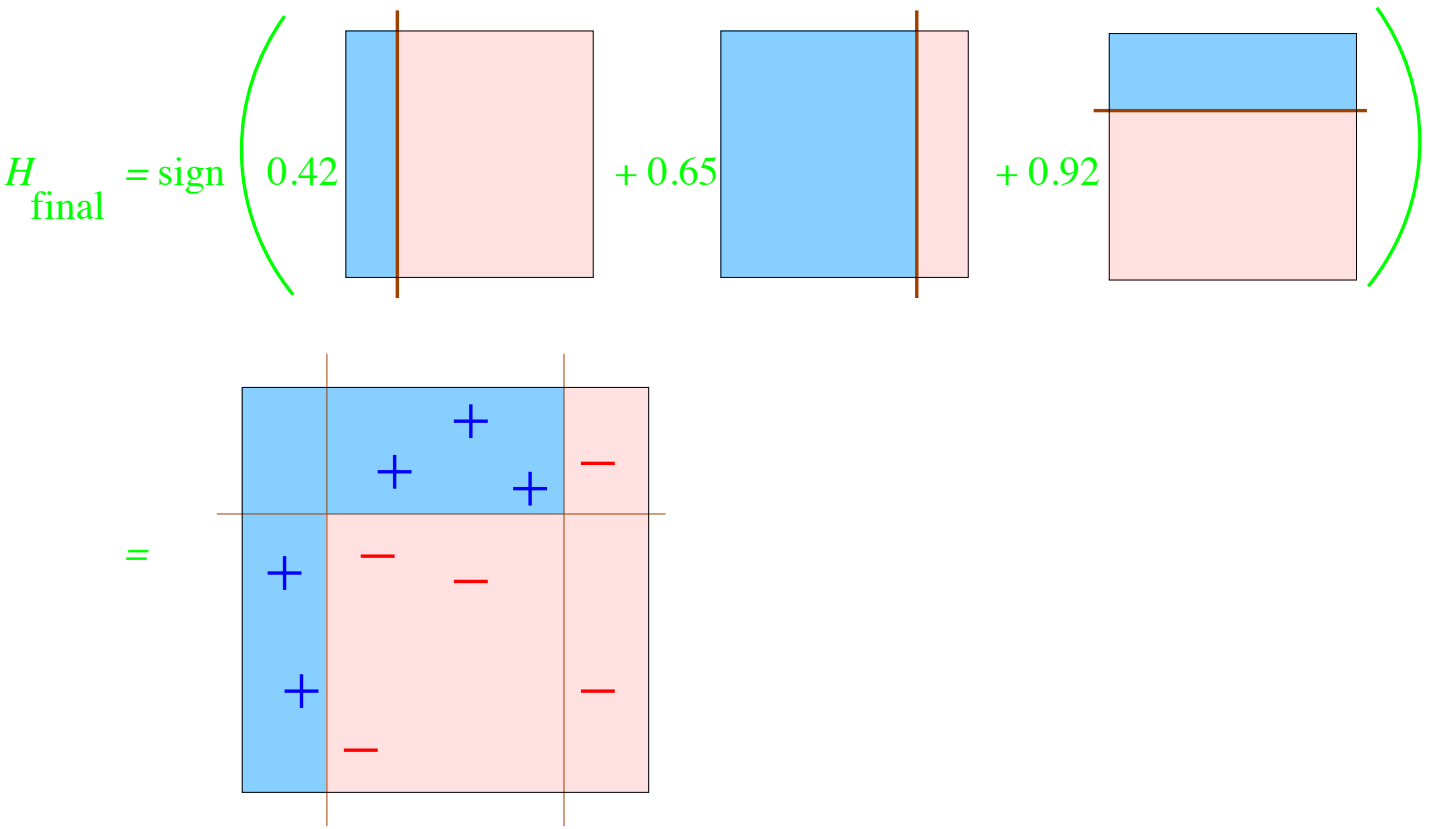
- 3 misclassified (circled): $\epsilon_2 = 0.21 \rightarrow \beta_2 = 0.65$.
- D_3 puts more weights on those examples

Round 3: $t = 3$



- again 3 misclassified (circled): $\epsilon_3 = 0.14 \rightarrow \beta_3 = 0.92$.

