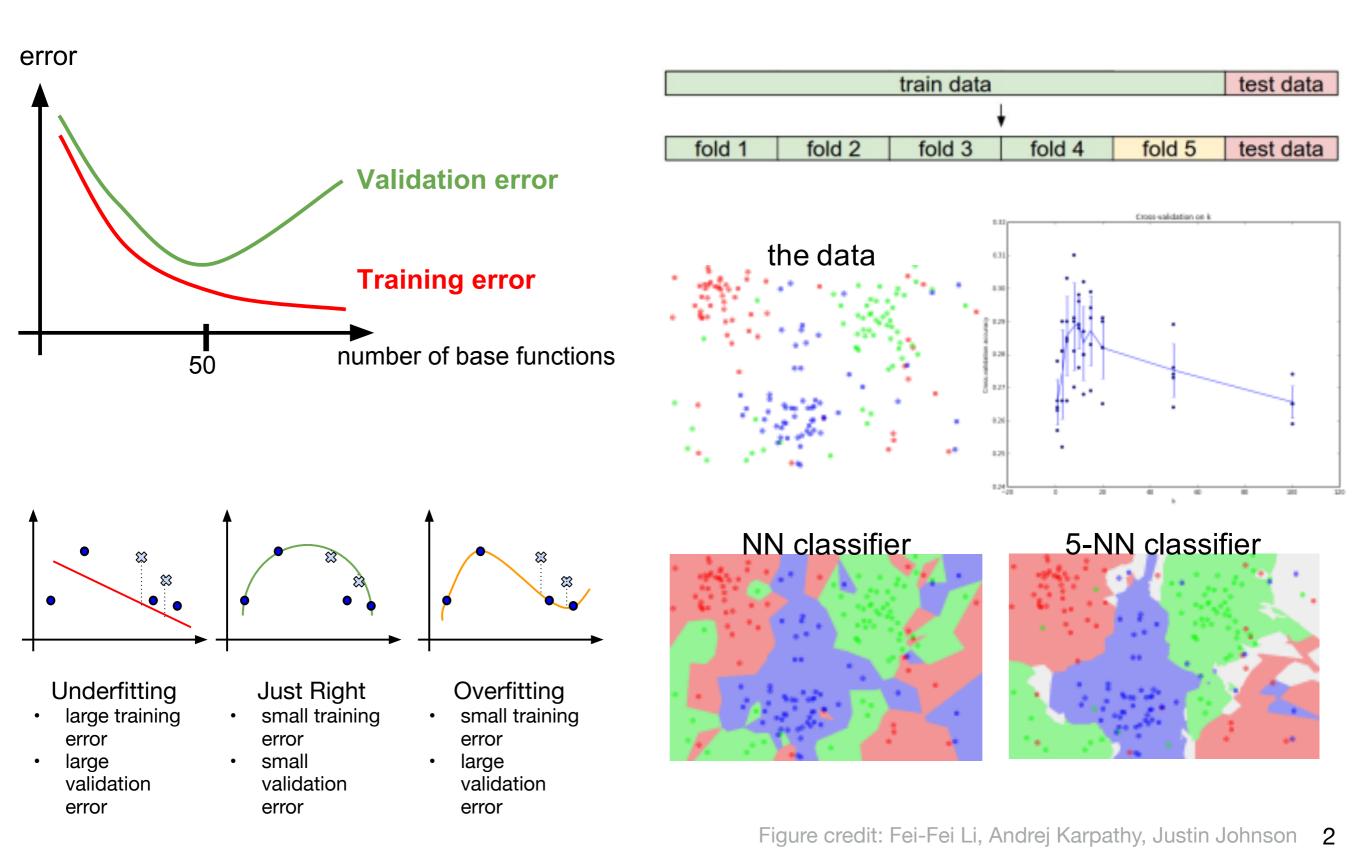
# AND THE AND TH

Learning theory Probability Review



Erkut Erdem // Hacettepe University // Fall 2023

#### Last time... Regularization, Cross-Validation



### Today

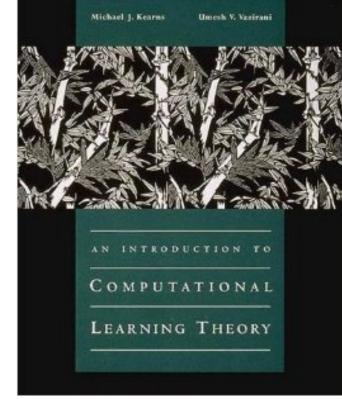
- Learning Theory
- Probability Review

### Learning Theory: Why ML Works

### Computational Learning Theory

- Entire subfield devoted to the mathematical analysis of machine learning algorithms
- Has led to several practical methods:
  - PAC (probably approximately correct) learning
     → boosting
  - VC (Vapnik–Chervonenkis) theory
    - → support vector machines

Annual conference: Conference on Learning Theory (COLT)



### The Role of Theory

- Theory can serve two roles:
  - It can justify and help understand why common practice works.
     theory after theory
  - It can also serve to suggest new algorithms and approaches that turn out to work well in practice.

#### Often, it turns out to be a mix!

### The Role of Theory

- Practitioners discover something that works surprisingly well.
- Theorists figure out why it works and prove something about it.
  - In the process, they make it better or find new algorithms.
- Theory can also help you understand what's possible and what's not possible.

### Learning and Inference

The inductive inference process:

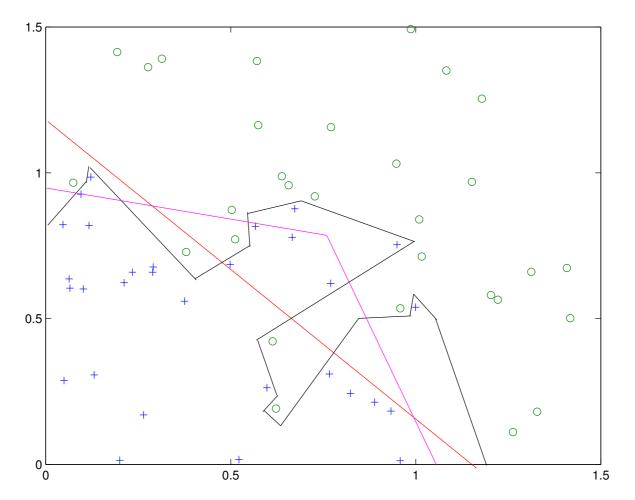
- 1. Observe a phenomenon
- 2. Construct a model of the phenomenon
- 3. Make predictions
- This is more or less the definition of natural sciences !
- The goal of Machine Learning is to automate this process
- The goal of Learning Theory is to formalize it.

### Pattern recognition

- We consider here the supervised learning framework for pattern recognition:
  - Data consists of pairs (instance, label)
  - Label is +1 or -1
  - Algorithm constructs a function (instance  $\rightarrow$  label)
  - Goal: make few mistakes on future unseen instances

