CSE 417T Introduction to Machine Learning

Lecture 4

Instructor: Chien-Ju (CJ) Ho

Logistics: HW1

- Due: Feb 19 (Friday), 2020
 - http://chienjuho.com/courses/cse417t/hw1.pdf
 - Strongly encouraged to work on it before the drop deadline
 - Two submission links: Report and Code
 - Report: Answer all questions, including the implementation question
 - Grades are based on the report
 - Code: Complete and submit hw1.py for Problem 2
 - The code will only be used for correctness checking (when in doubts) and plagiarism checking
 - Reserve time if you never used Gradescope.
 - Make sure to specify the pages for each problem. You won't get points otherwise.
- (Optional) Python session
 - See Piazza post

Logistics: Office Hours

Tentative schedule of TA office hours (starting next week)

Monday	10:00 AM to 11:20 AM (Oliver)	02:30 PM to 03:50 PM (Guanghui)	
Tuesday	02:30 PM to 03:50 PM (Quentin)	05:00 PM to 06:20 PM (Victoria)	
Wednesday	04:30 PM to 05:50 PM (Amrit)	08:00 PM to 09:20 PM (Cecilia)	
Thursday	10:00 AM to 11:20 AM (Aaron)	02:30 PM to 03:50 PM (Matthew)	
Friday	08:00 AM to 09:20 AM (Shohaib)	12:00 PM-1:20 PM (Tong)	04:10 PM to 05:30 PM (Flora)

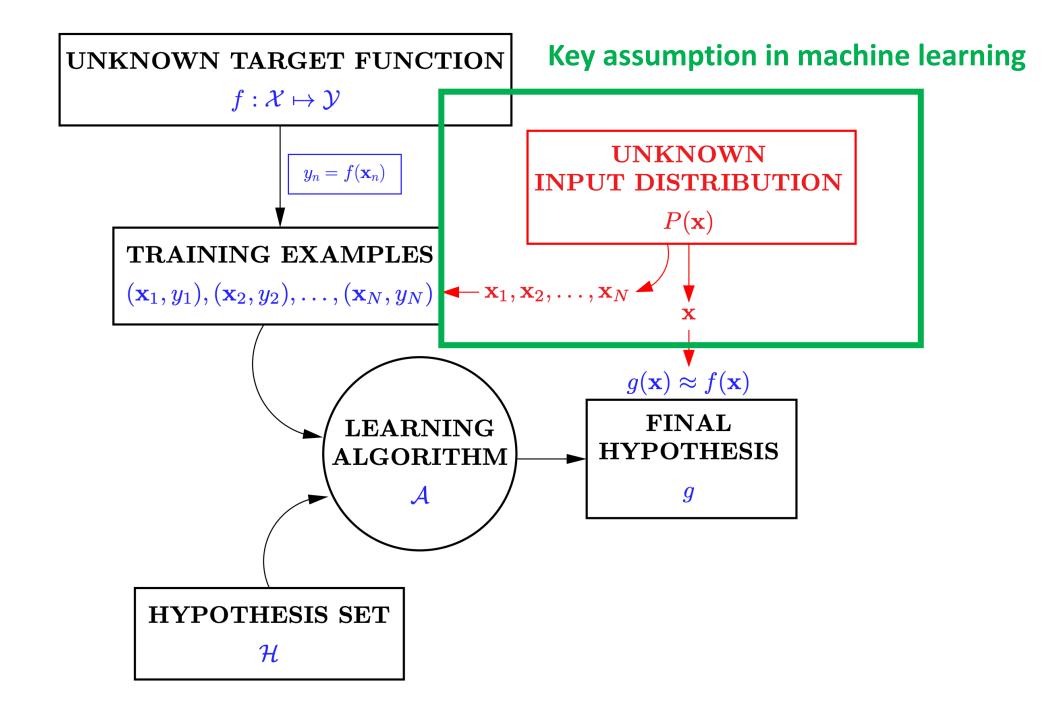
- Remote via Zoom
- Please follow Piazza for zoom links and potential updates
- Recommendation: Try to utilize the office hour early (way ahead of deadlines), you are likely to get more of TAs' time this way.