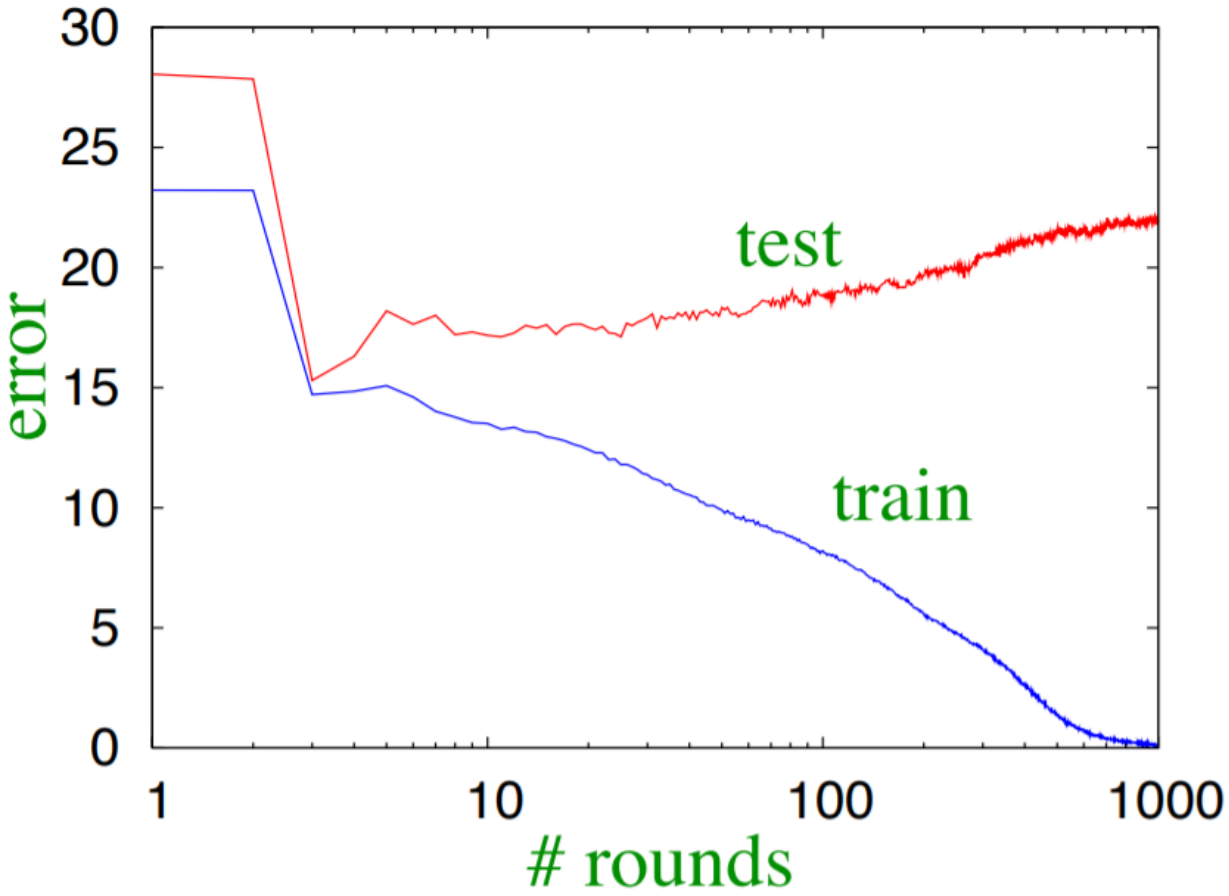
Final classifier: combining 3 classifiers



All data points are now classified correctly, even though each weak classifier makes 3 mistakes.

Overfitting

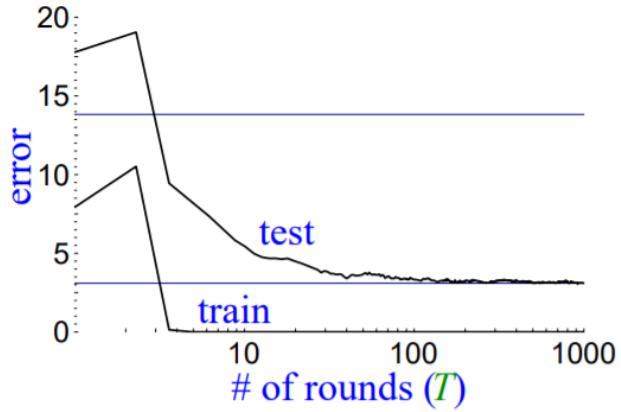
When T is large, the model is very complicated and overfitting can happen



(boosting "stumps" on heart-disease dataset)

Resistance to overfitting

However, *very often AdaBoost is resistant to overfitting*



(boosting C4.5 on "letter" dataset)

- test error does not increase, even after 1000 rounds
  - (total size $>$ 2,000,000 nodes)
- test error continues to drop even after training error is zero!

	# rounds		
	5	100	1000
train error	0.0	0.0	0.0
test error	8.4	3.3	3.1

Used to be a mystery, but by now rigorous theory has been developed to explain this phenomenon.

Why AdaBoost works?

In fact, *AdaBoost also follows the general framework of minimizing some surrogate loss*.

