# CSE 417T Introduction to Machine Learning

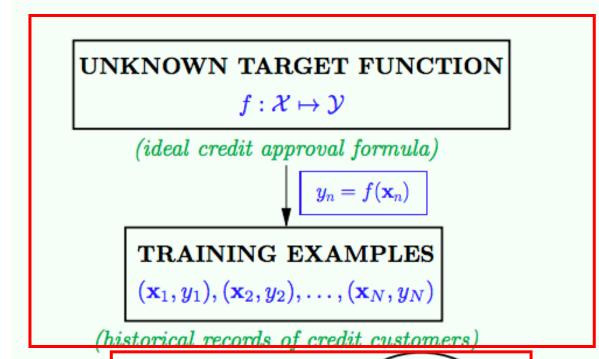
Lecture 3

Instructor: Chien-Ju (CJ) Ho

# Logistics

- Course website and Piazza
  - Website: http://chienjuho.com/courses/cse417t/
  - Piazza: <a href="http://piazza.com/wustl/fall2022/cse417t">http://piazza.com/wustl/fall2022/cse417t</a>
  - Make sure you follow both regularly
- Office hours
  - Will be announced later this week
  - Will start next week
- Homework 1
  - Will be announced later this week
  - A mixture of math questions and programming questions
    - Programming language: Python (We won't teach you how to program Python)

# Recap



 $\mathcal{H}$ 

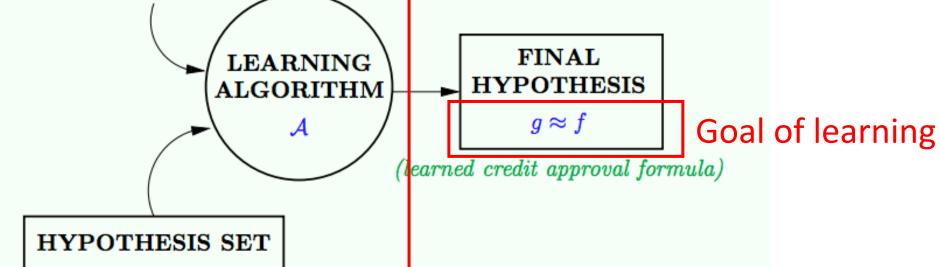
 $(set\ of\ candidate\ formulas)$ 

Given by the learning problem

learning model (example:

H: Perceptron

A: PLA)



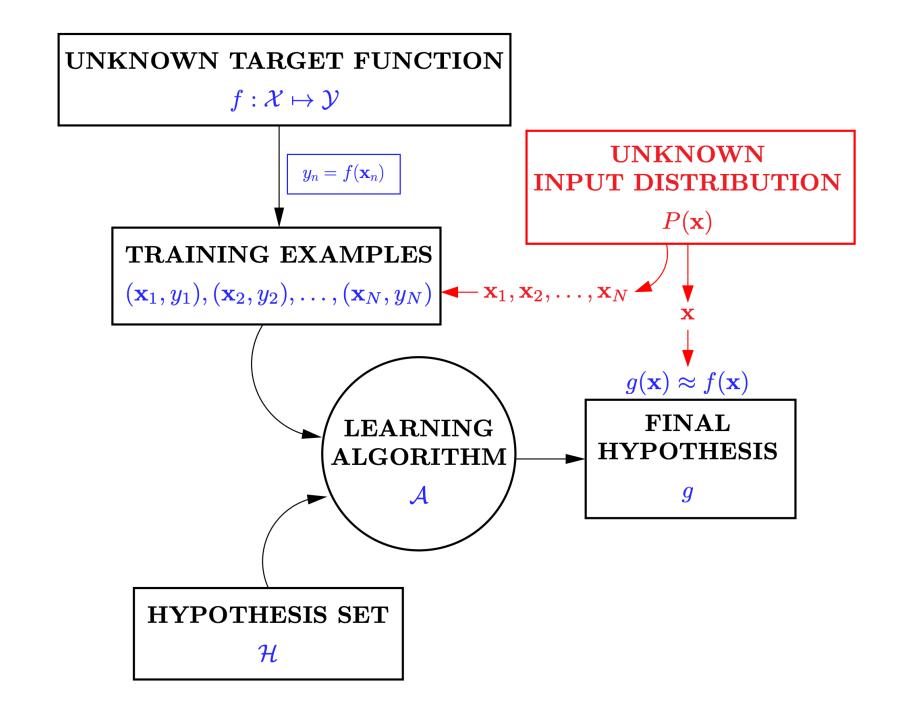
# Goal of Learning: Generalization

• Given training data, find  $g \approx f$  on the unseen test data.