Recap

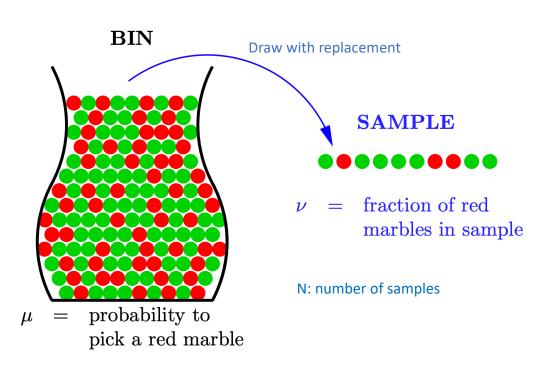
Common Notations

- Data point with augmented x_0 : $\vec{x} = (x_0, ..., x_d)$
 - We often use d to specify the dimensions of data points
 - We augment $x_0 = 1$ for each data point (Check Lecture 1 for the reasoning)
- Dataset: $D = \{(\vec{x}_1, y_1), ..., (\vec{x}_N, y_N)\}$
 - We often use N to specify the number of data points in the dataset
- Hypothesis set H
 - We use $h \in H$ to specify an arbitrary hypothesis
 - We use $g \in H$ to specify the hypothesis output by the learning algorithm
- Indicator variable:
 - $\mathbb{I}[\text{event}] = \begin{cases} 1 & \text{if event is true} \\ 0 & \text{if event is false} \end{cases}$

Example:
$$\mathbb{I}[h(\vec{x}) \neq f(\vec{x})] = \begin{cases} 1 & \text{if } h(\vec{x}) \neq f(\vec{x}) \\ 0 & \text{if } h(\vec{x}) = f(\vec{x}) \end{cases}$$



Hoeffding's Inequality



$$\Pr[|\mu - \nu| > \epsilon] \le 2e^{-2\epsilon^2 N}$$

Define $\delta = \Pr[|\mu - \nu| > \epsilon]$

- Fix δ , ϵ decreases as N increases
- Fix ϵ , δ decreases as N increases
- Fix N, δ decreases as ϵ increases

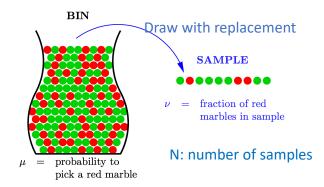
Informal intuitions of notations

N: # sample

 δ : probability of "bad" event

 ϵ : error of estimation

Connection to Learning



- Given dataset $D = \{(\vec{x}_1, y_1), ..., (\vec{x}_N, y_N)\}.$
- Fix a hypothesis h
 - $E_{in}(h) \stackrel{\text{def}}{=} \frac{1}{N} \sum_{n=1}^{N} \mathbb{I}[h(\vec{x}_n) \neq f(\vec{x}_n)]$ [In-sample error, analogy to ν]
 - $E_{out}(h) \stackrel{\text{def}}{=} \Pr_{\vec{x} \sim P(\vec{x})}[h(\vec{x}) \neq f(\vec{x})]$ [Out-of-sample error, analogy to μ]
- Apply Hoeffding's inequality

$$Pr[|E_{out}(h) - E_{in}(h)| > \epsilon] \leq 2e^{-2\epsilon^2 N}$$

• This is verification, not learning

Connection to "Real" Learning

- Given a finite hypothesis set $H = \{h_1, ..., h_M\}$
- Apply some learning algorithm on D, output a $g \in H$
- What can we say about $E_{out}(g)$ from $E_{in}(g)$?

$$Pr[|E_{out}(g) - E_{in}(g)| > \epsilon] \le 2Me^{-2\epsilon^2N}$$
 for any $\epsilon > 0$