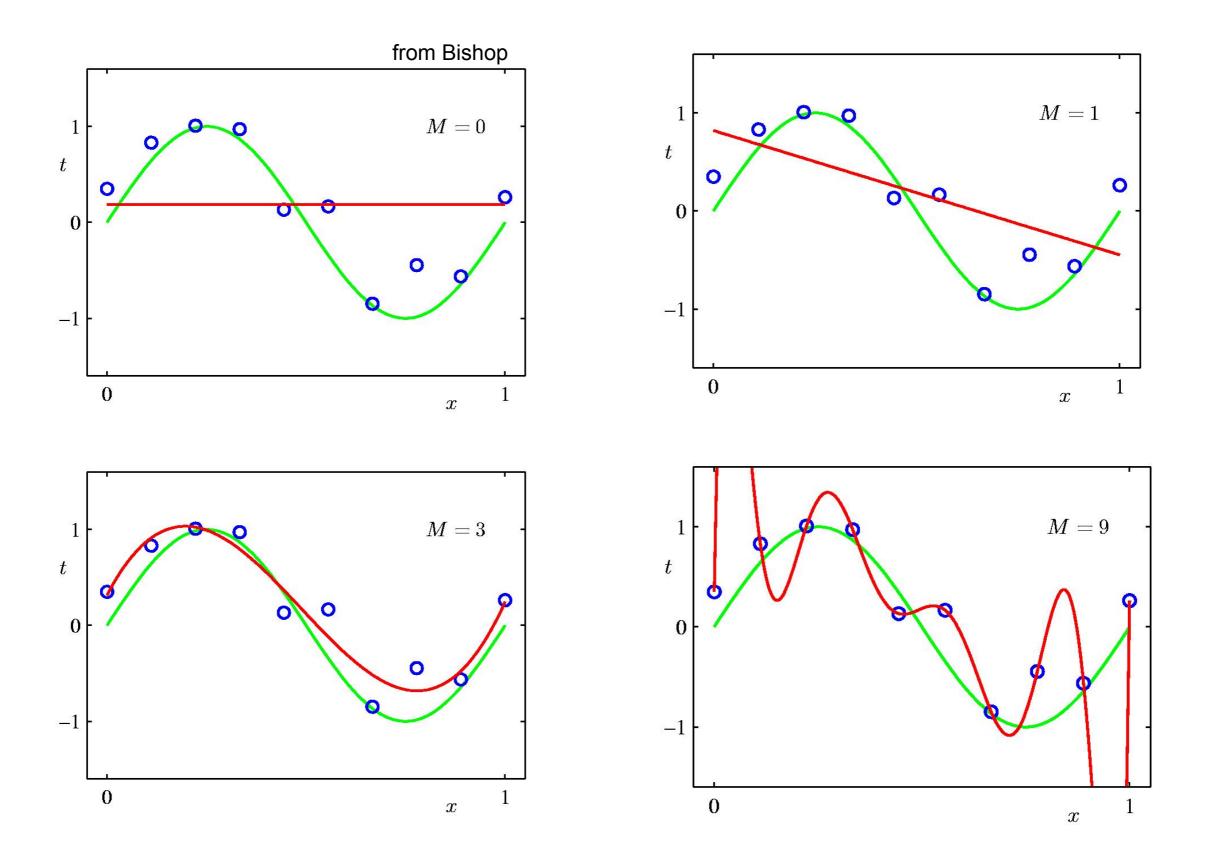
### Approximation/Interpolation

 It is always possible to build a function that fits exactly the data.



• But is it reasonable?

### Which Fit is Best?



11

### Occam's Razor

 Idea: look for regularities in the observed phenomenon

These can be **generalized** from the observed past to the future

⇒ choose the simplest consistent model

- How to measure simplicity ?
  - Physics: number of constants
  - Description length
  - Number of parameters



William of Occam (c. 1288 – c. 1348)

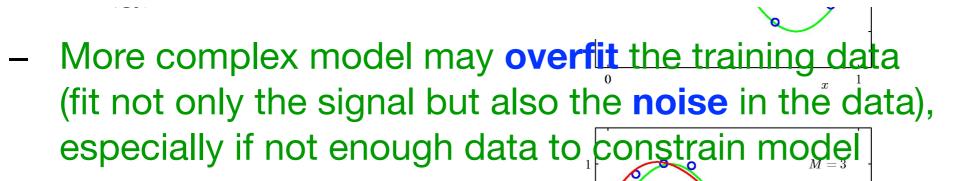
### No Free Lunch

#### • No Free Lunch

- if there is no assumption on how the **past** is related to the future, prediction is impossible
- if there is no restriction on the possible phenomena, generalization is impossible
- We need to make assumptions
- Simplicity is not absolute
- Data will never replace knowledge
- Generalization = data + knowledge

#### concepts

M = 1



 $w_0^\star$ 

 $w_1^\star w_2^\star$ 

 $w_3^\star w_4^\star$ 

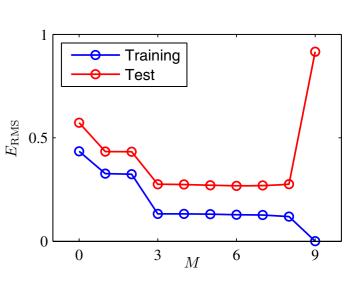
 $w_{6}^{\star} w_{7}^{\star} w_{8}^{\star}$ 

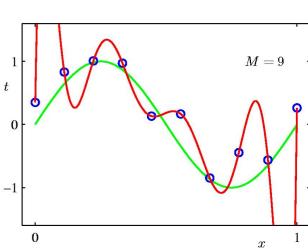
 $w_{q}^{\star}$ 

- One method of assessing fit:
  - test generalization = model's ability to predict the held out data  $\frac{1}{0}$
- Regularization

$$\widetilde{E}(\mathbf{w}) = \frac{1}{2} \sum_{n=1}^{N} \{y(x_n, \mathbf{w}) - t_n\}^2 + \frac{\lambda}{2} \|\mathbf{w}\|^2$$

$$\|\mathbf{w}\|^2 \equiv \mathbf{w}^{\mathrm{T}}\mathbf{w} = w_0^2 + w_1^2 + \ldots + w_M^2$$





### Probably Approximately Correct (PAC) Learning

- A formalism based on the realization that the best we can hope of an algorithm is that
  - It does a good job most of the time (probably approximately correct)

### Probably Approximately Correct (PAC) Learning

- Consider a hypothetical learning algorithm
  - We have 10 different binary classification data sets.
  - For each one, it comes back with functions  $f_1, f_2, \ldots, f_{10}$ .
    - For some reason, whenever you run *f*<sub>4</sub> on a test point, it crashes your computer. For the other learned functions, their performance on test data is always at most 5% error.
    - If this situtation is guaranteed to happen, then this hypothetical learning algorithm is a PAC learning algorithm.
      - It satisfies probably because it only failed in one out of ten cases, and it's approximate because it achieved low, but non-zero, error on the remainder of the cases.

### PAC Learning

**Definitions 1.** An algorithm A is an  $(\epsilon, \delta)$ -PAC learning algorithm if, for all distributions D: given samples from D, the probability that it returns a "bad function" is at most  $\delta$ ; where a "bad" function is one with test error rate more than  $\epsilon$  on D.

### PAC Learning

- Two notions of efficiency
  - Computational complexity: Prefer an algorithm that runs quickly to one that takes forever
  - Sample complexity: The number of examples required for your algorithm to achieve its goals

**Definition:** An algorithm  $\mathcal{A}$  is an efficient  $(\epsilon, \delta)$ -PAC learning algorithm if it is an  $(\epsilon, \delta)$ -PAC learning algorithm whose runtime is polynomial in  $\frac{1}{\epsilon}$  and  $\frac{1}{\delta}$ .

In other words, to let your algorithm to achieve 4% error rather than 5%, the runtime required to do so should not go up by an exponential factor!

#### Example: PAC Learning of Conjunctions

- Data points are binary vectors, for instance  $\mathbf{x} = \langle 0, 1, 1, 0, 1 \rangle$
- Some Boolean conjunction defines the true labeling of this data (e.g.  $x_1 \wedge x_2 \wedge x_5$ )
- There is some distribution  $\mathcal{D}_X$  over binary data points (vectors)  $\mathbf{x} = \langle x_1, x_2, \dots, x_D \rangle$ .
- There is a fixed concept conjunction c that we are trying to learn.
- There is no noise, so for any example *x*, its true label is simply  $y = c(\mathbf{x})$

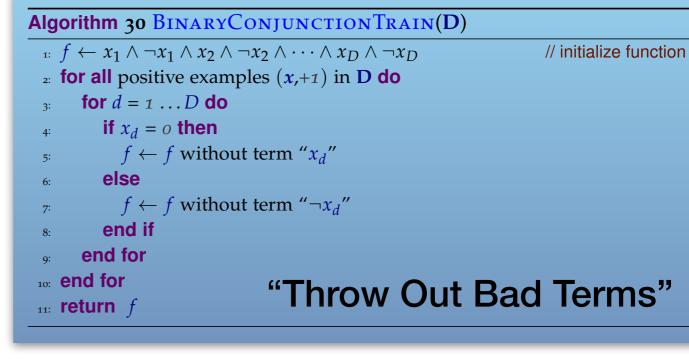
#### • Example:

- Clearly, the true formula cannot include the terms  $x_1, x_2, \neg x_3, \neg x_4$ 

y	$ x_1 $	$x_2$	<i>x</i> <sub>3</sub>	$x_4$
+1 +1	0			1
+1	0	1	1	1
-1	1	1	0	1

# Example: PAC Learning of Conjunctions

y	$x_1$	$x_2$	<i>x</i> <sub>3</sub>	$x_4$	
+1	0	0	1	1	
+1	0	1	1	1	
-1	1	1	0	1	

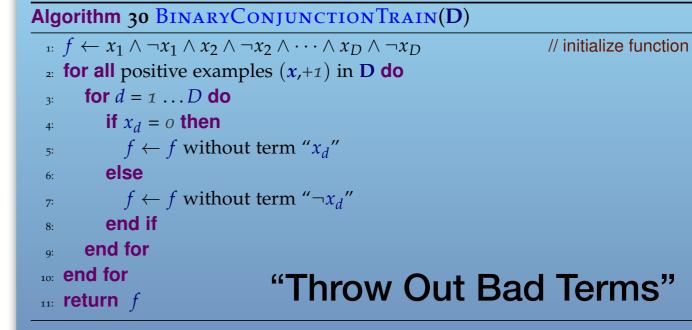


$$f^{0}(\mathbf{x}) = x_{1} \wedge \neg x_{1} \wedge x_{2} \wedge \neg x_{2} \wedge x_{3} \wedge \neg x_{3} \wedge x_{4} \wedge \neg x_{4}$$
$$f^{1}(\mathbf{x}) = \neg x_{1} \wedge \neg x_{2} \wedge x_{3} \wedge x_{4}$$
$$f^{2}(\mathbf{x}) = \neg x_{1} \wedge x_{3} \wedge x_{4}$$
$$f^{3}(\mathbf{x}) = \neg x_{1} \wedge x_{3} \wedge x_{4}$$

- After processing an example, it is guaranteed to classify that example correctly (provided that there is no noise)
  - Computationally very efficient
    - Given a data set of N examples in D dimensions, it takes O (ND) time to process the data. This is linear in the size of the data set.

