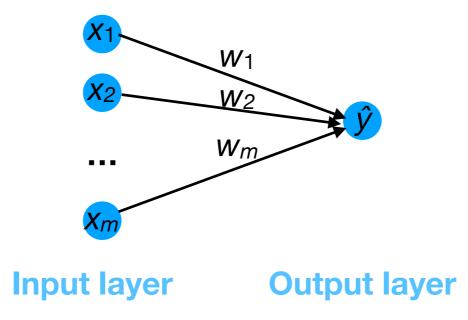
CS/DS 541: Class 3

Jacob Whitehill

Exercise

Gradient descent

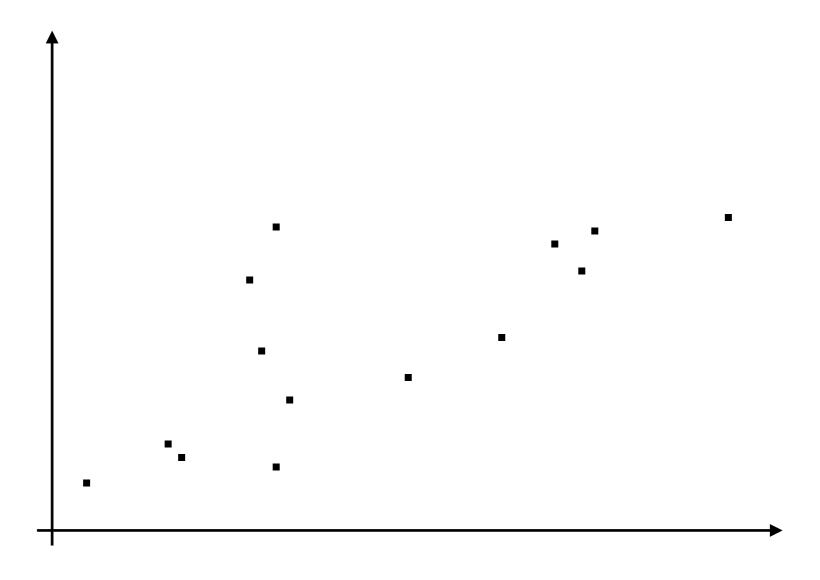
- For the 2-layer NN below, let m=2 and $\mathbf{w}^{(0)}=[1\ 0]^T$.
- Compute the updated weight vector w⁽¹⁾ after one iteration of gradient descent using MSE loss, a single training example (x,y)=([2, 3]^T, 4), and learning rate ε=0.1.



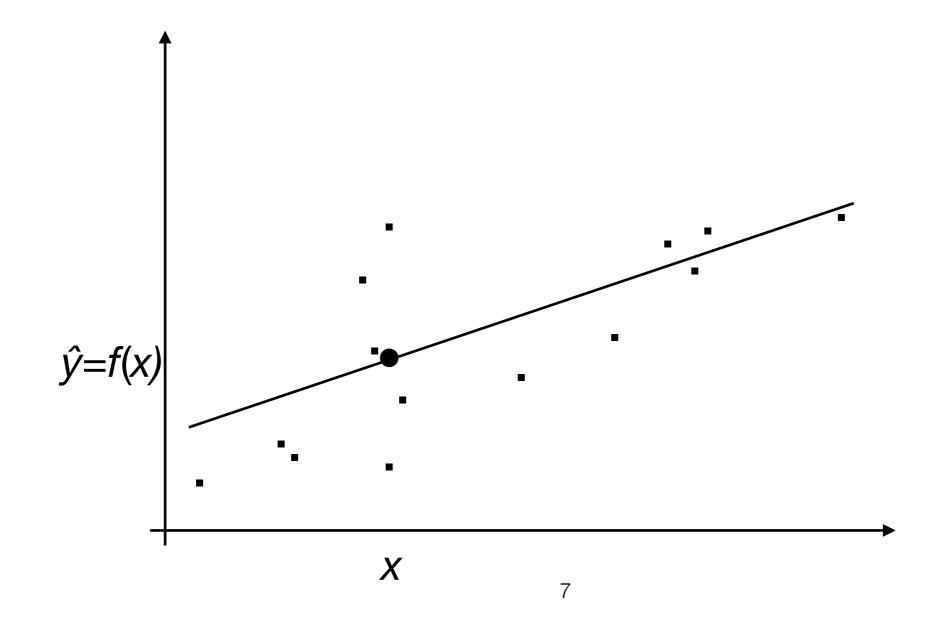
Solution

Probabilistic machine learning

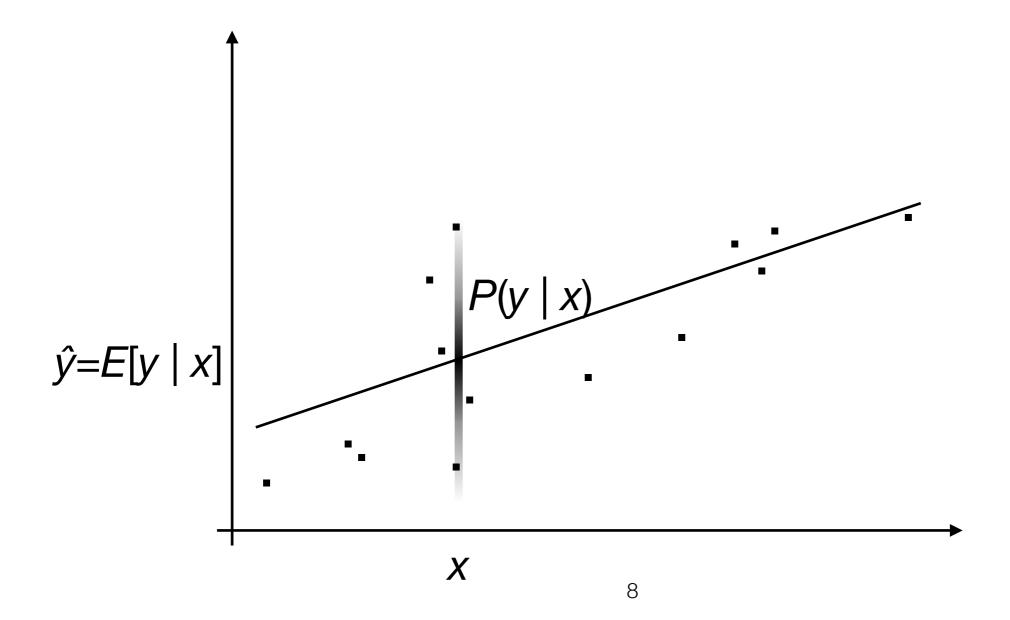
 Sometimes we may be very uncertain about our prediction of the target value y from the input x.



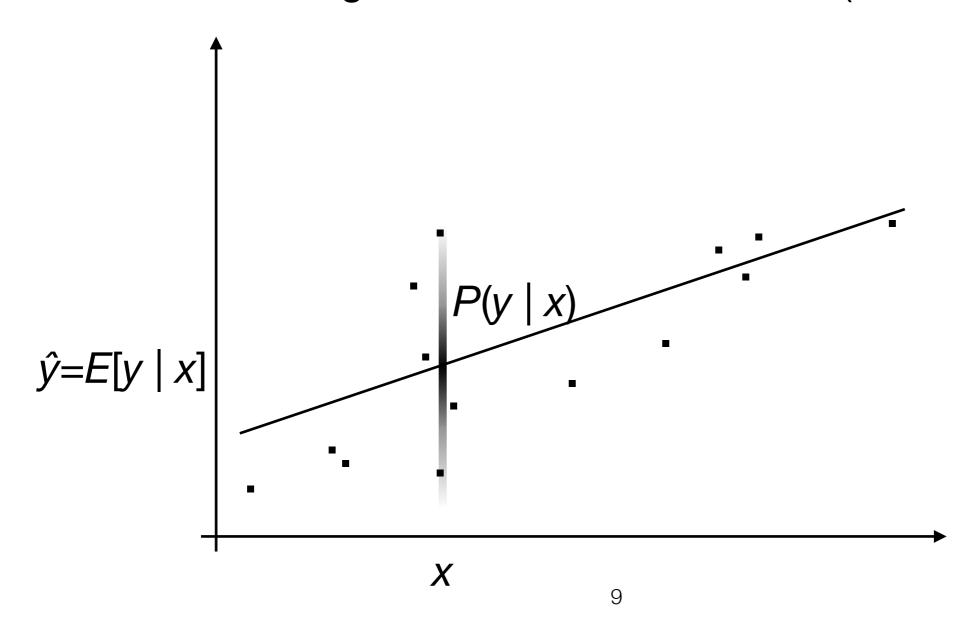
• Rather than just giving a point estimate $\hat{y}=f(x)...$



• ...we can estimate an entire probability distribution $P(y \mid x)$ that expresses our (conditional) uncertainty.