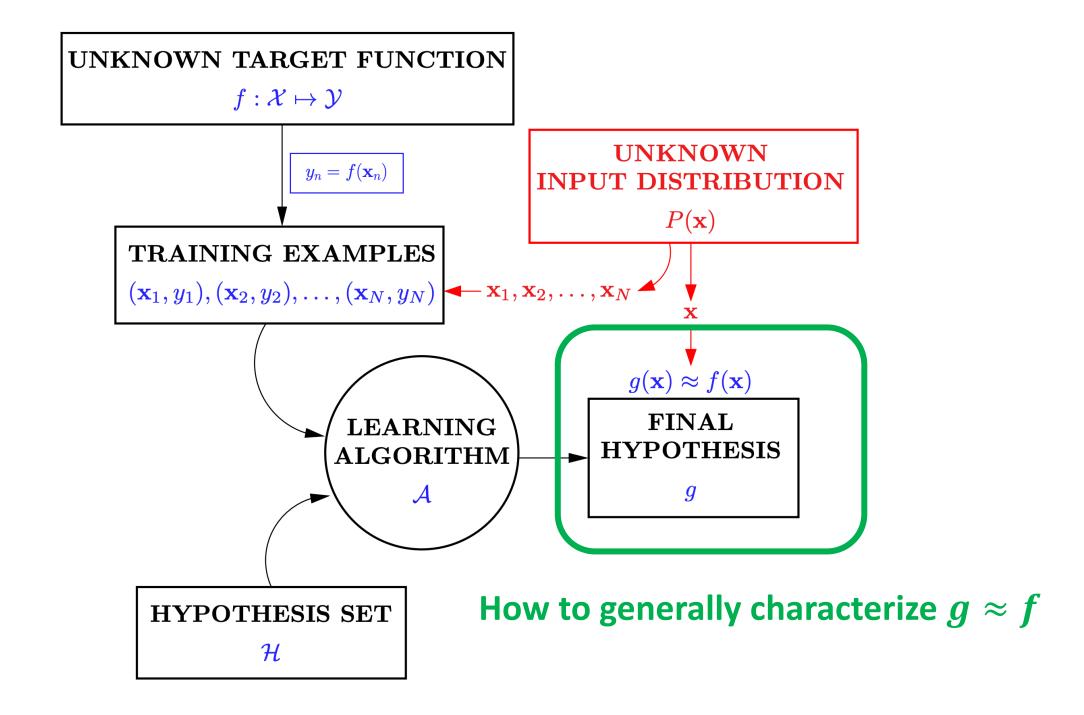
Intuitions:

- 1. Bad event $B(g) \subseteq B(h_1) \cup B(h_2) \dots \cup B(h_M)$ g is selected within $\{h_1, \dots, h_M\}$ => bad event of g is within the union of the bad events of h_1, \dots, h_M
- 2. $\Pr[B(g)] \leq \Pr[B(h_1)] + \dots + \Pr[B(h_M)]$ each of the $\Pr[B(h_m)]$ follows Hoeffding's inequality

Today's Lecture

The notes are not intended to be comprehensive. They should be accompanied by lectures and/or textbook. Let me know if you spot errors.

Revisit the learning problem



Goal: $g \approx f$

- A general approach:
 - Define an error function E(h, f) that quantify how far away h is to f
 - choose $g = \underset{h \in \mathcal{H}}{\operatorname{argmin}} E(h, f)$
- A major component of ML is optimization
- E is usually defined in terms of a pointwise error function $e(h(\vec{x}), f(\vec{x}))$
 - Binary error (classification): $e(h(\vec{x}), f(\vec{x})) = \mathbb{I}[h(\vec{x}_n) \neq f(\vec{x}_n)]$
 - Squared error (regression): $e(h(\vec{x}), f(\vec{x})) = (f(\vec{x}) h(\vec{x}))^2$

$$E_{in}(h) = \frac{1}{N} \sum_{n=1}^{N} e(h(\vec{x}_n), f(\vec{x}_n))$$

$$E_{out}(h) = \mathbb{E}_{\vec{x}}[e(h(\vec{x}), f(\vec{x}))]$$

The discussion on the Hoeffding's inequality applies for general (bounded) error functions.