# Example: PAC Learning of Conjunctions

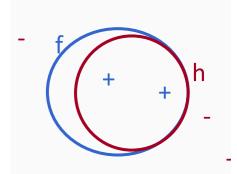
y	$x_1$	$x_2$	$x_3$	$x_4$	
+1	0	0	1	1	
+1	0	1	1	1	
-1	1	1	0	1	



• Is this an efficient  $(\varepsilon, \delta)$ -PAC learning algorithm?

 $f = x_2 \land x_3 \land x_4 \land x_5 \land x_{100} \qquad \qquad h = x_1 \land x_2 \land x_3 \land x_4 \land x_5 \land x_{100}$ 

- Claim 1: Any hypothesis  $c_{0}$  sistent with the training training data will only make mistakes on positive future examples. Why?
- A mistake will occur only if some literal z (in our example x<sub>1</sub>) is present in h but not in f



- This mistake can cause a positive example to be predicted as negative by h. Specifically: x<sub>1</sub> = 0, x<sub>2</sub> =1, x<sub>3</sub>=1, x<sub>4</sub>=1, x<sub>5</sub>=1, x<sub>100</sub>=1
- The reverse situation can never happen
  - For an example to be predicted as positive in the training set, every relevant literal must have been present

• **Theorem:** Suppose we are learning a conjunctive concept with n dimensional Boolean features using m training examples. If

$$m > \frac{n}{\epsilon} \left( \log(n) + \log\left(\frac{1}{\delta}\right) \right)$$

Poly in n, 1/ $\delta$ , 1/ $\epsilon$ 

then, with probability > 1 -  $\delta$ , the error of the learned hypothesis err<sub>D</sub>(h) will be less than  $\epsilon$ .

If we see these many training examples, then the algorithm will produce a conjunction that, with high probability, will make few errors

• Theorem: Suppose we are learning a conjunctive concept with n dimensional Boolean features using m training examples. If n((1))

$$m > \frac{n}{\epsilon} \left( \log(n) + \log\left(\frac{1}{\delta}\right) \right)$$

then, with probability > 1 -  $\delta$ , the error of the learned hypothesis err<sub>D</sub>(h) will be less than  $\epsilon$ .

Let's prove this assertion

### **Proof Intuition**

 $f = x_2 \wedge x_3 \wedge x_4 \wedge x_5 \wedge x_{100} \qquad \qquad h = x_1 \wedge x_2 \wedge x_3 \wedge x_4 \wedge x_5 \wedge x_{100}$ 

- What kinds of examples would drive a hypothesis to make a mistake?
- Positive examples, where  $x_1$  is absent
  - f would say true and h would say false
- None of these examples appeared during training
  - Otherwise x<sub>1</sub> would have been eliminated
- If they never appeared during training, maybe their appearance in the future would also be rare!
  - Let's quantify our surprise at seeing such examples

- Let p(z) be the probability that, in an example drawn from D, the feature z is absent but the example has a positive label
  - That is, after training is done, p(z) is the probability that in a randomly drawn example, the literal z causes a mistake
  - For any z in the target function, p(z) = 0

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Remember that there will only be mistakes on positive examples for this toy problem

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<(0,1,1,1,1,0,...0,1,1), 1>

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$$h = x_1 \wedge x_2 \wedge x_3 \wedge x_4 \wedge x_5 \wedge x_{100}$$

<(0,1,1,1,1,0,...0,1,1), 1 >  $p(x_1)$ : Probability that this situation occurs

- Let p(z) be the probability that, in an example drawn from D, the feature z is absent but the example has a positive label
  - That is, after training is done, p(z) is the probability that in a randomly drawn example, the literal z causes a mistake
  - For any z in the target function, p(z) = 0

We know that 
$$err_D(h) \leq \sum_{z \in h} p(z)$$

via direct application of the union bound

Union bound

For a set of events, probability that at least one of them happens < the sum of the probabilities of the individual events

• Call a literal z bad if  $p(z) > \frac{\epsilon}{n}$ 

n = dimensionality

- Intuitively, a bad literal is one that has a significant probability of not appearing with a positive example
  - (And, if it appears in all positive training examples, it can cause errors)

If there are no bad literals, then  $err_D(h) < \epsilon$ 

$$err_D(h) \le \sum_{z \in h} p(z)$$

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  - (And, if it appears in all positive training examples, it can cause errors)
  - If there are no bad literals, then  $err_D(h) < \epsilon$ 
    - Why? Because

$$err_{D}(h) \leq \sum_{z \in \mathcal{D}} p(z) \\ p(z)$$

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- n = dimensionality
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 $z \in h$ 

- (And, if it appears in all positive training examples, it can cause errors)
- If there are no bad literals, then  $err_D(h) < \epsilon$
- Why? Because  $err_D(h) \leq \sum_{z \in \mathcal{V}} p(z)$  $err_D(h) \leq \sum_{z \in \mathcal{V}} p(z)$

Let us try to see when this will not happen

• Call a literal z bad if  $p(z) > \frac{\epsilon}{n}$ 

n = dimensionality

- Intuitively, a bad literal is one that has a significant probability of not appearing with a positive example
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What if there are bad literals?

$$err_D(h) \le \sum_{z \in h} p(z)$$

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#### What if there are bad literals?

- Let z be a bad literal
- What is the probability that for will not (see eliminated by one training example?  $z \in h$

•

• Call a literal z bad if  $p(z)_{p(z)} \ge \frac{\epsilon}{\pi}$ 

n = dimensionality

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#### What if there are bad literals?

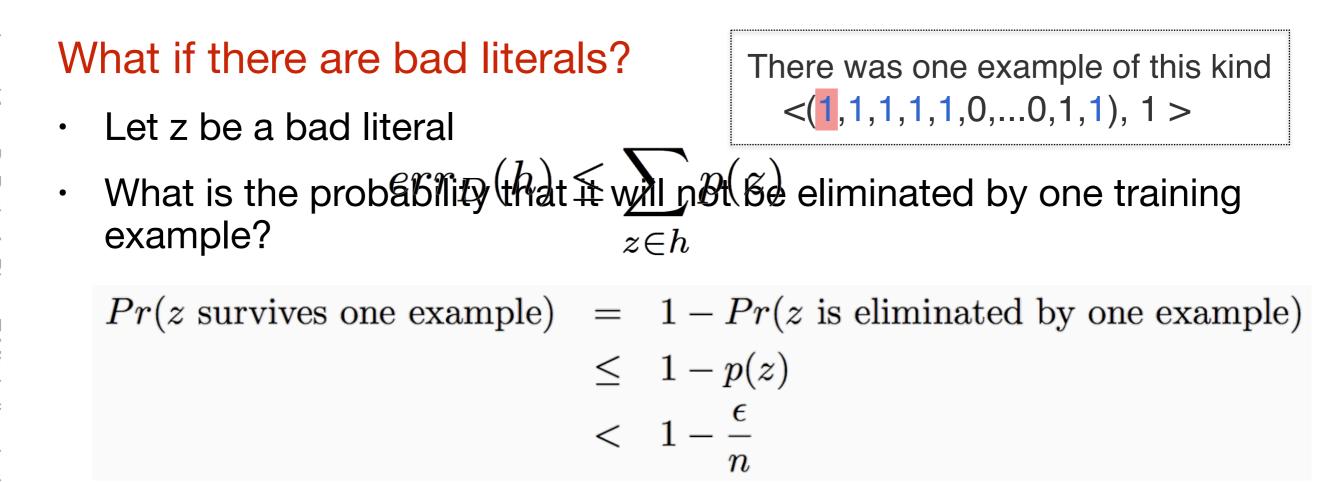
- Let z be a bad literal
- What is the probability that for will not (see eliminated by one training example?  $z \in h$