 Indeed, it turns out that the optimal parameters for a conditional Gaussian probability model are exactly the same as for linear regression with minimal MSE (i.e., 2-layer NN).



Probabilistic deep learning

- Neural networks can be used in various ways to make probabilistic predictions:
 - For regression, estimate both the expected value and the variance of the prediction.
 - Model a high-dimensional distribution using a probabilistic latent variable model (LVM) — akin to factor analysis but deeper.

Random variables

- A random variable* (RV) X (with sample space Ω) has a value we are unsure about, maybe because (a) it is decided by some random process or (b) it is hidden from us.
- Types of sample spaces Ω:
 - Finite, e.g., { 0, 1 }, { red, blue, green }.
 - Countable, e.g., $\mathbb{Z}_{>0}$
 - Uncountable, e.g., $\mathbb R$

^{*} This is a practical definition for the purposes of this dourse, not a formal definition based on measure theory.

Random variables

- RVs are typically written as capital letters, e.g., X, Y.
- Once the value of RV X has been "sampled", "selected", "instantiated", or "realized" (by a random number generator, generative process, God, etc.), it takes a specific value from the sample space.
- The values the RV can take are typically written as lowercase letters, e.g., x, y.

Random variables

- The probability that a random variable X takes a particular value is determined by a:
 - Probability mass function (PMF) for finite or countable sample spaces.
 - Probability density function (PDF) for uncountable sample spaces.

PMF

• Example 1 (finite):



- Let RV X be the outcome of rolling a 6-sided die.
- If X is fair, then: $P(X=i)=\frac{1}{6} \quad \forall i \in \{1,\dots,6\}$

PMF

• Example 2 (countable):



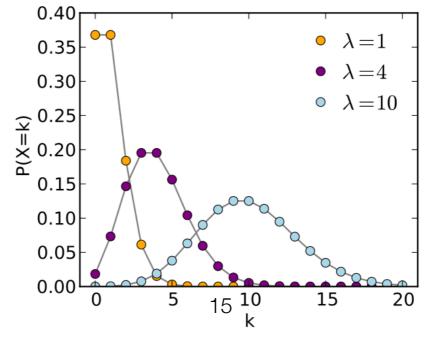
 Let RV X be the number of TCP/IP packets that arrive in 1 second.

• We can model the count of packets with a Poisson distribution: $\lambda k = -\lambda$

$$P(X = k) = \frac{\lambda^k e^{-\lambda}}{k!}$$

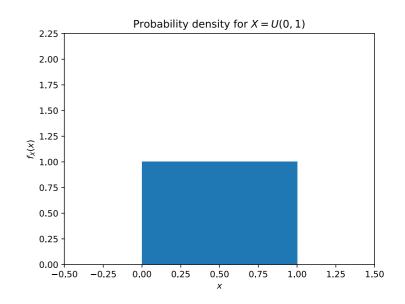
where parameter λ specifies the *rate* of the packet

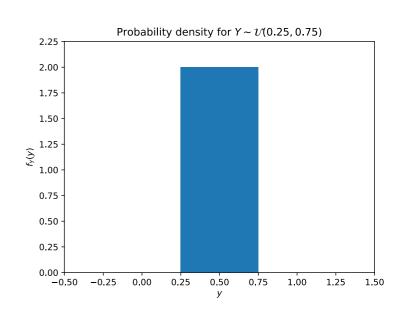
arrivals.



PDF

- Example 1:
 - Let X be a uniformly-distributed RV over Ω =[0,1].
 - Then $f_X(x) = 1 \quad \forall x \in \Omega$
- Example 2:
 - Let Y be a uniformly-distributed RV over Ω =[1/4, 3/4].
 - Then $f_Y(y)=2 \quad \forall y \in \Omega$ Note that the PDF can exceed 1.



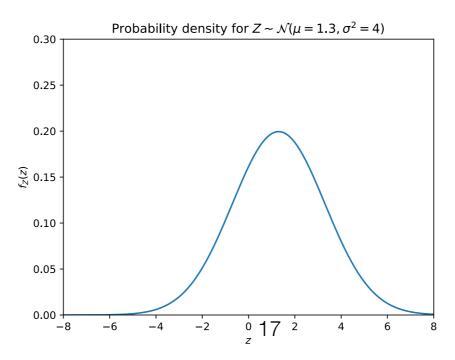


PDF

- Example 3:
 - Let Z be a **normally** (aka **Gaussian**) distributed RV with mean 1.5 and variance 4, i.e., $Z \sim \mathcal{N}(z; \mu = 1.5, \sigma^2 = 4)$.
 - Then

$$f_Z(z) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(z-\mu)^2}{2\sigma^2}\right)$$

where μ is the mean and σ^2 is the variance.



Probability distributions

- In this course, we will relax notation and use "probability distribution" to mean either the PDF or PMF of a RV (as appropriate).
- As a notational shortcut, we use P(x) to mean P(X=x) or $f_X(x)$.

- For multiple random variables X, Y, ..., we can construct the joint probability distribution P(x, y, ...) to mean the probability that $(X=x) \land (Y=y) \land$
- Note that P must still sum to 1 over all possible joint values (x, y, ...).

• Example in 2-D — crayons:



- Let X be the color (red, blue, green, white).
- Let Y be the intensity (low, medium, high).

	Red	Blue	Green	White
Low	0.1	0.05	0.025	0.2
Med	0.075	0.05	0.1	0
High	0.25	0.05	0.075	0.025

Exercise:



 What is the overall probability of obtaining a white crayon?

	Red	Blue	Green	White
Low	0.1	0.05	0.025	0.2
Med	0.075	0.05	0.1	0
High	0.25	0.05	0.075	0.025

• Example in 2-D — crayons:



• From the joint distribution we can compute the marginal distributions P(x) and P(y).

$$P(x) = \sum_{y} P(x, y) \quad P(y) = \sum_{x} P(x, y)$$

	Red	Blue	Green	White	P(y)
Low	0.1	0.05	0.025	0.2	0.375
Med	0.075	0.05	0.1	0	0.225
High	0.25	0.05	0.075	0.025	0.4
P(x)	0.425	0.15	0.2	0.225	

Law of total probability

- This is also called the law of total probability:
 - For any RVs X and Y:

$$P(x) = \sum_{y} P(x, y)$$

- In machine learning, we often use joint distributions of many variables that are part of a collection, e.g.:
 - Sequence $(W_1, W_2, ..., W_T)$ of words in a sentence.
 - W_t is the t^{th} RV in the sequence.
 - Grid $(I_{11}, ..., I_{1M}, ..., I_{N1}, ..., I_{NM})$ of the pixels in an $N \times M$ image.
 - I_{ij} is the RV corresponding to location (i, j).

- Sometimes the value of one RV is predictive of the value of another RV.
- Examples:
 - If I know a person's height H, then I have some information about their weight W.
 - If I know how much cholesterol C a person eats, then I have some information about their chance of coronary heart disease D.