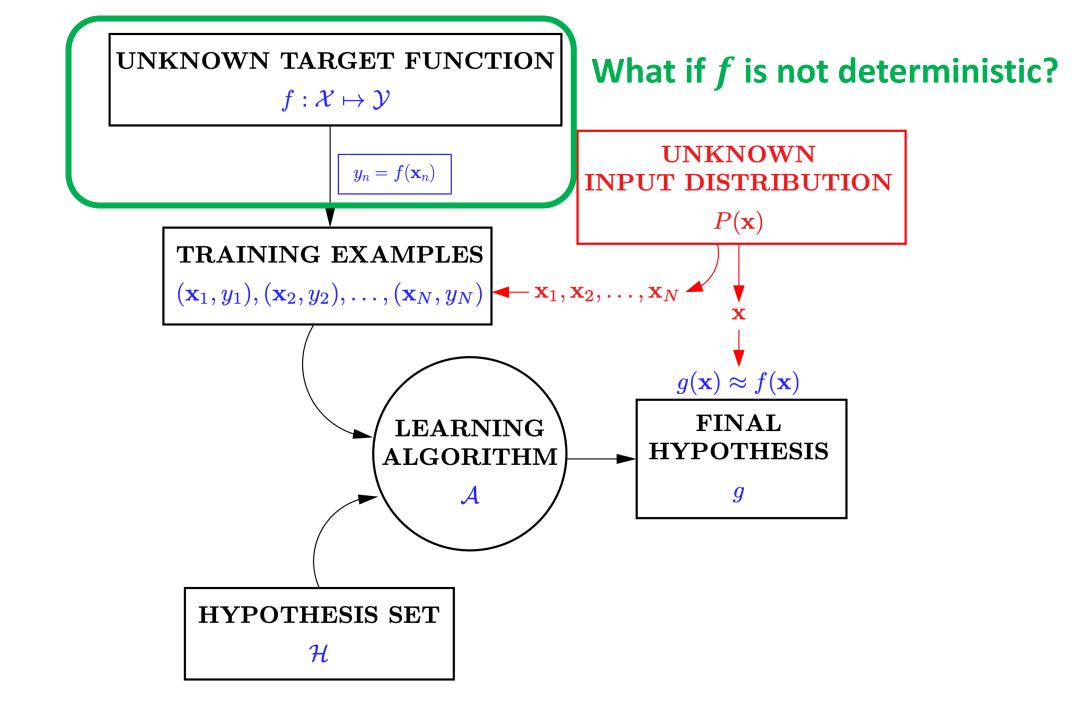
How to Choose Error Function

Depending on domain application

Cuparr	markat	$f(\vec{x})$	
Supermarket		+1	-1
h (♂)	+1	0	Small
$h(\vec{x})$	-1	Large	0

-	D I	$f(\vec{x})$	
FBI		+1	-1
$h(\vec{x})$	+1	0	Large
$h(\vec{x})$	-1	Small	0

- Make it "easier" to optimize
 - One natural ML algorithm: $g = \underset{h \in \mathcal{H}}{\operatorname{argmin}} E_{in}(h)$
 - (e.g., if E_{in} is convex, continuous, differentiable, we usually have efficient algorithms.)

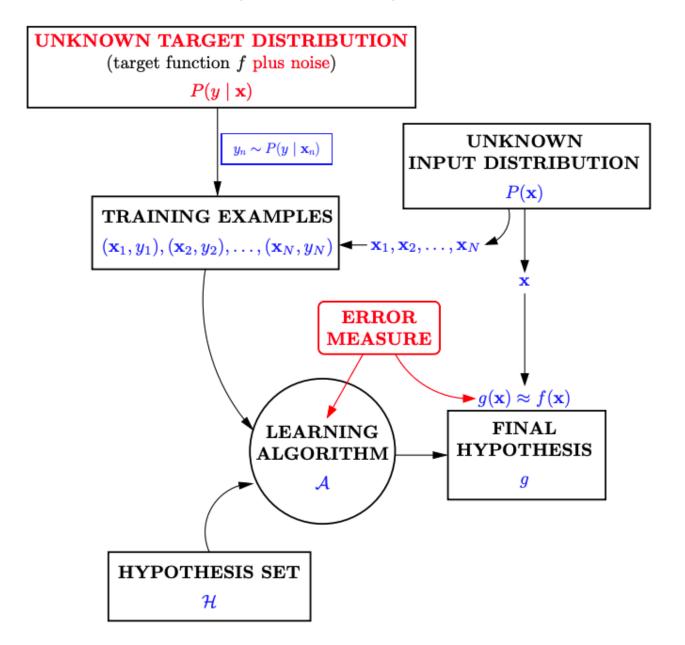


Noisy Target

- What if there doesn't exist f such that $y = f(\vec{x})$?
 - *f* is stochastic instead of deterministic
 - (even if two customers have exactly the same attributes, one might be a good customer for bank, and the other might not be)
- Common approach
 - Instead of a target function, define a target distribution
 - Instead of $y = f(\vec{x})$, y is drawn from a conditional distribution $P(y|\vec{x})$
 - $y = f(\vec{x}) + \epsilon$
 - $f(\vec{x})$ is the mean of the distribution
 - ϵ is zero-mean noise

The discussion on the Hoeffding's inequality applies for noisy targets.

General Setup of (Supervised) Learning



Theory of Generalization

Revisit the "Multi-Hypothesis" Bound

- Given a finite hypothesis set $H = \{h_1, ..., h_M\}$
- Apply some learning algorithm on D, output a $g \in H$
- What can we say about $E_{out}(g)$ from $E_{in}(g)$?

$$Pr[|E_{out}(g) - E_{in}(g)| > \epsilon] \le 2Me^{-2\epsilon^2N}$$
 for any $\epsilon > 0$

What if *M* is infinite?