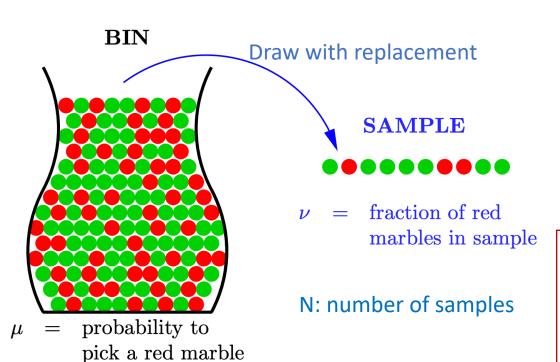
This goal is generally impossible without assumptions.

#### Key assumption of ML

Training data points and test data points are i.i.d. drawn from the same (unknown) distribution



# A Thought Experiment about Probability



What can we say about  $\mu$  from  $\nu$ ?

Law of large numbers

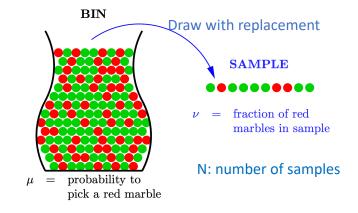
• When  $N \to \infty$ ,  $\nu \to \mu$ 

#### **Hoeffding's Inequality**

•  $\Pr[|\mu - \nu| > \epsilon] \le 2e^{-2\epsilon^2 N}$  for any  $\epsilon > 0$ 

### Connection to Learning

- Let each marble represent a point  $\vec{x}$ , drawn from unknown  $P(\vec{x})$ 
  - Dataset  $D = \{(\vec{x}_1, y_1), ..., (\vec{x}_N, y_N)\}$
  - Recall that  $y_n = f(\vec{x}_n)$  (will discuss noisy target function f later in the semester)
- "Fix" a hypothesis h
  - For each marble  $\vec{x}$ , color it as below
    - If  $h(\vec{x}) = f(\vec{x})$ , color it as green marble [h is correct on  $\vec{x}$ ]
    - If  $h(\vec{x}) \neq f(\vec{x})$ , color it as red marble  $[h \text{ is wrong on } \vec{x}]$



With the above coloring

$$\mu = \Pr_{\vec{x} \sim P(\vec{x})} [h(\vec{x}) \neq f(\vec{x})]$$

$$\stackrel{\text{def}}{=} E_{out}(h) \quad \text{[Out-of-sample error of } h \text{]}$$

$$\nu = \frac{1}{N} \sum_{n=1}^{N} \mathbb{I}[h(\vec{x}_n) \neq f(\vec{x}_n)]$$

$$\stackrel{\text{def}}{=} E_{in}(h) \quad \text{[in-sample error of } h\text{]}$$

# Connection to Learning

- $E_{out}(h)$ : What we really want to know but unknown to us
- $E_{in}(h)$ : What we can calculate from dataset

• Fixed a h, What can we say about  $E_{out}(h)$  from  $E_{in}(h)$ ?

#### **Hoeffding's Inequality**

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

• This is verification, not learning!

# Verification vs. Learning

#### Verification

- I have a hypothesis h
- I know  $E_{in}(h)$ , i.e., how well h performs in my dataset
- I can infer what  $E_{out}(h)$  (how well h will perform for unseen data) might be

#### Learning

- Given a dataset D and hypothesis set H
- Apply some learning algorithm, that outputs a  $g \in H$
- Know  $E_{in}(g)$
- Want to infer  $E_{out}(g)$

# Connection to "Real" Learning

- Given a finite hypothesis set  $H = \{h_1, ..., h_M\}$ 
  - Will discuss the infinite case in the next few lectures.
- Apply some learning algorithm on D, output a  $g \in H$ 
  - For example, choosing the hypothesis that minimizes in-sample error
    - $g = argmin_{h \in H} E_{in}(h)$
- Can we apply Hoeffding's inequality and claim

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

#### No!

# Today's Lecture

The notes are not intended to be comprehensive. Let me know if you spot errors.