 We can form a conditional probability distribution of RV X given the value of RV Y:

$$P(x \mid y)$$

- Examples:
 - Height given weight: P(h | w)
 - Heart disease given cholesterol: P(d | c)

 We can form a conditional probability distribution of RV X given the value of RV Y:

$$P(x \mid y)$$
 "given"

- Examples:
 - Height given weight: P(h | w)
 - Heart disease given cholesterol: $P(d \mid c)$

• More generally, we can form a conditional probability distribution of $X_1, ..., X_n$ given the values of $Y_1, ..., Y_m$:

$$P(x_1,\ldots,x_n\mid y_1,\ldots,y_m)$$

 A conditional probability distribution is related to the joint probability distribution as follows:

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

It follows that:

$$P(x \mid y, z)P(y \mid z) = P(x, y \mid z)$$

More generally:

$$P(x_1, \ldots, x_n \mid y_1, \ldots, y_m) P(y_1, \ldots, y_m) = P(x_1, \ldots, x_n, y_1, \ldots, y_m)$$

And also:

$$P(x_1, ..., x_n \mid y_1, ..., y_m, z_1, ..., z_p) P(y_1, ..., y_m \mid z_1, ..., z_p)$$

$$= P(x_1, ..., x_n, y_1, ..., y_m \mid z_1, ..., z_p)$$

 Note that the same joint probability can be factored in different ways, e.g.:

$$P(x, y, z) = P(x, y \mid z)P(z)$$
$$= P(x \mid y, z)P(y, z)$$

Exercises:

1.
$$P(a, b, c, d) = P(a, c) * ?$$

2.
$$P(w_1, w_2, w_3) = P(w_3 | w_1) * ? * ?$$

3.
$$P(x_1, x_2, x_3) = P(x_1) * ? * P(x_3 | x_1, x_2)$$

4.
$$P(x_1, ..., x_n) = P(x_1) * ? * ? * ... * ?$$

n-1 terms

Independence

- RVs X and Y are independent iff $P(x,y) = P(x)P(y) \ \forall x,y$, i.e., the joint distribution equals the product of the marginal distributions.
- Note that this implies that $P(x \mid y) = P(x)$ and $P(y \mid x) = P(y)$ since $P(x,y) = P(x \mid y)P(y) = P(y \mid x)P(x)$ by definition of conditional probability.

Conditional independence

 RVs X and Y are conditionally independent given RV Z iff:

$$P(x, y \mid z) = P(x \mid z)P(y \mid z) \ \forall x, y, z$$

Note that this implies:

$$P(x \mid y, z) = P(x \mid z)$$

In words: "If I already know the value of Z, then knowing Y tells me nothing further about X".

Conditional independence

• More generally: $X_1, ..., X_n$ and $Y_1, ..., Y_m$ are conditionally independent given $Z_1, ..., Z_p$ iff:

$$P(x_1, \dots, x_n, y_1, \dots, y_m \mid z_1, \dots, z_p)$$

= $P(x_1, \dots, x_n \mid z_1, \dots, z_p) P(y_1, \dots, y_m \mid z_1, \dots, z_p)$

- It is often useful to compute $P(x \mid y)$ in terms of $P(y \mid x)$.
 - For example, if X represents a student's skill level, and
 Y is their test score, it's often easier to compute P(y | x).
 But given a student's test score Y, we really want to
 know P(x | y).
- Bayes' rule:

$$P(x \mid y) = \frac{P(x,y)}{P(y)} = \frac{P(y|x)P(x)}{P(y)}$$

 We can also generalize Bayes' rule to cases where we always condition on a tertiary variable Z:

$$P(x \mid y, z) = \frac{P(y \mid x, z)P(x \mid z)}{P(y|z)}$$

- It is sometimes possible and more convenient to work with unnormalized probabilities.
- For instance, it might suffice to know that $[P(y^{(1)} | x), P(y^{(2)} | x), P(y^{(3)} | x)] \propto [3.5, 7, 0.04]$ rather than their exact (normalized) values.

 In this case, instead of needing to compute the denominator of...

$$P(x \mid y) = \frac{P(y \mid x)P(x)}{P(y)}$$

$$= \frac{P(y \mid x)P(x)}{\int P(x,y)dx}$$

$$= \frac{P(y \mid x)P(x)}{\int P(y \mid x)P(x)dx}$$

 In this case, instead of needing to compute the denominator of...

$$P(x \mid y) = \frac{P(y \mid x)P(x)}{P(y)}$$

$$= \frac{P(y \mid x)P(x)}{\int P(x,y)dx}$$

$$= \frac{P(y \mid x)P(x)}{\int P(y \mid x)P(x)dx}$$

Normalizing constant

In this case, instead of needing to compute the denominator of...

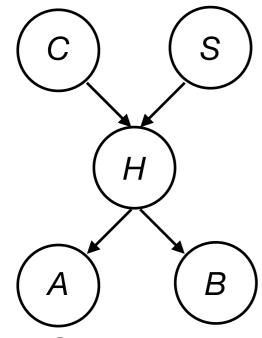
$$P(x \mid y) = \frac{P(y \mid x)P(x)}{P(y)}$$
 ... we might only need to compute the numerator:
$$= \frac{P(y \mid x)P(x)}{\int P(x,y)dx}$$

$$= \frac{P(y \mid x)P(x)}{\int P(y \mid x)P(x)dx}$$

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

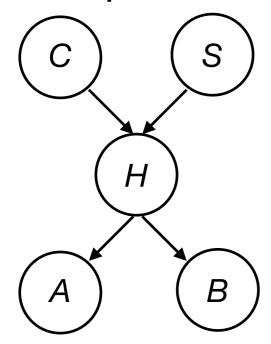
- To express the conditional independence relationships between multiple RVs, it is useful to represent their dependencies in a graph.
- A formal theory of probabilistic graphical models (Pearl 1998) has been devised.
 - Conditional independence can be determined via the principle of d-separation (beyond the scope of this course).

Example 1 — medical diagnosis:



- C: whether the patient eats caviar.
- S: the patient's **s**ex
- H: whether the patient has high cholesterol
- A: whether the patient will have a heart attack.
- B: whether the patient has shortness of breath.

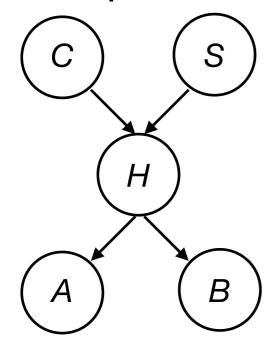
Example 1 — medical diagnosis:



This model implies that:

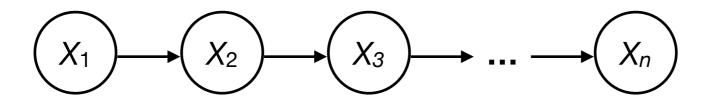
$$P(a, b \mid h, c, s) = P(a, b \mid h)$$
 and $P(c, s \mid h, a, b) = P(c, s \mid h)$

Example 1 — medical diagnosis:



• In words, "If I want to know the probability the patient will have a heart attack A, and I already know the patient has high cholesterol H, then the patient's sex and whether she/he eats caviar C is irrelevant."

Example 2 — Markov chain:

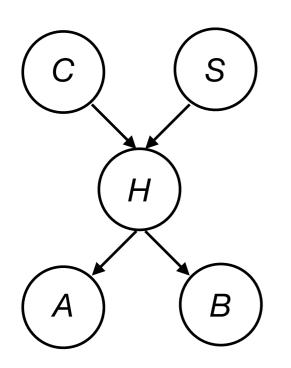


• This chain-like model of $X_1, ..., X_n$ implies that:

$$P(x_i \mid x_1, \dots, x_{i-1}) = P(x_i \mid x_{i-1})$$
 and $P(x_i \mid x_{i+1}, \dots, x_n) = P(x_i \mid x_{i+1})$

• In words, "If I want to know the value of X_i and I already know X_{i-1} , then the values of any 'earlier' X's are irrelevant."