 $Pr[|E_{out}(g) - E_{in}(g)| > \epsilon] \le 2Me^{-2\epsilon^2N}$ don't seem to carry any meanings

Key Intuitions in the Multi-Hypothesis Analysis

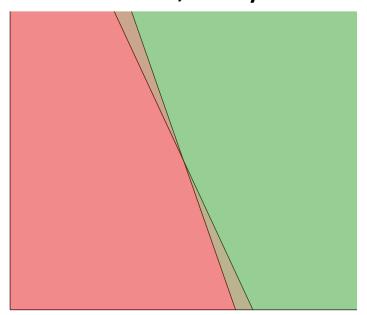
- Define "bad event of h" B(h) as $|E_{out}(h) E_{in}(h)| > \epsilon$
- If g is selected from $\{h_1, h_2\}$
 - $B(g) \subseteq B(h_1) \cup B(h_2)$
 - $\Pr[B(g)] \le \Pr[B(h_1) \text{ or } B(h_2)]$ $\le \Pr[B(h_1)] + \Pr[B(h_2)]$ (Union Bound)

 $B(h_1)$ $B(h_2)$

Union bound considers the worst case: Bad events don't overlap

Do Bad Events Overlap?

Oftentimes, they overlap a lot!



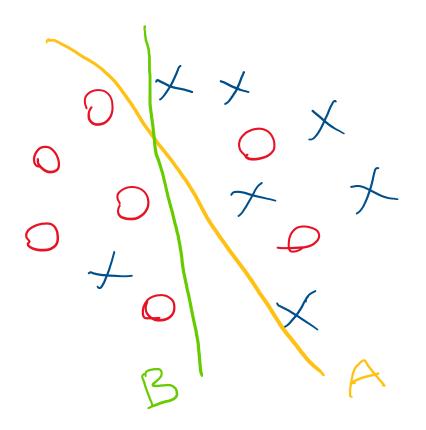
The two linear separators on the left make the same predictions for most points.

If it's a bad event for one, it's likely to be a bad event for the other.

"bad event of h" B(h): $|E_{out}(h) - E_{in}(h)| > \epsilon$

Recall: Informally, you can interpret "bad event of h" as the event that we draw a "unrepresentative dataset D" that makes the in-sample errors of h to be far away from out-of-sample error of h

What Can We Do?



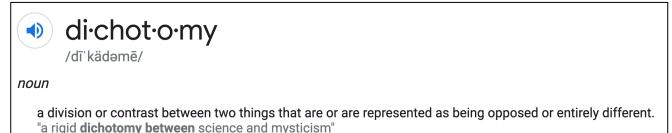
Any difference between A and B?

For this dataset, probably not.

They make the same predictions for every data point in this dataset.

What Can We Do?

• Let's define "data-dependent" hypothesis, call it dichotomy.



- A hypothesis $h: X \to \{-1, +1\}$
- A dichotomy for a set of data points $(\vec{x}_1, ..., \vec{x}_N)$:
 - Assign either +1 or -1 for each of the data points (divide the data points into two groups)
- Why dichotomies?
 - It helps us count "effective number of hypothesis" (to replace M)

More Formal Definitions

Dichotomies

- Informally, consider a dichotomy as "data-dependent" hypothesis
- Characterized by both hypothesis set H and N data points $(\vec{x}_1, ..., \vec{x}_N)$

$$H(\vec{x}_1, ... \vec{x}_N) = \{h(\vec{x}_1), ..., h(\vec{x}_N) | h \in H\}$$

• The set of possible prediction combinations $h \in H$ can induce on $\vec{x}_1, \dots, \vec{x}_N$

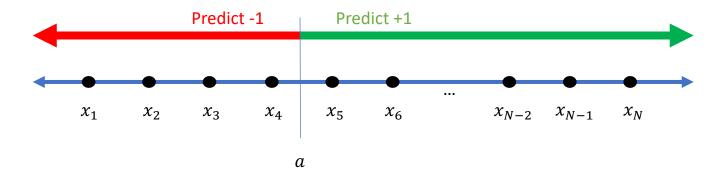
Growth function

• Largest number of dichotomies H can induce across all possible data sets of size N

$$m_H(N) = \max_{(\vec{x}_1, ..., \vec{x}_N)} |H(\vec{x}_1, ..., \vec{x}_N)|$$

Example: H = Positive Rays

- Data points are in one-dimensional space
- Positive rays: h(x) = sign(x a)



• What is $H(\vec{x}_1, ..., \vec{x}_N)$?