- Three fair coins, numbered by 1, 2, 3. Flip each 10 times.
- Question: (choosing from >5, =5, or <5)</li>
- Ans: = 5 For coin 1, what's the expected number of heads among 10 flips?
- Ans: = 5 Randomly choose a coin, what's the expected number of heads for this coin?
- Ans: >5 Look at the realized flips and choose the coin with the largest number of heads. What is the expected number of heads for the coin?
- Ans: = 5 Without observing the flips, choose the coin anyway you like, what is the expected number of heads of the 10 flips for this coin?
  - You will simulate this process (with 1,000 coins) in HW1.

• If you toss a fair coin 10 times, the prob that you get heads 10 times is

$$2^{-10} = \frac{1}{1024}$$

• If you toss 1000 fair coins 10 times each, the probability that at least one coin comes up heads 10 times is

$$1 - \left(\frac{1023}{1024}\right)^{1000} \approx 62.36\%$$

- If each hypothesis is doing random guessing (i.e., tossing a fair coin), if we have 1000 hypothesis with 10 data points, more than 60% chance there will be at least one hypothesis with zero in-sample error
  - But that hypothesis is still random guessing and has 50% out-of-sample error

- Connects to learning
  - Coin -> Hypothesis
  - Coin flips -> Performance of hypothesis in training data D
- Choosing the hypothesis "before" or "after" looking at the data (knowing the realization of the data drawing) makes a big difference!

#### **Hoeffding's Inequality**

•  $\Pr[|\mu - \nu| > \epsilon] \le 2e^{-2\epsilon^2 N}$  for any  $\epsilon > 0$ 

Some graphical explanations

WE FOUND NO LINK BETWEEN PURPLE JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN BROWN JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN PINK JELLY BEANS AND ACNE (P>0.05).



WE FOUND NO LINK BETWEEN BLUE JELLY BEANS AND ACNE (P>0.05).



WE FOUND NO LINK BETWEEN TEAL JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN SALMON JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN RED JELLY BEANS AND ACNE (P > 0.05)



WE FOUND NO LINK BETWEEN TURQUOISE JELLY BEANS AND ACNE (P>0.05).



WE FOUND NO LINK BETWEEN MAGENTA JELLY BEANS AND ACNE (P > 0.05)



WE FOUND NO LINK BETWEEN YELLOW JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN GREY JELLY BEANS AND ACNE (P>0.05).



WE FOUND NO LINK BETWEEN TAN JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN CYAN JELLY BEANS AND ACNE (P > 0.05),



WE FOUND A LINK BETWEEN GREEN JELLY BEANS AND ACNE (P<0.05).



JE FOUND NO LINK BETWEEN MAUVE JELLY BEANS AND AONE (P>0.05),



WE FOUND NO LINK BETWEEN BEIGE JELLY BEANS AND ACNE (P>0.05).



WE FOUND NO LINK BETWEEN LILAC JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN BLACK JELLY BEANS AND ACNE (P > 0.05),

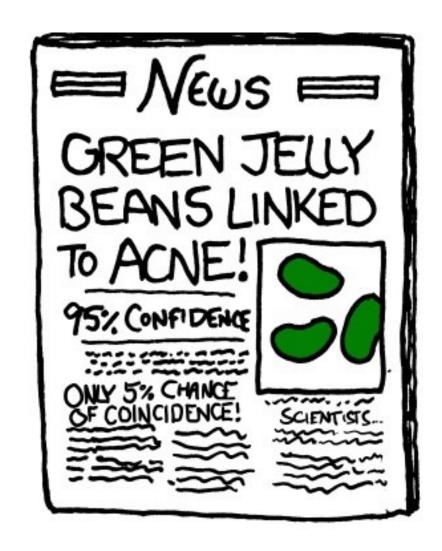


WE FOUND NO LINK BETWEEN PEACH JELLY BEANS AND ACNE (P > 0.05).



WE FOUND NO LINK BETWEEN ORANGE JELLY BEANS AND ACNE (P > 0.05).





# What Can We Do?

# Connection to "Real" Learning

- Given a finite hypothesis set  $H = \{h_1, ..., h_M\}$
- Apply some learning algorithm on D, output a  $g \in H$ 
  - For example, choosing the hypothesis that minimizes in-sample error
    - $g = argmin_{h \in H} E_{in}(h)$
- Question: What can we say about  $E_{out}(g)$  from  $E_{in}(g)$ ?