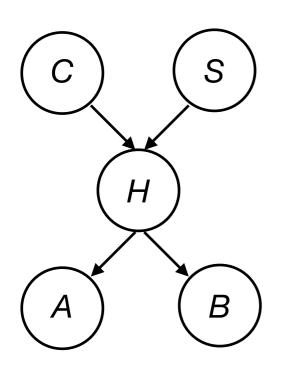
- Given a model with multiple RVs and how they are related to each other, we can **infer** the values of other RVs.
- For the medical diagnosis example, suppose we knew the conditional probability distributions:



P(H=0)		
	C=0	C=1
S=Ma		
S=Fe		

P(H=1)		
C=0 C=1		
S=Ma	0.3	0.6
S=Fe	0.2	0.25

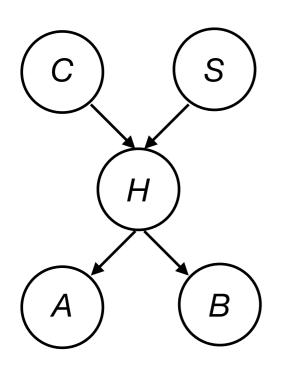
- Given a model with multiple RVs and how they are related to each other, we can infer the values of other RVs.
- For the medical diagnosis example, suppose we knew the conditional probability distributions:



P(H=0)		
	C=0	C=1
S=Ma	0.7	0.4
S=Fe	8.0	0.75

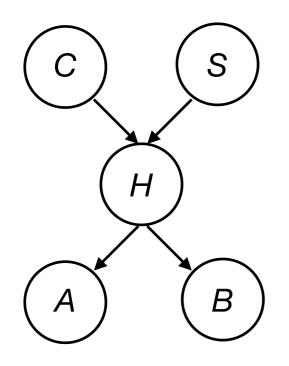
P(H=1)		
C=0 C=1		
S=Ma	0.3	0.6
S=Fe	0.2	0.25

- Given a model with multiple RVs and how they are related to each other, we can **infer** the values of other RVs.
- For the medical diagnosis example, suppose we knew the conditional probability distributions:



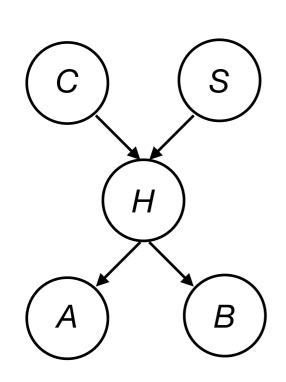
P(A	i=0)	P(A	=1)
H=0	0.75	H=0	0.25
<i>H</i> =1	0.2	H=1	0.8

- Given a model with multiple RVs and how they are related to each other, we can infer the values of other RVs.
- For the medical diagnosis example, suppose we knew the conditional probability distributions:



P(B	2=0)	P(B	=1)
<i>H</i> =0	0.9	H=0	0.
<i>H</i> =1	0.1	H=1	0

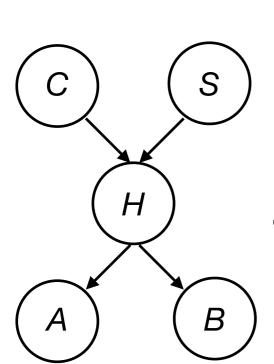
- Suppose we meet a male patient who eats caviar.
- What is the posterior probability that H=1, i.e., P(H=1 | C=1, S=Ma)? (Posterior means after observing C, S.)



P(H=0)		
C=0 C=1		C=1
S=Ma	0.7	0.4
S=Fe	8.0	0.75

P(H=1)		
C=0 C=1		C=1
S=Ma	0.3	0.6
S=Fe	0.2	0.25

- Suppose we meet a male patient who eats caviar.
- What is the posterior probability that H=1, i.e., P(H=1 | C=1, S=Ma)? (Posterior means after observing C, S.)

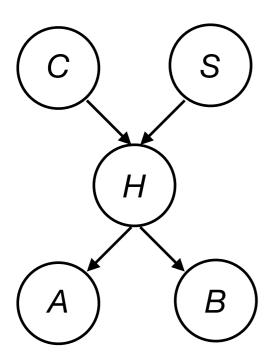


P(H=0)		
C=0 C=1		C=1
S=Ma	0.7	0.4
S=Fe	8.0	0.75

P(H=1)		
C=0 C=1		C=1
S=Ma	0.3	0.6

Just consult the conditional probability distribution.

What if we also know that the patient is short of breath?



P(H=0)		
C=0 C=1		C=1
S=Ma	0.7	0.4
S=Fe	8.0	0.75

P(B=0)		
<i>H</i> =0	0.9	
<i>H</i> =1	0.1	

P(H=1)			
	C=0 C=1		
S=Ma	0.3	0.6	
S=Fe	0.2	0.25	

P(B=1)		
H=0	0.1	
<i>H</i> =1	0.9	

$$P(b \mid h, c, s) = P(b \mid h)$$

$$P(H=1 \mid C=1, S={\rm Ma}, B=1) \\ = \frac{P(B=1 \mid H=1, C=1, S={\rm Ma})P(H=1 \mid C=1, S={\rm Ma})}{P(B=1 \mid C=1, S={\rm Ma})}$$
 Bayes' rule

	P(H=0)		ŀ	P(H=1)	
	C=0	C=1		C=0	C=1
S=Ma	0.7	0.4	S=Ma	0.3	0.6
S=Fe	0.8	0.75	S=Fe	0.2	0.25

$$P(B=0)$$
 $P(B=1)$ $H=0$ 0.1 $H=1$ 0.9 $H=1$ 0.9

$$P(b | h, c, s) = P(b | h)$$

$$P(H=1 \mid C=1, S={\rm Ma}, B=1)$$

$$= \frac{P(B=1 \mid H=1, C=1, S={\rm Ma})P(H=1 \mid C=1, S={\rm Ma})}{P(B=1 \mid C=1, S={\rm Ma})}$$
 Bayes' rule
$$= \frac{P(B=1 \mid H=1)P(H=1 \mid C=1, S={\rm Ma})}{P(B=1 \mid C=1, S={\rm Ma})}$$
 Conditional independence

	P(H=0)			P(H=1)	
	C=0	C=1		C=0	C=1
S=Ma	0.7	0.4	S=Ma	0.3	0.6
S=Fe	0.8	0.75	S=Fe	0.2	0.25

$$P(B=0)$$
 $P(B=1)$ $H=0$ 0.1 $H=1$ 0.9 $H=1$ 0.9

$$P(b \mid h, c, s) = P(b \mid h)$$

$$P(H=1 \mid C=1, S=\text{Ma}, B=1)$$

$$= \frac{P(B=1 \mid H=1, C=1, S=\text{Ma})P(H=1 \mid C=1, S=\text{Ma})}{P(B=1 \mid C=1, S=\text{Ma})}$$
Bayes' rule
$$= \frac{P(B=1 \mid H=1)P(H=1 \mid C=1, S=\text{Ma})}{P(B=1 \mid C=1, S=\text{Ma})}$$
Conditional independence
$$P(B=1 \mid C=1, S=\text{Ma})$$

$$0.9*0.6$$

 $= \frac{1}{\sum_{h=0}^{1} P(B=1, H=h \mid C=1, S=Ma)}$

Law of total probability

	P(H=0)		ı	P(H=1)	
	C=0	C=1		C=0	C=1
S=Ma	0.7	0.4	S=Ma	0.3	0.6
S=Fe	8.0	0.75	S=Fe	0.2	0.25