Pr(z survives one example) = 1 - Pr(z is eliminated by one example)

$$\leq 1 - p(z) \ < 1 - rac{\epsilon}{n}$$

• Call a literal z bad if  $p(z) > \frac{\epsilon}{\pi}$ 

- n = dimensionality
- Intuitively, a bad literal is one that has a significant probability of not appearing with a positive example
  - (And, if it appears in all positive training examples, it can cause errors)



• What we know so far:

n = dimensionality

 $Pr(A \text{ bad literal is not eliminated by one example}) < 1 - \frac{\epsilon}{n}$ 

 $Pr(A \text{ bad literal survives } m \text{ examples}) < \left(1 - \frac{\epsilon}{n}\right)^m$ 

 $Pr(\text{Any bad literal survives } m \text{ examples}) < n\left(1 - \frac{\epsilon}{n}\right)^m$ 

• What we know so far:

n = dimensionality

 $P_r(A_{bad}, i_{eral}, i_{snot}, f_{efiminated}) > (A_{bad}, i_{eral}, i_{snot}, f_{eral}, f_{eral}, i_{snot}, f_{eral}, f_{eral},$ 

But say we have m training examples. Then

 $Pr(A \text{ bad literal survives } m \text{ examples}) < \left(1 - \frac{\epsilon}{n}\right)^m$ 

$$P_{n}(A_{ny}bad literal survives mexamples}) < n((1 - \frac{\epsilon}{n}))^{n}$$

What we know so far:

n = dimensionality

 $Pr(A \text{bad}|\text{iteral}; \text{is not eliminated by one example}) \leq 1 - \frac{\epsilon}{n} \frac{1}{n} \frac{1}{\epsilon}$  $Pr(A \text{ bad literal}; \text{is not eliminated by one example}) < 1 - \frac{\epsilon}{n} \frac{1}{n} \frac{1}{\epsilon}$ 

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There are at most n bad literals. So  $P_{r}(Any bad literal survives mexamples) < n((1-\frac{\epsilon}{n})^{m} P_{r}(Any bad literal survives m examples) < n((1-\frac{\epsilon}{n})^{m})^{m}$ 

#### $Pr(A \text{ bad literal survives } m \text{ examples}) < (1 - \frac{c}{n})^m$ Learning Conjunctions: Analysis

 $Pr(\text{Any bad literal survives } m \text{ examples}) < n\left(1 - \frac{\epsilon}{m}\right)^m$ 

- We want this probability to be small
- Why? So that we can choose enough training examples so that the probability that any z survives all of them is less than some  $\delta$ .

 $Pr(A \text{ bad literal survives } m \text{ examples}) < \left(1 - \frac{\epsilon}{n}\right)^{m}$  Learning Conjunctions: Analysis  $Pr(Any \text{ bad literal survives } m \text{ examples}) < n \left(1 - \frac{\epsilon}{n}\right)^{m}$   $Pr(Any \text{ bad literal survives } m \text{ examples}) < n \left(1 - \frac{\epsilon}{n}\right)^{m}$ 

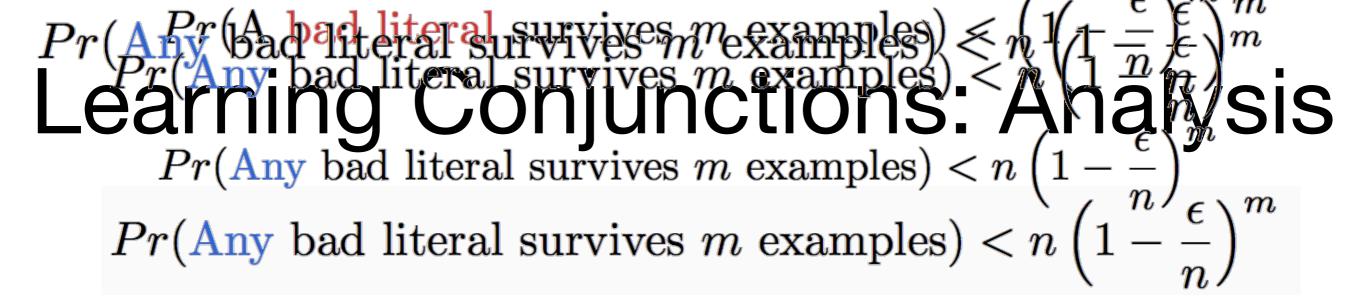
- We want this probability to be small
- Why? So that we can choose enough training examples so that the probability that any z survives all of them is less than some  $\delta$ .

That is, we want 
$$n\left(1-\frac{\epsilon}{n}\right)^m < \delta$$

$$ne^{-\frac{m\epsilon}{n}} < \delta$$

$$\begin{array}{l} Pr(\operatorname{Any} (\operatorname{Any} \operatorname{bad} \operatorname{diteral survives} m \operatorname{examples}) \leqslant n \left(1 + \frac{\epsilon}{n}\right)^{m} \\ \begin{array}{l} \mathsf{Learning} & \mathsf{Conjunctions:} & \mathsf{Analysis} \\ Pr(\operatorname{Any} \operatorname{bad} \operatorname{literal survives} m \operatorname{examples}) < n \left(1 - \frac{\epsilon}{n}\right)^{m} \\ \end{array} \\ \begin{array}{l} Pr(\operatorname{Any} \operatorname{bad} \operatorname{literal survives} m \operatorname{examples}) < n \left(1 - \frac{\epsilon}{n}\right)^{m} \end{array} \end{array}$$

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- Why? So that we can choose enough training examples so that the probability that any *ε* survives all of them is less than some δ.
   That is, we want n (1 n)
  - We know that  $1 x < e^{-x}$ . So it is sufficient to require  $\frac{ne^{-\frac{m\epsilon}{n}} < \delta}{ne^{-\frac{n}{n}} < \delta}$



- We want this probability to be small
- Why? So that we can choose enough training examples so that the probability that any *z* survives all of them is less than some δ.
   *n* n1(1--+) < δ</li>
   That is, we want n(1--+) < δ</li>
  - We know that  $1 x < e^{-x}$ . So it is sufficient to require  $ne^{-\frac{me}{n}} < \delta$

Or equivalently,

$$m > \frac{n}{\epsilon} \left( \log(n) + \log\left(\frac{1}{\delta}\right) \right)$$

• To guarantee a probability of failure (i.e, error >  $\epsilon$ ) that is the set of that the set of the the set of the set of the number of examples we need is

$$m > \frac{n}{\epsilon} \left( \log(n) + \log\left(\frac{1}{\delta}\right) \right)$$
 [Pole

Poly in n, 1/
$$\delta$$
, 1/ $\epsilon$ 

- That is, if m has this property, then
  - With probability 1  $\delta$ , there will be no bad literals,
  - Or equivalently, with probability 1  $\delta$ , we will have err<sub>D</sub>(h) <  $\epsilon$

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 Poly in n, 1/ $\delta$ , 1/ $\epsilon$ 

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#### What does this mean:

If  $\epsilon = 0.1$  and  $\delta = 0.1$ , then for n = 100, we need 6908 training examples

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 Poly in

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- If  $\epsilon = 0.1$  and  $\delta = 0.1$ , then for n = 100, we need 6908 training examples
- If  $\epsilon = 0.1$  and  $\delta = 0.1$ , then for n = 10, we need only 461 examples

•

n, 1/ $\delta$ , 1/ $\epsilon$ 

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#### What does this mean:

- If  $\epsilon = 0.1$  and  $\delta = 0.1$ , then for n = 100, we need 6908 training examples
- If  $\epsilon = 0.1$  and  $\delta = 0.1$ , then for n = 10, we need only 461 examples
- If  $\epsilon = 0.1$  and  $\delta = 0.01$ , then for n = 10, we need 691 examples

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• To guarantee a probability of failure (i.e, error >  $\epsilon$ ) that is the set of that the set of the the set of the set of the number of examples we need is

$$m > \frac{n}{\epsilon} \left( \log(n) + \log\left(\frac{1}{\delta}\right) \right)$$

Poly in n, 1/
$$\delta$$
, 1/ $\epsilon$ 

- That is, if m has this property, then
  - With probability 1  $\delta$ , there will be no bad literals,
  - Or equivalently, with probability 1  $\delta$ , we will have err<sub>D</sub>(h) <  $\epsilon$

What we have here is a PAC guarantee

Our algorithm is **Probably Approximately Correct.** 

#### Vapnik-Chervonenkis (VC) Dimension

- A classic measure of complexity of infinite hypothesis classes based on this intuition.
- The VC dimension is a very classification-oriented notion of complexity
  - The idea is to look at a finite set of unlabeled examples
  - no matter how these points were labeled, would we be able to find a hypothesis that correctly classifies them
- The idea is that as you add more points, being able to represent an arbitrary labeling becomes harder and harder.