#### Derivations

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

For each fixed  $h \in H$ , we have  $\Pr[B(h)] \leq 2e^{-2\epsilon^2 N}$ 

- Recall g is selected from H (it could be any  $h \in H$ )
- What can we say about Pr[B(g)]?

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If g is selected from \{h_1, h_2\} B(g) \subseteq B(h_1) \cup B(h_2) \Pr[B(g)] \leq \Pr[B(h_1) \text{ or } B(h_2)] \leq \Pr[B(h_1)] + \Pr[B(h_2)] (Union Bound)
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For each fixed  $h \in H$ , we have  $\Pr[B(h)] \leq 2e^{-2\epsilon^2 N}$ 

- Recall g is selected from H (it could be any  $h \in H$ )
- What can we say about Pr[B(g)]?

$$\Pr[B(g)] \le \Pr[B(h_1) \text{ or } B(h_2) \text{ or } \dots \text{ or } B(h_M)]$$
  
 $\le \Pr[B(h_1)] + \Pr[B(h_2)] + \dots + \Pr[B(h_M)]$   
 $\le M \ 2e^{-2\epsilon^2 N}$ 

# Connection to "Real" Learning

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

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

- M can be considered as a proxy of the "complexity" of the hypothesis set
  - Will talk about what happens when  $M \to \infty$  in the next few lectures

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

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

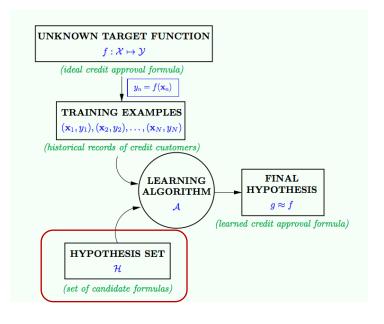
- Playing around with the math
  - Define  $\delta = \Pr[|E_{out}(g) E_{in}(g)| > \epsilon]$
  - We have  $\delta \le 2Me^{-2\epsilon^2N} \implies \epsilon \le \sqrt{\frac{1}{2N}\ln\frac{2M}{\delta}}$
- This means, with probability at least  $1-\delta$

• 
$$E_{out}(g) \le E_{in}(g) + \epsilon \le E_{in}(g) + \sqrt{\frac{1}{2N} \ln \frac{2M}{\delta}}$$

#### More Discussion

• With probability at least  $1-\delta$ 

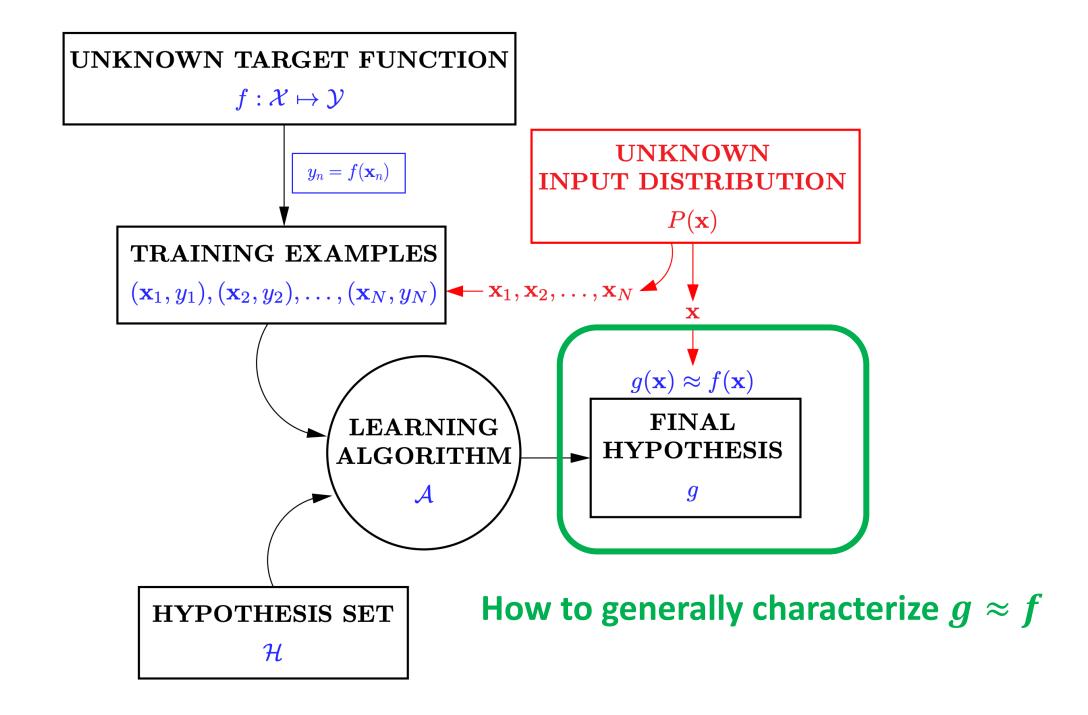
$$E_{out}(g) \le E_{in}(g) + \sqrt{\frac{1}{2N}} \ln \frac{2M}{\delta}$$