P(B	3=0)	P(B	=1)
H=0	0.9	H=0	0.1
<i>H</i> =1	0.1	H=1	0.9

$$P(b \mid h, c, s) = P(b \mid h)$$

$$P(H = 1 \mid C = 1, S = \text{Ma}, B = 1)$$

$$= \frac{P(B = 1 \mid H = 1, C = 1, S = \text{Ma})P(H = 1 \mid C = 1, S = \text{Ma})}{P(B = 1 \mid C = 1, S = \text{Ma})}$$
Bayes' rule
$$= \frac{P(B = 1 \mid H = 1)P(H = 1 \mid C = 1, S = \text{Ma})}{P(B = 1 \mid C = 1, S = \text{Ma})}$$
Conditional independence
$$= \frac{0.9 * 0.6}{\sum_{h=0}^{1} P(B = 1, H = h \mid C = 1, S = \text{Ma})}$$
Law of total probability
$$= \frac{0.54}{\sum_{h=0}^{1} P(B = 1 \mid H = h, C = 1, S = \text{Ma})P(H = h \mid C = 1, S = \text{Ma})}$$
Def. of cond. prob.

$$P(H=1 \mid C=1, S=\text{Ma}, B=1)$$

$$= \frac{P(B=1 \mid H=1, C=1, S=\text{Ma})P(H=1 \mid C=1, S=\text{Ma})}{P(B=1 \mid C=1, S=\text{Ma})} \quad \text{Bayes' rule}$$

$$= \frac{P(B=1 \mid H=1)P(H=1 \mid C=1, S=\text{Ma})}{P(B=1 \mid C=1, S=\text{Ma})} \quad \text{Conditional independence}$$

$$= \frac{0.9*0.6}{\sum_{h=0}^{1} P(B=1, H=h \mid C=1, S=\text{Ma})} \quad \text{Law of total probability}$$

$$= \frac{0.54}{\sum_{h=0}^{1} P(B=1 \mid H=h, C=1, S=\text{Ma})P(H=h \mid C=1, S=\text{Ma})} \quad \text{Ond. prob.}$$

$$= \frac{0.54}{\sum_{h=0}^{1} P(B=1 \mid H=h)P(H=h \mid C=1, S=\text{Ma})} \quad \text{Conditional independence}$$

$$P(H=1 \mid C=1, S=\text{Ma}, B=1) \\ = \frac{P(B=1 \mid H=1, C=1, S=\text{Ma})P(H=1 \mid C=1, S=\text{Ma})}{P(B=1 \mid C=1, S=\text{Ma})} \quad \text{Bayes' rule} \\ = \frac{P(B=1 \mid H=1)P(H=1 \mid C=1, S=\text{Ma})}{P(B=1 \mid C=1, S=\text{Ma})} \quad \text{Conditional independence} \\ = \frac{0.9*0.6}{\sum_{h=0}^{1} P(B=1, H=h \mid C=1, S=\text{Ma})} \quad \text{Law of total probability} \\ = \frac{0.54}{\sum_{h=0}^{1} P(B=1 \mid H=h, C=1, S=\text{Ma})P(H=h \mid C=1, S=\text{Ma})} \quad \text{Conditional independence} \\ = \frac{0.54}{\sum_{h=0}^{1} P(B=1 \mid H=h)P(H=h \mid C=1, S=\text{Ma})} \quad \text{Conditional independence} \\ = \frac{0.54}{0.1*0.4+0.9*0.6}$$

$$\begin{split} &P(H=1\mid C=1,S=\text{Ma},B=1)\\ &=\frac{P(B=1\mid H=1,C=1,S=\text{Ma})P(H=1\mid C=1,S=\text{Ma})}{P(B=1\mid C=1,S=\text{Ma})} \quad \text{Bayes' rule}\\ &=\frac{P(B=1\mid H=1)P(H=1\mid C=1,S=\text{Ma})}{P(B=1\mid C=1,S=\text{Ma})} \quad \text{Conditional independence}\\ &=\frac{P(B=1\mid C=1,S=\text{Ma})}{P(B=1\mid H=h\mid C=1,S=\text{Ma})} \quad \text{Law of total probability}\\ &=\frac{O.54}{\sum_{h=0}^{1}P(B=1\mid H=h,C=1,S=\text{Ma})P(H=h\mid C=1,S=\text{Ma})} \quad \text{Def. of cond. prob.}\\ &=\frac{O.54}{\sum_{h=0}^{1}P(B=1\mid H=h)P(H=h\mid C=1,S=\text{Ma})} \quad \text{Conditional independence}\\ &=\frac{O.54}{0.1*0.4+0.9*0.6}\\ &=\frac{O.54}{0.04+0.54} \quad \text{Conditional independence} \end{split}$$

 ≈ 0.93

$$P(H=h\mid C=1,S=\mathrm{Ma},B=1)$$

$$\propto P(B=1\mid H=h,C=1,S=\mathrm{Ma})P(H=h\mid C=1,S=\mathrm{Ma})$$
 Bayes' rule
$$=P(B=1\mid H=h)P(H=h\mid C=1,S=\mathrm{Ma})$$
 Conditional independence

$$P(H = 1 \mid C = 1, S = Ma, B = 1)$$

 $\propto P(B = 1 \mid H = 1)P(H = 1 \mid C = 1, S = Ma)$
 $= 0.9 * 0.6$

$$P(H = 1 \mid C = 1, S = Ma, B = 1)$$

 $\propto P(B = 1 \mid H = 1)P(H = 1 \mid C = 1, S = Ma)$
 $= 0.9 * 0.6$
 $P(H = 0 \mid C = 1, S = Ma, B = 1)$
 $\propto P(B = 1 \mid H = 0)P(H = 0 \mid C = 1, S = Ma)$
 $= 0.1 * 0.4$

$$P(H=1 \mid C=1, S=\mathrm{Ma}, B=1)$$

$$\propto P(B=1 \mid H=1)P(H=1 \mid C=1, S=\mathrm{Ma})$$

$$= 0.9*0.6$$

$$P(H=0 \mid C=1, S=\mathrm{Ma}, B=1)$$

$$\propto P(B=1 \mid H=0)P(H=0 \mid C=1, S=\mathrm{Ma})$$

$$= 0.1*0.4$$

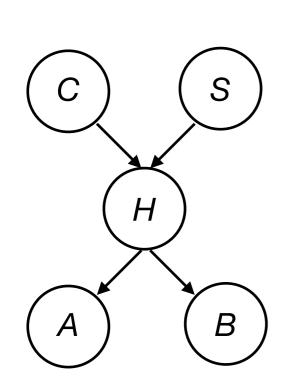
$$\Longrightarrow$$

$$P(H=1 \mid C=1, S=\mathrm{Ma}, B=1) = \frac{0.9*0.6}{0.9*0.6+0.1*0.4} = 0.93$$

$$P(H=0 \mid C=1, S=\mathrm{Ma}, B=1) = \frac{0.1*0.4}{0.9*0.6+0.1*0.4} = 0.07$$
 since the probabilities must sum to 1.

Parameters in probability distributions

- Most probabilistic models have parameters we want to estimate.
- For example, the conditional probabilities for medical diagnosis are all parameters that must be learned.



P(H=0)		
C=0 C=1		
S=Ma	0.7	0.4
S=Fe	8.0	0.75

<i>i</i> (<i>i i</i> – 1)			
	C=0	C=1	
S=Ma	0.3	0.6	
S=Fe	0.2	0.25	

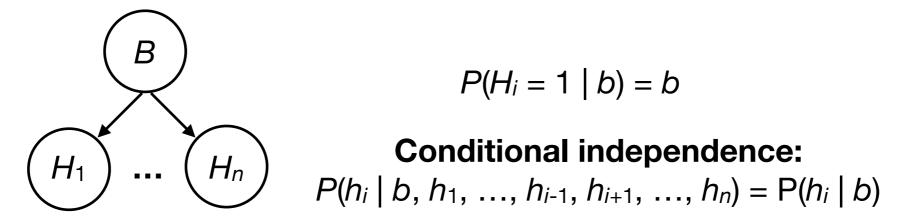
P(H=1)

P(B	(=0)
<i>H</i> =0	0.9
<i>H</i> =1	0.1

P(B=1)		
<i>H</i> =0	0.1	
<i>H</i> =1	0.9	

Parameters in probability distributions

- Most probabilistic models have parameters we want to estimate.
- As another example, we might want to estimate the bias B of a coin after observing n coin flips $H_1, ..., H_n$:



• What is a principled approach to estimating *B*?