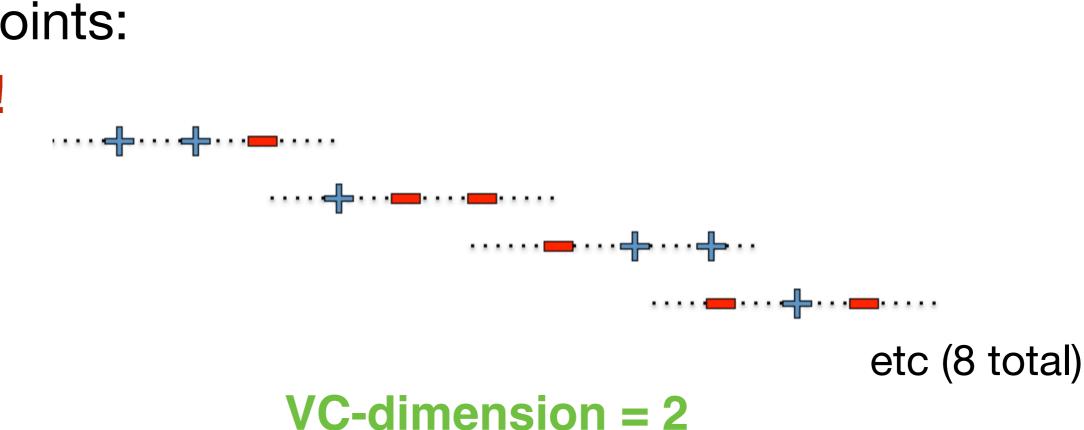
**Definitions 2.** For data drawn from some space X, the VC dimension of a hypothesis space H over X is the maximal K such that: there exists a set  $X \subseteq X$  of size |X| = K, such that for all binary labelings of X, there exists a function  $f \in H$  that matches this labeling.

#### How many points can a linear boundary classify exactly? (1-D)

• 2 points:

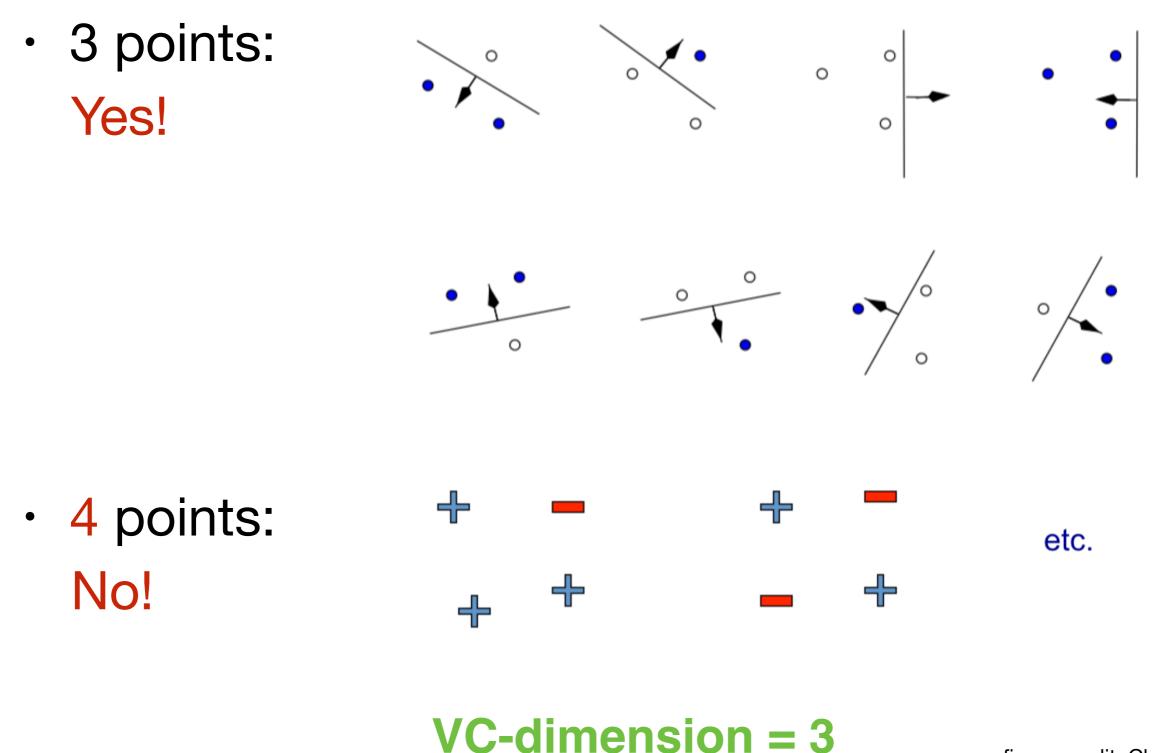
Yes!





.....

# How many points can a linear boundary classify exactly? (2-D)



#### Basic Probability Review

### Probability

- A is non-deterministic event
   Can think of A as a boolean-valued variable
- Examples
  - A = your next patient has cancer
  - A = Max Verstappen wins United States
     Grand Prix 2023



## Interpreting Probabilities

If I flip this coin, the probability that it will come up heads is 0.5

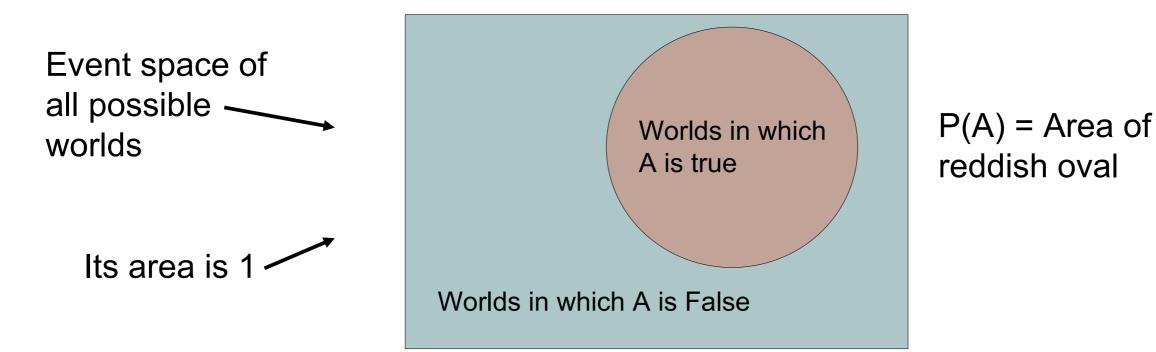
- Frequentist Interpretation: If we flip this coin many times, it will come up heads about half the time. Probabilities are the expected frequencies of events over repeated trials.
- Bayesian Interpretation: I believe that my next toss of this coin is equally likely to come up heads or tails. Probabilities quantify subjective beliefs about single events.
- Viewpoints play complementary roles in machine learning:
  - Bayesian view used to build models based on domain knowledge, and automatically derive learning algorithms
  - Frequentist view used to analyze worst case behavior of learning algorithms, in limit of large datasets
  - From either view, basic mathematics is the same!