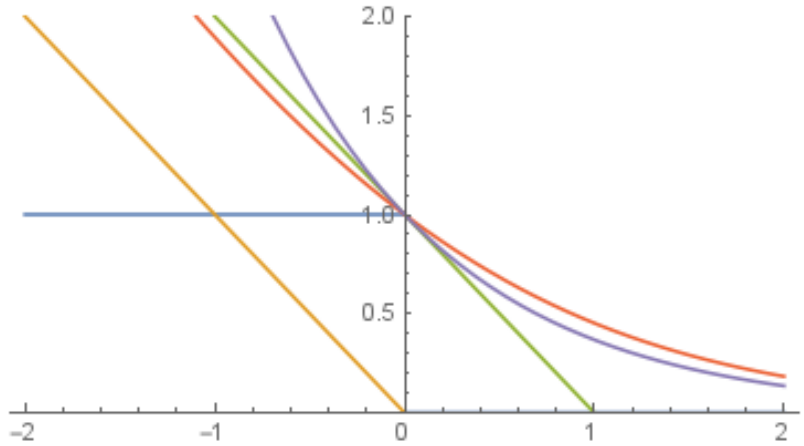
Step 1: the model that AdaBoost considers is

$$\left\{ \text{sgn}\left(f(\cdot)\right) \;\middle|\; f(\cdot) = \sum_{t=1}^{T} \beta_t h_t(\cdot) \text{ for some } \beta_t \geq 0 \text{ and } h_t \in \mathcal{H} \right\}$$

where $\mathcal{H}$ is the set of models considered by the base algorithm

Step 2: the loss that AdaBoost minimizes is the exponential loss

$$\sum_{n=1}^{\mathrm{N}} \exp\left(-y_n f(\boldsymbol{x}_n)\right)$$



Greedy minimization

Step 3: the way that AdaBoost minimizes exponential loss is by a greedy approach, that is, find β_t, h_t one by one for $t = 1, \ldots, T$.

Specifically, let $f_t = \sum_{\tau=1}^{t} \beta_\tau h_\tau$. Suppose we have found f_{t-1}, *what should f_t be?* Greedily, we want to find β_t, h_t to minimize

$$\begin{aligned}\sum_{n=1}^{\mathsf{N}} \exp\left(-y_n f_t(\boldsymbol{x}_n)\right) &= \sum_{n=1}^{\mathsf{N}} \exp\left(-y_n f_{t-1}(\boldsymbol{x}_n)\right) \exp\left(-y_n \beta_t h_t(\boldsymbol{x}_n)\right) \\ &\propto \sum_{n=1}^{\mathsf{N}} D_t(n) \exp\left(-y_n \beta_t h_t(\boldsymbol{x}_n)\right)\end{aligned}$$

where the last step is by the definition of weights

$$D_t(n) \propto D_{t-1}(n) \exp\left(-y_n \beta_{t-1} h_{t-1}(\boldsymbol{x}_n)\right) \propto \cdots \propto \exp\left(-y_n f_{t-1}(\boldsymbol{x}_n)\right)$$

Greedy minimization

So the goal becomes finding $\beta_t, h_t \in \mathcal{H}$ that minimize

$$\begin{aligned}
&\sum_{n=1}^{\mathsf{N}} D_t(n) \exp\left(-y_n \beta_t h_t(\boldsymbol{x}_n)\right) \\
&= \sum_{n:y_n \neq h_t(\boldsymbol{x}_n)} D_t(n) e^{\beta_t} + \sum_{n:y_n = h_t(\boldsymbol{x}_n)} D_t(n) e^{-\beta_t} \\
&= \epsilon_t e^{\beta_t} + (1-\epsilon_t) e^{-\beta_t} \qquad \text{(recall } \epsilon_t = \textstyle\sum_{n:y_n \neq h_t(\boldsymbol{x}_n)} D_t(n)\text{)} \\
&= \epsilon_t (e^{\beta_t} - e^{-\beta_t}) + e^{-\beta_t}
\end{aligned}$$

It is now clear we should find h_t to minimize its the weighted classification error ϵ_t, *exactly what the base algorithm should do intuitively!*

This greedy step is abstracted out through a base algorithm.

Greedy minimization

When h_t (and thus ϵ_t) is fixed, we then find β_t to minimize

$$\epsilon_t(e^{\beta_t} - e^{-\beta_t}) + e^{-\beta_t},$$

which gives (by setting the derivative to 0):

$$\beta_t = \frac{1}{2} \ln \left(\frac{1 - \epsilon_t}{\epsilon_t} \right)$$

Keep doing this greedy minimization gives the AdaBoost algorithm.

Summary for boosting

Key idea of boosting is to **combine weak predictors into a strong one**.

There are many boosting algorithms: AdaBoost is the most classic one (XGBoost is the most popular in practice)

AdaBoost is **greedily minimizing the exponential loss**.

AdaBoost is often **resistant to overfitting**.