- Maximum likelihood estimation (MLE):
 - The value of a latent variable is estimated as the one that makes the observed data as likely (probable) as possible.
- The **likelihood** of $H_1, ..., H_n$ given B is:

$$P(h_1, \dots, h_n \mid b) = P(h_1 \mid b) \prod_{i=2}^n P(h_i \mid b, h_1, \dots, h_{i-1})$$

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$$=\prod_{i=2}^n P(h_i\mid b)$$

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$$P(h_i \mid b) = b^{h_i} (1 - b)^{1 - h_i}$$

= b if $h_i = 1$ or
 $(1 - b)$ if $h_i = 0$

The exponent "chooses" the correct probability for $H_i=1$ or $H_i=0$.

- We seek to maximize the probability of $h_1, ..., h_n$ by optimizing b.
- It's often easier instead to optimize the log-likelihood.

$$\arg\max_{b} P(h_1, \dots, h_n \mid b) = \arg\max_{b} \log P(h_1, \dots, h_n \mid b)^*$$

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^{*} assuming the probability is never exactly 0.

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 due to conditional independence

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$$= \sum_{i=1}^n \log b^{h_i} (1-b)^{1-h_i}$$

$$= \sum_{i=1}^n h_i \log b + (1-h_i) \log(1-b)$$

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$$= \sum_{i=1}^n h_i \log b + (1-h_i) \log(1-b)$$

$$= n_1 \log b + (n-n_1) \log(1-b)$$

 We can now differentiate w.r.t. b, set to 0, and solve to obtain the MLE of B:

$$\nabla_b \left[n_1 \log b + (n - n_1) \log(1 - b) \right] = \frac{n_1}{b} - \frac{(n - n_1)}{1 - b}$$

$$b = ?$$

 We can now differentiate w.r.t. b, set to 0, and solve to obtain the MLE of B:

$$\nabla_b \left[n_1 \log b + (n - n_1) \log(1 - b) \right] = \frac{n_1}{b} - \frac{(n - n_1)}{1 - b}$$

$$(1 - b)n_1 - b(n - n_1) = 0$$

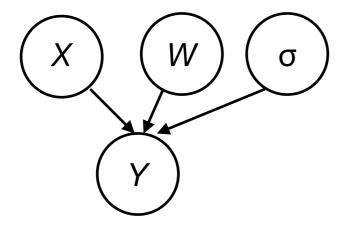
$$n_1 - bn_1 - bn + bn_1 = 0$$

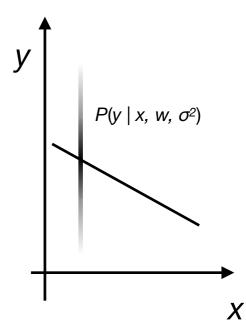
$$n_1 = bn$$

$$b = \frac{n_1}{n}$$

The MLE for B is the fraction of coin flips that are heads.

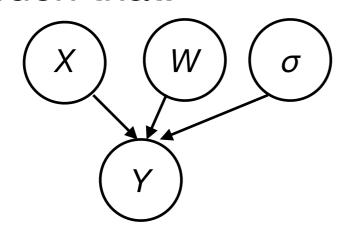
Let's consider a different model that contains real-valued
 RVs (not just from a finite sample space).

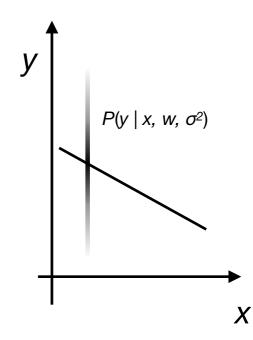




- X is some feature vector (e.g., face image).
- Y is some outcome variable (e.g., age).
- W is a vector of weights that characterize how Y is related to X.
- σ expresses how uncertain we are about Y after seeing X.

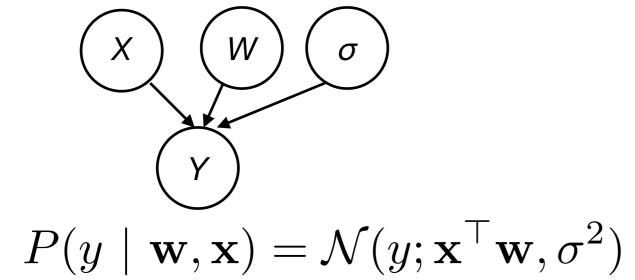
 Suppose we model the relationship between X, W, σ, and Y such that:

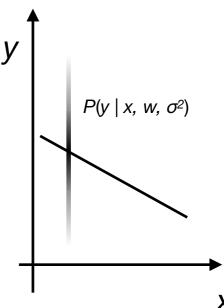




- Y is a normal/Gaussian random variable.
- The expected value of Y is x^Tw .
- The variance of Y is constant (σ^2) for all possible x.

 Suppose we model the relationship between X, W, σ, and Y such that:





• If we collect a dataset $\mathcal{D} = \{(\mathbf{x}^{(i)}, y^{(i)})\}_{i=1}^n$, what is the MLE for W and σ ?

$$P(y \mid \mathbf{x}, \mathbf{w}, \sigma^2) = \mathcal{N}(y; \mathbf{x}^\top \mathbf{w}, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(y - \mathbf{x}^\top \mathbf{w})^2}{2\sigma^2}\right)$$

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$$P(\mathcal{D} \mid \mathbf{w}, \sigma^2) = \prod^n P(y^{(i)} \mid \mathbf{x}^{(i)}, \mathbf{w}, \sigma^2)$$
 Conditional independence

$$P(y \mid \mathbf{x}, \mathbf{w}, \sigma^2) = \mathcal{N}(y; \mathbf{x}^\top \mathbf{w}, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(y - \mathbf{x}^\top \mathbf{w})^2}{2\sigma^2}\right)$$

$$P(\mathcal{D} \mid \mathbf{w}, \sigma^2) = \prod_{i=1}^n P(y^{(i)} \mid \mathbf{x}^{(i)}, \mathbf{w}, \sigma^2)$$
 Conditional independence

$$\log P(\mathcal{D} \mid \mathbf{w}, \sigma^2) = \log \prod_{i=1}^n P(y^{(i)} \mid \mathbf{x}^{(i)}, \mathbf{w}, \sigma^2)$$

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$$= \sum_{i=1}^{n} \log P(y^{(i)} \mid \mathbf{x}^{(i)}, \mathbf{w}, \sigma^2)$$

MLE for w:

$$\mathbf{w} = \left(\sum_{i=1}^{n} \mathbf{x}^{(i)} \mathbf{x}^{(i)}^{\top}\right)^{-1} \left(\sum_{i=1}^{n} \mathbf{x}^{(i)} y^{(i)}\right)$$

This is the same solution as for linear regression, but derived as the MLE of a probabilistic model (instead of the minimum MSE).

• MLE for σ^2 :

$$\sigma^2 = \frac{1}{n} \sum_{i=1}^{n} (\mathbf{x}^{(i)}^{\mathsf{T}} \mathbf{w} - y^{(i)})^2$$

This is the sum of squared residuals of the predictions w.r.t. ground-truth.

L₂ Regularization

Regularization

- The larger the coefficients (weights) w are allowed to be, the more the neural network can overfit.
- If we "encourage" the weights to be small, we can reduce overfitting.
- This is a form of regularization any practice designed to improve the machine's ability to generalize to new data.