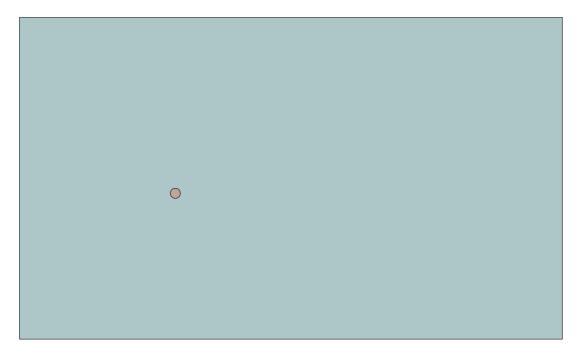
#### Axioms of Probability

- 0<= P(A) <= 1
- P(empty-set) = 0
- P(everything) = 1
- P(A or B) = P(A) + P(B) P(A and B)

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The area of A can t get any smaller than 0

And a zero area would mean no world could ever have A true

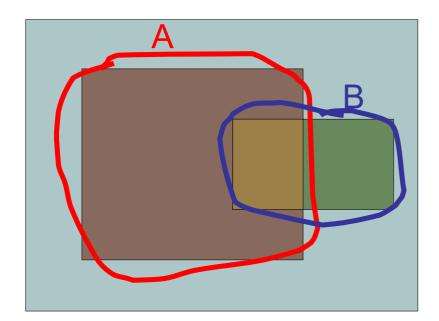
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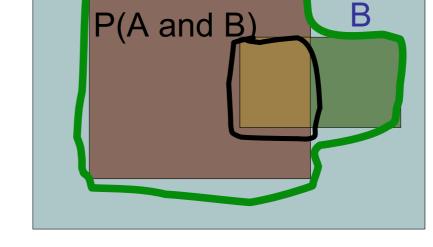


The area of A can t get any bigger than 1

And an area of 1 would mean all worlds will have A true

- 0<= P(A) <= 1
- P(empty-set) = 0
- P(everything) = 1
- P(A or B) = P(A) + P(B) P(A and B)





P(A or B)

Simple addition and subtraction

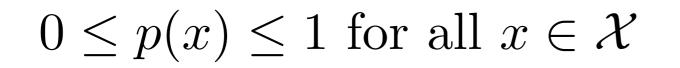
#### **Discrete Random Variables**

discrete random variable

sample space of possible outcomes,

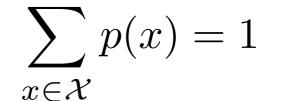
which may be finite or countably infinite

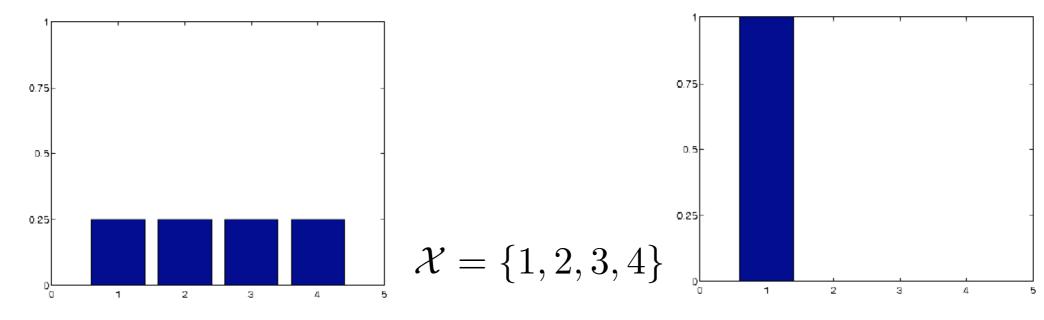
 $x \in \mathcal{X} \longrightarrow$  outcome of sample of discrete random variable



p(X = x)

p(x)



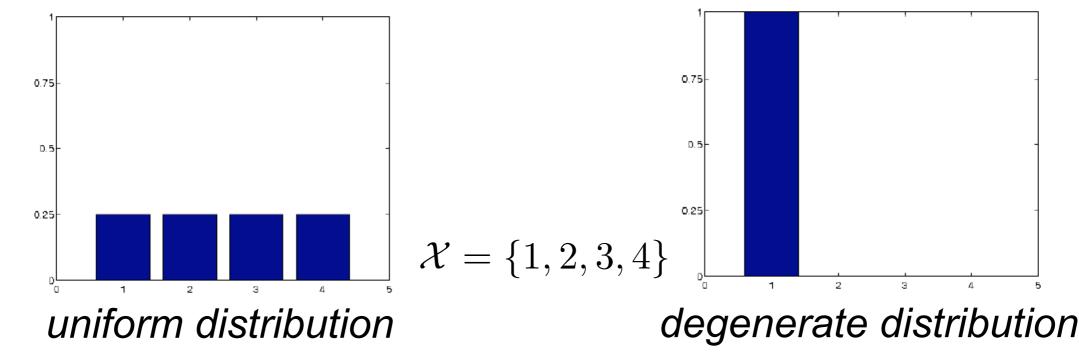


#### **Discrete Random Variables**

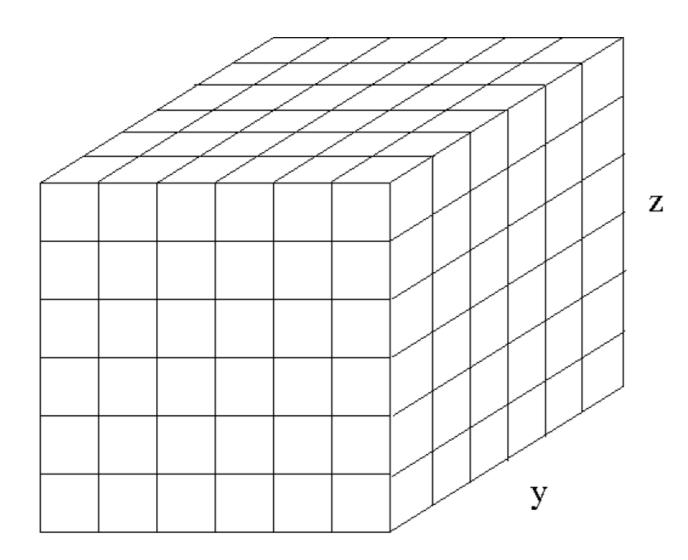
- discrete random variable
  - sample space of possible outcomes,
  - which may be finite or countably infinite
- $x \in \mathcal{X} \longrightarrow$  outcome of sample of discrete random variable  $p(X = x) \longrightarrow$  probability distribution (probability mass function)
  - $p(x) \longrightarrow$  shorthand used when no ambiguity

$$0 \le p(x) \le 1$$
 for all  $x \in \mathcal{X}$ 

$$\sum_{x \in \mathcal{X}} p(x) = 1$$



#### Joint Distribution

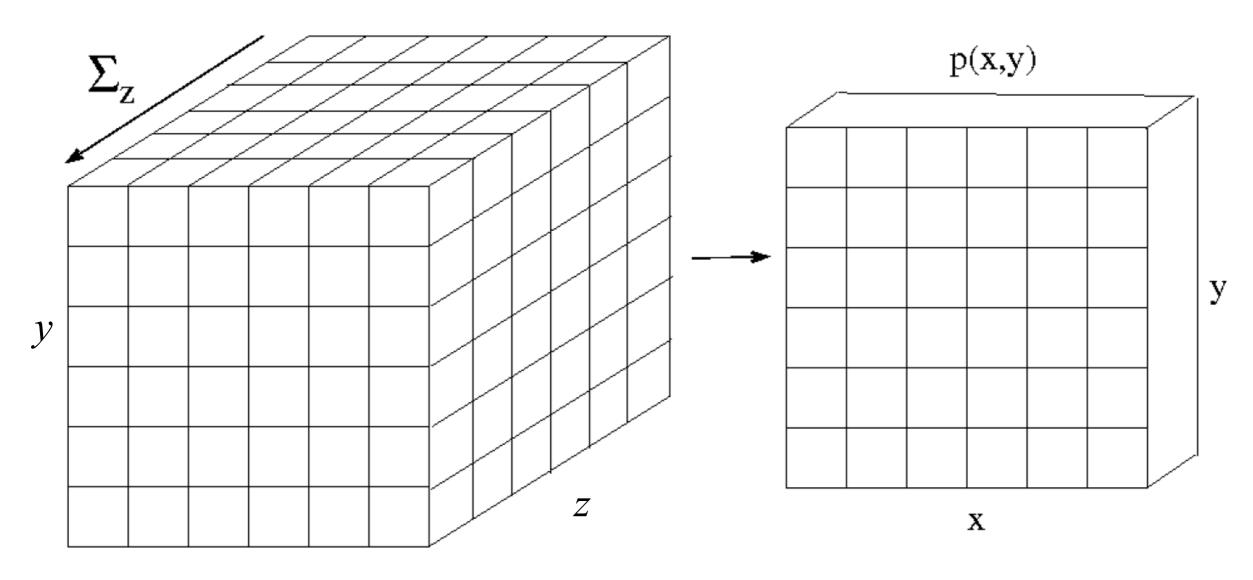


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#### Marginalization

- Marginalization
  - Events: P(A) = P(A and B) + P(A and not B)
  - Random variables  $P(X = x) = \sum_{y} P(X = x, Y = y)$

#### Marginal Distributions



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$$p(x,y) = \sum_{z \in \mathcal{Z}} p(x,y,z)$$

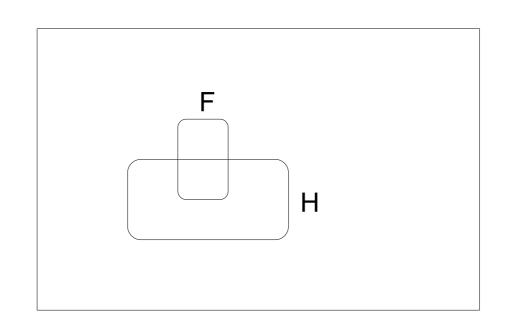
 $p(x) = \sum_{y \in \mathcal{Y}} p(x, y)$ 

#### **Conditional Probabilities**

- P(Y=y | X=x)
- What do you believe about Y=y, if I tell you X=x?
- P(Max Verstappen winning the 2024 United States Grand Prix)?
- What if I tell you:
  - He has won the Formula One World Champion title for 2021, 2022, and 2023.
  - He has won the United States Grand Prix 3/8 he has raced there.