Consider M as a proxy measure on the "complexity" of H

- Our ultimate goal is to have a small  $E_{out}(g)$ 
  - There is a tradeoff of choosing M (what "learning model" to use)
    - Increase  $M \rightarrow \text{Smaller } E_{in}(g)$  (more hypothesis to "fit" the training data)
    - Increase  $M \rightarrow Larger \epsilon$
  - It also depends on N, the number of data points you have
    - A small number of data points => use simple models (e.g., linear models)
    - Complex models (e.g., deep learning) work when you have a lot of data

# 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  $g = \underset{h \in \mathcal{H}}{\operatorname{argmin}} E(h, f)$
- 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 the error function?

- Consideration 1: Properties of domain applications
- Example: Fingerprint recognition
  - Input: fingerprints
  - Outputs: whether the person is authorized

		$f(\overrightarrow{x})$		
		+1	-1	
$h(\vec{x})$	+1	No error	False positive	
	-1	False negative	No error	

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

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

#### How to choose the error function?

Consideration 1: Properties of application problems

- Consideration 2: Computation
  - ML Algorithm is essentially doing optimization (finding g with smallest error)

$$g = \underset{h \in \mathcal{H}}{\operatorname{argmin}} E(h, f)$$

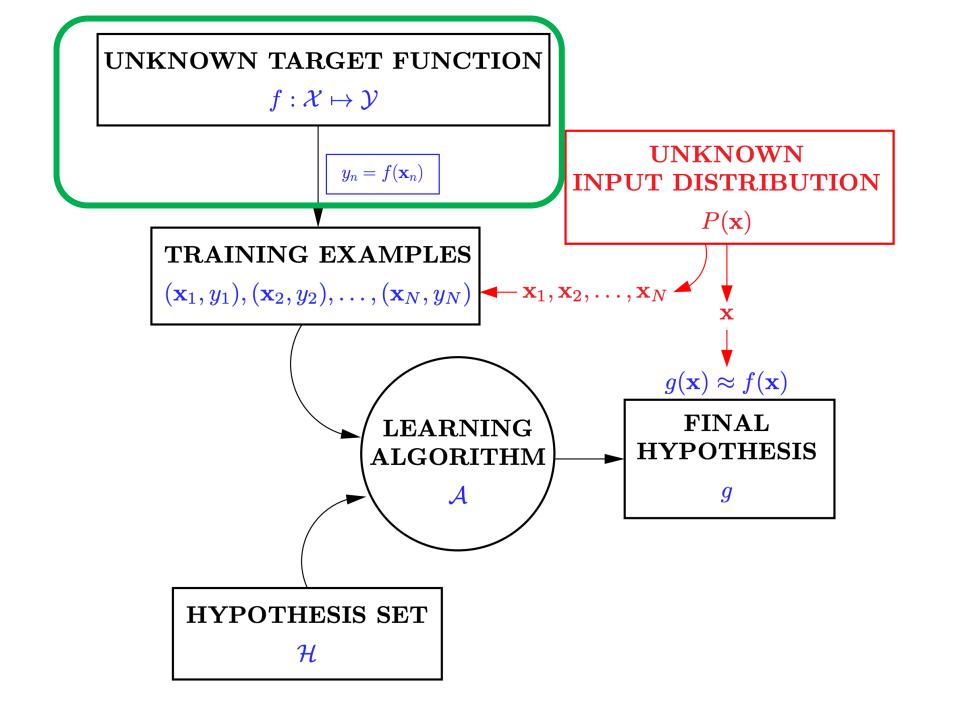
- Choosing the error that is "easier" to optimize
  - e.g., if the error function is convex, continuous, differentiable, we usually have efficient algorithms