Regularization

- One of the simplest and oldest regularization techniques is to penalize large weights in the cost function.
- The "unregularized" f_{MSE} is:

$$f_{\text{MSE}}(\mathbf{w}) = \frac{1}{2n} \sum_{i=1}^{n} (y^{(i)} - \hat{y}^{(i)})^2$$

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The L₂-regularized f_{MSE} becomes:

$$f_{\text{MSE}}(\mathbf{w}) = \frac{1}{2n} \sum_{i=1}^{n} (y^{(i)} - \hat{y}^{(i)})^2 + \frac{\alpha}{2n} \mathbf{w}^{\top} \mathbf{w}$$

Hyperparameter tuning

Hyperparameter tuning

- The values we optimize when training a machine learning model — e.g., w and b for linear regression — are the parameters of the model.
- There are also values related to the training process itself e.g., learning rate ε , batch size \tilde{n} , regularization strength α which are the **hyperparameters** of training.

Hyperparameter tuning

- Both the parameters and hyperparameters can have a huge impact on model performance on test data.
- When estimating the performance of a trained model, it is important to tune both kinds of parameters in a principled way:
 - Training/validation/testing sets
 - Double cross-validation

- In an application domain with a large dataset (e.g., 100K examples), it is common to partition it into three subsets:
 - Training (typically 70-80%): optimization of parameters
 - Validation (typically 5-10%): tuning of hyperparameters
 - Testing (typically 5-10%): evaluation of the final model
- For comparison with other researchers' methods, this partition should be fixed.

- Hyperparameter tuning works as follows:
 - 1.For each hyperparameter configuration h:
 - Train the parameters on the training set using h.
 - Evaluate the model on the validation set.
 - If performance is better than what we got with the best h so far (h*), then save h as h*.
 - 2. Train a model with h^* , and evaluate its accuracy A on the **testing** set. (You can train either on training data, or on training+validation data).

To what machine does the reported accuracy A correspond?

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 - 1.For each hyperparameter configuration h:
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- When working with smaller datasets, cross-validation is commonly used so that we can use all data for training.
- Suppose we already know the best hyperparameters h*.
- We partition the data into k folds of equal sizes.
- Over k iterations, we train on (k-1) folds and test on the remaining fold.
- We then compute the average accuracy over the k testing folds.

To what machine does the reported accuracy A correspond?

```
    # D=dataset, k=# folds, h=hyperparameter configuration.
    CrossValidation (D, k, h):

            Partition D into k folds F<sub>1</sub>, ..., F<sub>k</sub>
             For i = 1, ..., k:
                test = F<sub>i</sub>
               train = D \ F<sub>i</sub>
                  Train the model on train using h
                 acc[i] = Evaluate NN on test
                  A = Avg[acc]
                 return A
                  Tresult of the property of the
```

To what machine does the reported accuracy A correspond?

None of them!

Training/validation/testing sets

- Cross-validation does not measure the accuracy of any single machine.
- Instead, cross-validation gives the expected accuracy of a classifier that is trained on (k-1)/k of the data.

Training/validation/testing sets

- Cross-validation does not measure the accuracy of any single machine.
- Instead, cross-validation gives the expected accuracy of a classifier that is trained on (k-1)/k of the data.
- However, we can train another model M using h* on the entire dataset, and then report A as its accuracy.
- Since *M* is trained on more data than any of the cross-validation models, its *expected* accuracy should be >= *A*.

Cross-validation

- But how do we find the best hyperparameters h* for each fold?
- The typical approach is to use double cross-validation, i.e.:
 - For each of the k "outer" folds, run cross-validation in an "inner" loop to determine the best hyperparameter configuration h* for the kth fold.

Double cross-validation

 # D=dataset, k=# folds, H=set of hyperparameter configurations. DoubleCrossValidation (D, k, H): Partition *D* into *k* folds $F_1, ..., F_k$ For i = 1, ..., k: $test = F_i$ $train = D \setminus F_i$ $A^* = -\infty$ For h in H: A = CrossValidation(train, k, h)if $A > A^*$: $A^* = A$ $h^* = h$ Train the model on *train* using *h** accs[i] = Evaluate the model on test A = Avg[accs]return A

For your reference...

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 Partition D into k folds F₁, ..., F_k
 For i = 1, ..., k:

 $test = F_i$ $train = D \setminus F_i$

 $A^* = -\infty$

For h in H:

A = CrossValidation(train, k, h)

if $A > A^*$:

 $A^* = A$

 $h^* = h$

Train the model on *train* using h^* accs[i] = Evaluate the model on *test*

A = Avg[accs] return A

None of them!

For your reference...

CrossValidation (D, k, h):

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Training/validation/testing sets

- In contrast to (single) cross-validation, it's not obvious how to train a model M with accuracy >= A.
- One strategy: return an ensemble model whose output is the average of the k models' predictions...but this is rarely done.

Subject independence

 In many machine learning settings, the data are not completely independent from each other — they are linked in some way.

Example:

 Predict multiple grades for each student based on their Canvas clickstream features (# logins, # forum posts, etc.).

Subject independence

- We could partition the data into folds in different ways:
 - We could randomize across all the data.
 - However, if grades are correlated within each student, then one (or more) training folds can leak information about the testing fold.

	Quiz 1	Quiz 2	Quiz 3
Student 1	45	48	42
Student 2	96	93	93
Student 3	86	86	87
Student 4	10	30	50

Subject independence

- We could partition the data into folds in different ways:
 - Alternatively, we can stratify across students, i.e., no student appears in more than 1 fold.
 - With this partition, the cross-validation accuracy estimates the model's performance on a subject not used for training.

	Quiz 1	Quiz 2	Quiz 3
Student 1	45	48	42
Student 2	96	93	93
Student 3	86	86	87
Student 4	10	30	50

- Gradient descent is guaranteed to converge to a local minimum (eventually) if the learning rate is small enough relative to the steepness of f.
- A function $f: \mathbb{R}^m \to \mathbb{R}$ is Lipschitz-continuous if:

$$\exists L : \forall \mathbf{x}, \mathbf{y} \in \mathbb{R}^m : ||f(\mathbf{x}) - f(\mathbf{y})||_2 \le L||\mathbf{x} - \mathbf{y}||_2$$

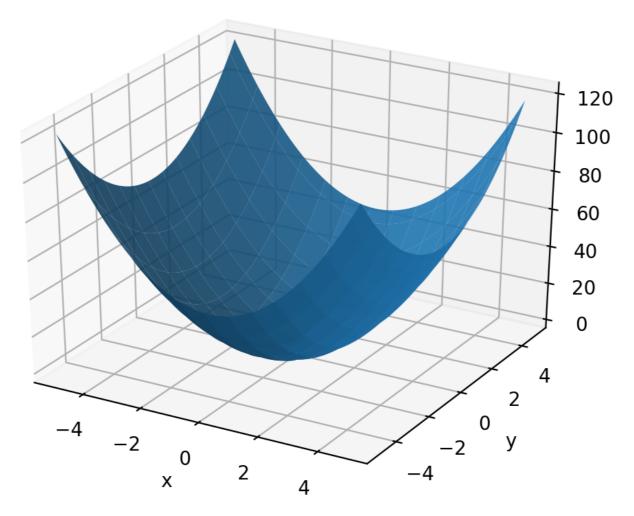
L is essentially an upper bound on the absolute slope of f.

- Gradient descent is guaranteed to converge to a local minimum (eventually) if the learning rate is small enough relative to the steepness of f.
- A function $f: \mathbb{R}^m \to \mathbb{R}$ is Lipschitz-continuous if:

$$\exists L : \forall \mathbf{x}, \mathbf{y} \in \mathbb{R}^m : ||f(\mathbf{x}) - f(\mathbf{y})||_2 \le L||\mathbf{x} - \mathbf{y}||_2$$

- L is essentially an upper bound on the absolute slope of f.
- For learning rate $\epsilon \leq \frac{1}{L}$, gradient descent will converge to a local minimum linearly, i.e., the error is O(1/k) in the iterations k.