#### **Conditional Probabilities**

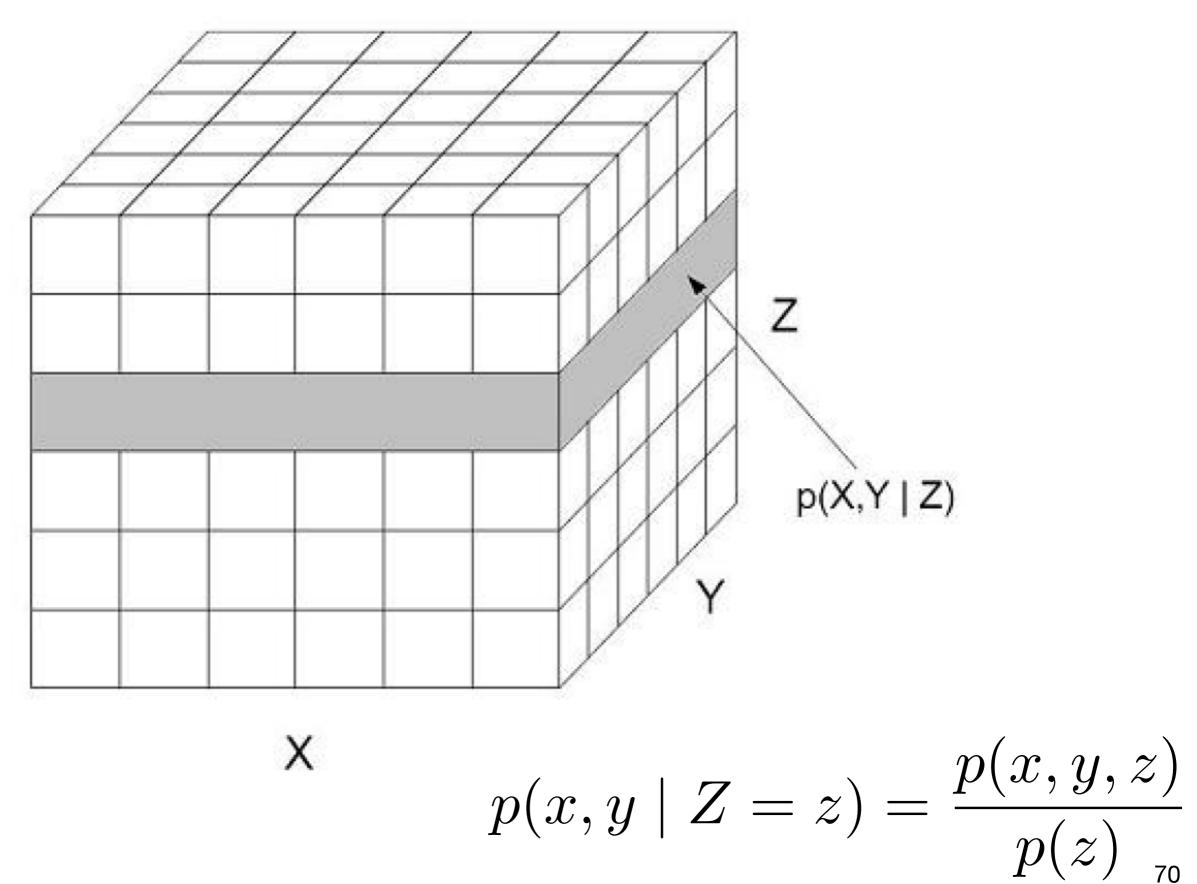
- P(A | B) = In worlds that where B is true, fraction where A is true
- Example
  - H: "Have a headache"
  - F: "Coming down with Flu"



P(H) = 1/10 P(F) = 1/40 P(H|F) = 1/2

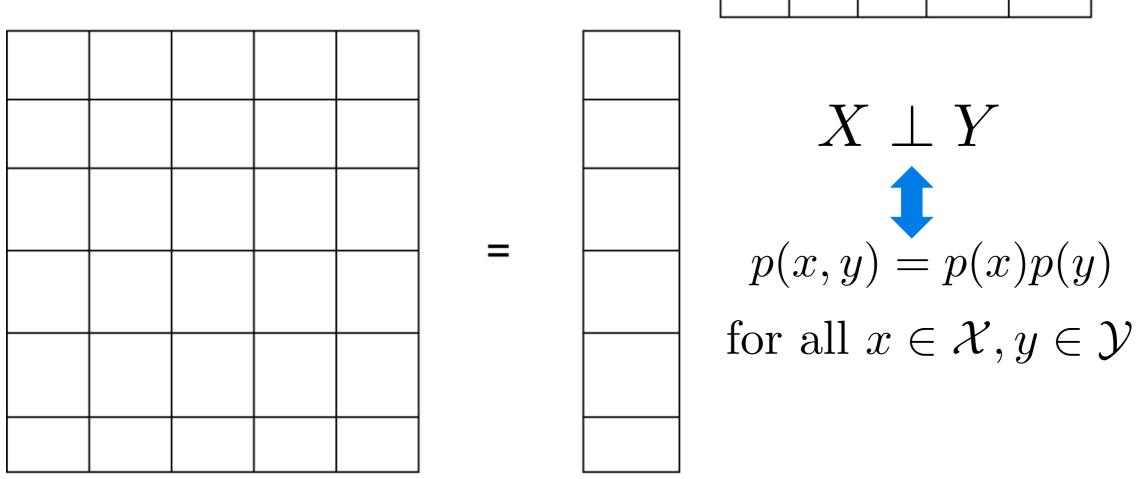
Headaches are rare and flu is rarer, but if you re coming down with flu there s a 50-50 chance you II have a headache.

#### **Conditional Distributions**



#### Independent Random Variables

P(x,y)



Equivalent conditions on conditional probabilities:

 $p(x \mid Y = y) = p(x) \text{ and } p(y) > 0 \text{ for all } y \in \mathcal{Y}$  $p(y \mid X = x) = p(y) \text{ and } p(x) > 0 \text{ for all } x \in \mathcal{X}$ 

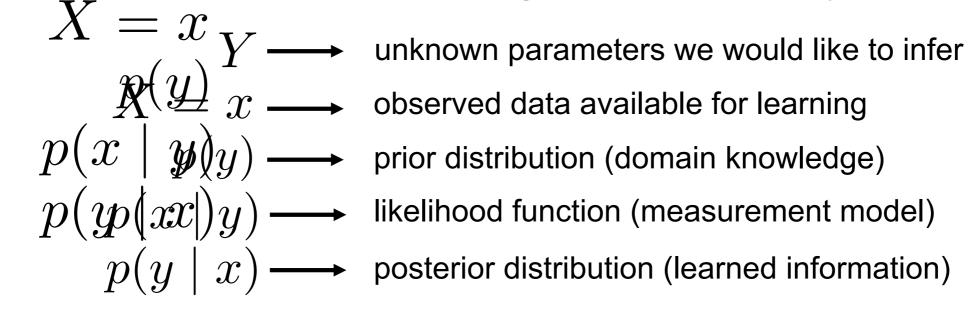
### Bayes Rule (Bayes Theorem)

$$p(x, y) = p(x)p(y \mid x) = p(y)p(x \mid y)$$

$$p(y \mid x) = \frac{p(x, y)}{p(x)} = \frac{p(x \mid y)p(y)}{\sum_{y' \in \mathcal{Y}} p(y')p(x \mid y')}$$

$$\propto p(x \mid y)p(y)$$

- A basic identity from the definition of conditional for the province of the
- Used in ways that have no thing to do with Bayesian statistics!
- Typical application to learning and data analysis:



#### Binary Random Variables

 Bernoulli Distribution: Single toss of a (possibly biased) coin

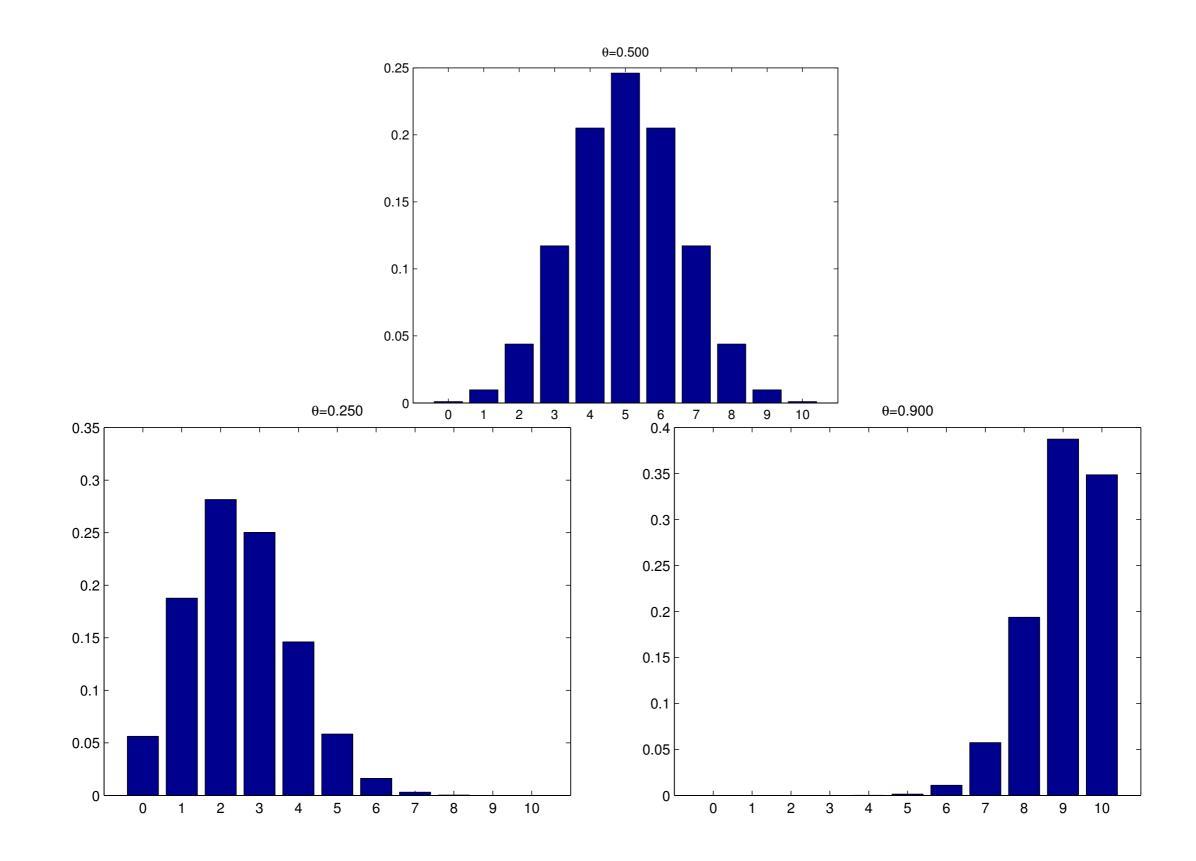
$$\begin{aligned} \mathcal{X} &= \{0, 1\} \\ \mathcal{X} &= \{0, 1\} \\ 0 &\leq \theta \leq 1 \\ \theta &\leq \theta \leq 1 \\ \theta &= \theta^{\delta(x, 1)}(1 - \theta)^{\delta(x, 0)} \\ \end{array}$$



• Binomial Distribution: Toss a single (possibly biased) coin *n* times, and report the number k of times it comes up  $\mathcal{K} = \{0, 1, 2, \dots, n\}$  $0 < \theta < 1$ 

$$\operatorname{Bin}(k \mid n, \theta) = \binom{n}{k} \theta^k (1 - \theta)^{n-k} \quad \binom{n}{k} = \frac{n!}{(n-k)!k!} \frac{\frac{n!}{k!k!}}{(n-k)!k!}$$

#### **Binomial Distributions**



#### Bean Machine (Sir Francis Galton)



http://en.wikipedia.org/wiki/ Bean machine

#### Categorical Random Variables

Multinoulli Distribution: Single roll of a (possibly biased) die

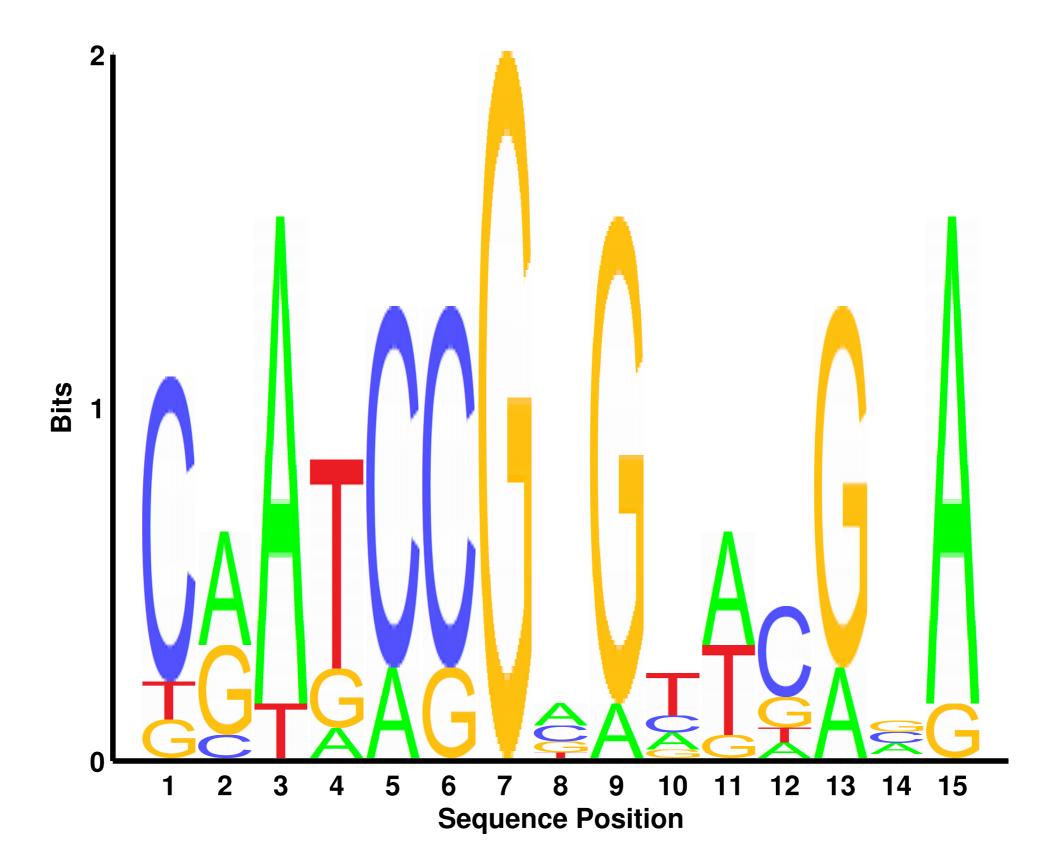
$$\mathcal{X} \overset{K_{K}}{=} \{ \{0,0\} \}_{j}^{K_{K}}, \sum_{\substack{k \in \mathbb{T}^{1} \\ k \in \mathbb{T}^{1}}} x y_{k} = \mathbf{1} \qquad \begin{array}{c} \text{binary vector} \\ \text{encoding} \\ \text{encoding} \\ \\ \mathcal{K} \\ \mathcal$$

• Multinomial Distribution: Roll a single (possibly biased) die *n* times, and report the number  $n_k$  of each possible outcome  $Mu(x \mid n, \theta) = \begin{pmatrix} n \\ n_1 \dots n_K \end{pmatrix} \prod_{k=1}^{K} \theta_k^{n_k} \qquad n_k = \sum_{i=1}^{n} x_{ik}$ 

#### Aligned DNA Sequences

cgatacggggtcgaa caatccgagatcgca caatccgtgttggga caatcggcatgcgg cgagccgcgtacgaa catacggagcacgaa taatccgggcatgta cgagccgagtacaga ccatccgcgtaagca ggatacgagatgaca

#### Multinomial Model of DNA



### Next Lecture: Maximum Likelihood Estimation (MLE)