#### How to choose the error function?

Consideration 1: Properties of application problems

Consideration 2: Computation

- Specifying the error function is part of setting up the learning problem
  - It impacts what you eventually learn



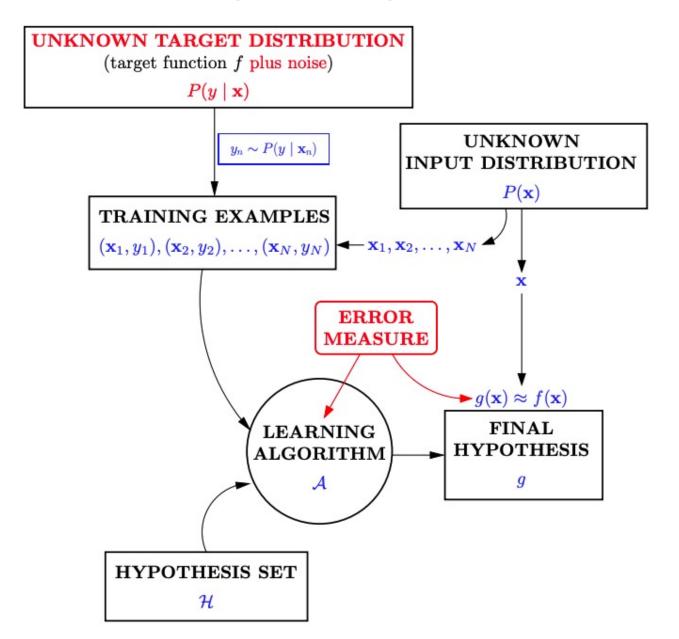
# Noisy Target

- What if there doesn't exist f such that  $y = f(\vec{x})$ ?
  - f is stochastic instead of deterministic

- Common approach
  - Instead of a target function, define a target <u>distribution</u>
  - Instead of  $y = f(\vec{x})$ , y is drawn from a conditional distribution  $P(y|\vec{x})$
  - $y = f(\vec{x}) + \epsilon$  where  $\epsilon$  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

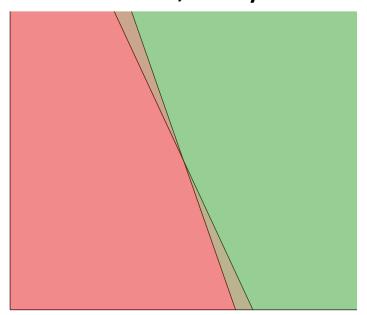
- Define "bad event of h" B(h) as  $|E_{out}(h) E_{in}(h)| > \epsilon$
- If g is selected from  $\{h_1, h_2\}$ 
  - $B(g) \subseteq B(h_1) \cup B(h_2)$
  - $\Pr[B(g)] \le \Pr[B(h_1) \text{ or } B(h_2)]$  $\le \Pr[B(h_1)] + \Pr[B(h_2)]$  (Union Bound)

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

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

# Do Bad Events Overlap?

Oftentimes, they overlap a lot!



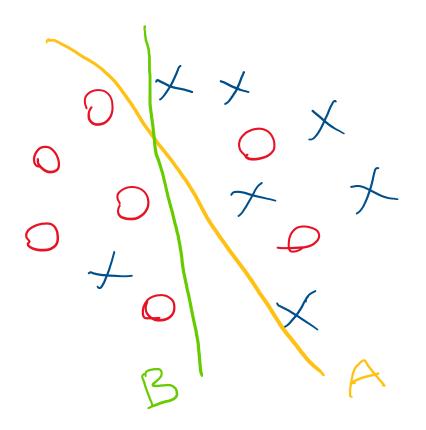
The two linear separators on the left make the same predictions for most points.

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?



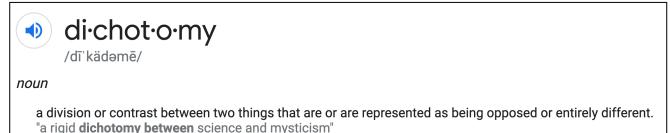
Any difference between A and B?

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

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