<u>Dichotomies</u>

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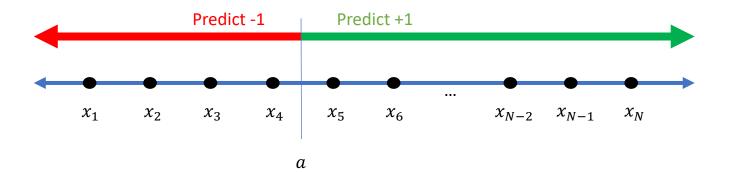
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• What is $m_H(N)$?

Example: H = Positive Rays

- Data points are in one-dimensional space
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• What is $H(\vec{x}_1, ..., \vec{x}_N)$?

$$H(\vec{x}_1, ..., \vec{x}_N) = \{(+1, +1, ..., +1), (-1, +1, ..., +1), ... (-1, -1, ..., -1)\}$$

Dichotomies

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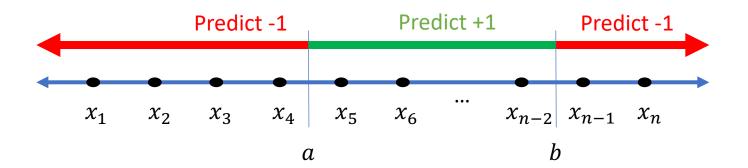
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• What is $m_H(N)$?

$$m_H(N) = N + 1$$

What is $m_H(N)$?

- H = Positive Intervals
 - Data points are in one-dimensional space
 - Choose two thresholds. Predict +1 within the interval, -1 outside



- H = Convex Sets
 - Data points are in 2-dimensional space
 - Hypothesis is represented by a convex set

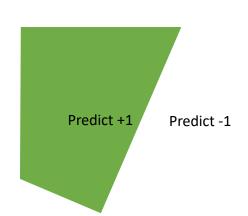


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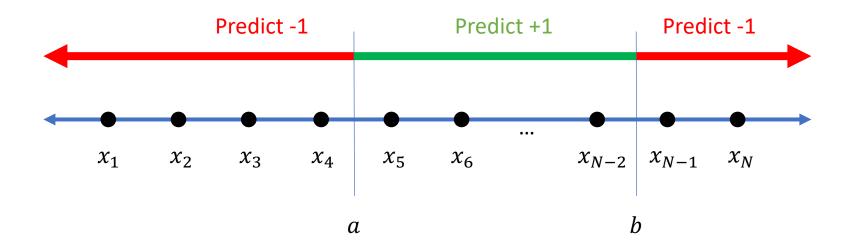
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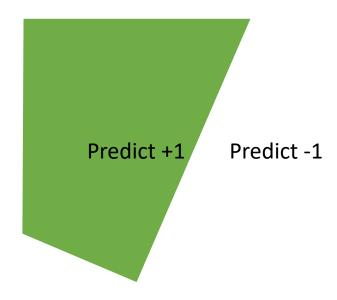


Example: H = Positive Intervals



- What is $m_H(N)$?
 - $m_H(N) = {N+1 \choose 2} + 1 = \frac{N^2}{2} + \frac{N}{2} + 1$

Example: H = Convex Sets



- What is $m_H(N)$?
 - $m_H(N) = 2^N$

Note: $m_H(N) \le 2^N$ for all H and all N (There are only 2^N possible label combinations for N points)

Why Growth Function?

- Growth function $m_H(N)$
 - Largest number of "effective" hypothesis H can induce on N data points
 - A more precise "complexity" measure for H
 - Goal: Replace M in finite-hypothesis analysis with $m_H(N)$

• With prob at least
$$1 - \delta$$
, $E_{out}(g) \le E_{in}(g) + \sqrt{\frac{1}{2N} ln \frac{2M}{\delta}}$

• Theorem: VC Inequality (1971)

With prob at least $1 - \delta$

$$E_{out}(g) \le E_{in}(g) + \sqrt{\frac{8}{N}} \ln \frac{4m_H(2N)}{\delta}$$

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Growth Functions for Other *H*

- H = 2-D Perceptron
 - What is $m_H(3)$
 - What is $m_H(4)$

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- Exactly calculate growth function is generally hard!
- Next lecture
 - Discuss how we can "bound" the growth function
 - Introduce the notion of VC dimension