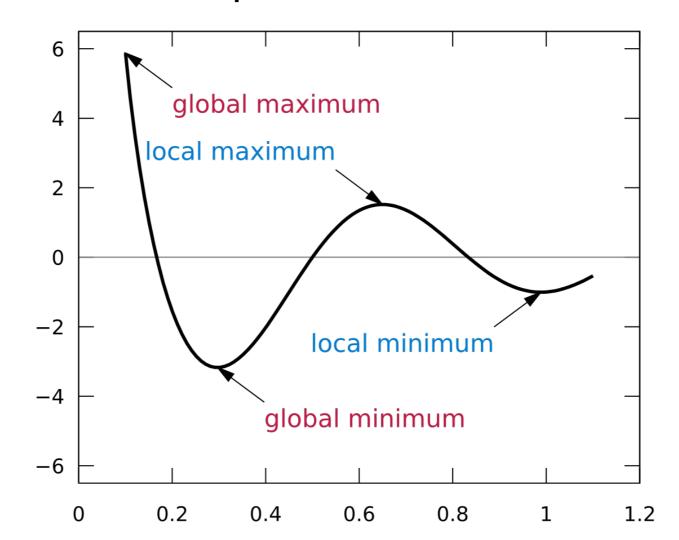
• With linear regression, the cost function f_{MSE} has a single local minimum w.r.t. the weights **w**:



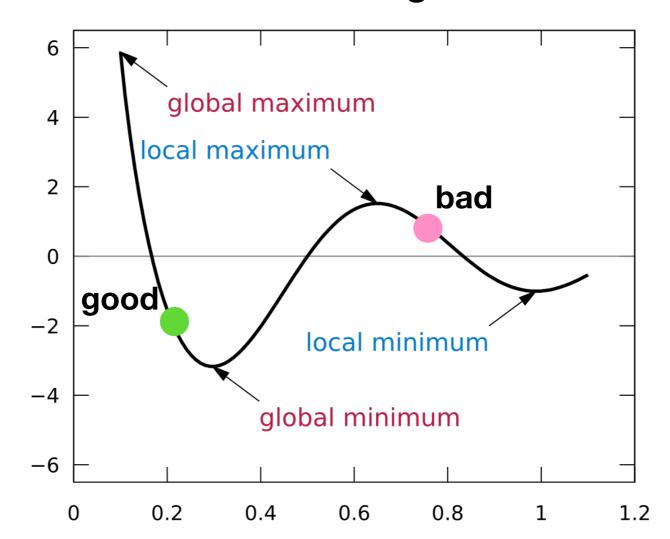
 As long as our learning rate is small enough, we will find the optimal w.

 In general ML and DL models, optimization is usually not so simple, due to:

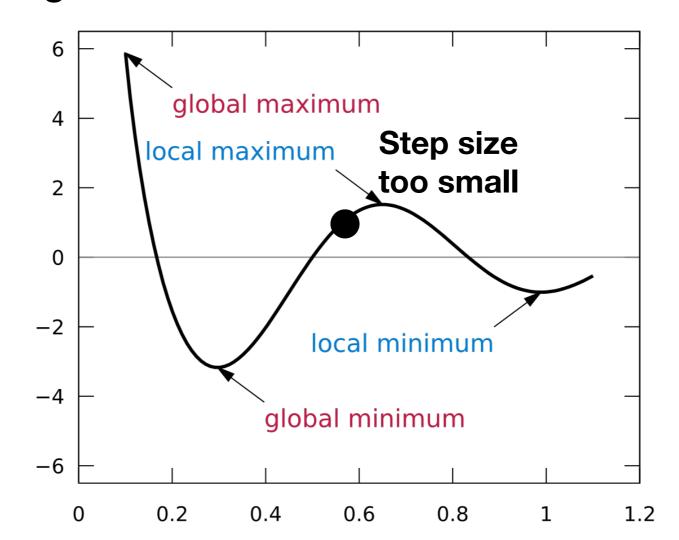
- In general ML and DL models, optimization is usually not so simple, due to:
 - 1. Presence of multiple local minima



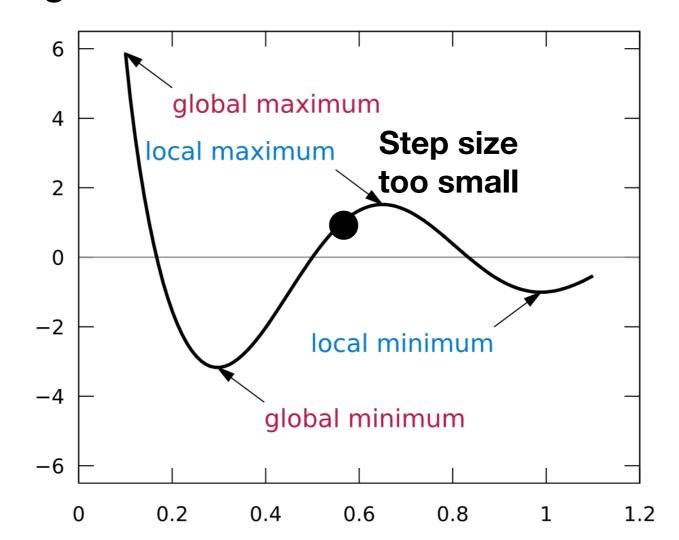
- In general ML and DL models, optimization is usually not so simple, due to:
 - 2. Bad initialization of the weights w.



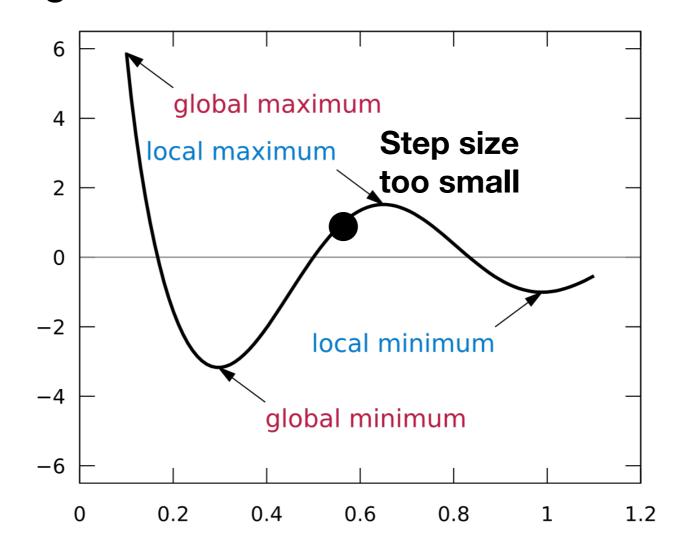
- In general ML and DL models, optimization is usually not so simple, due to:
 - 3. Learning rate is too small.



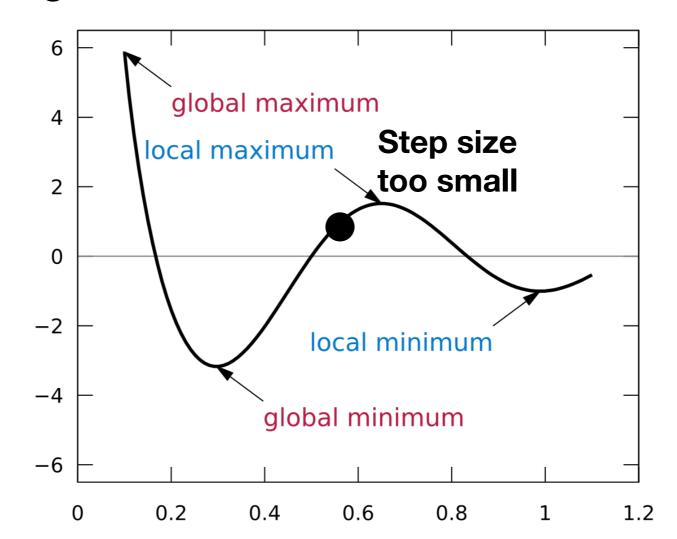
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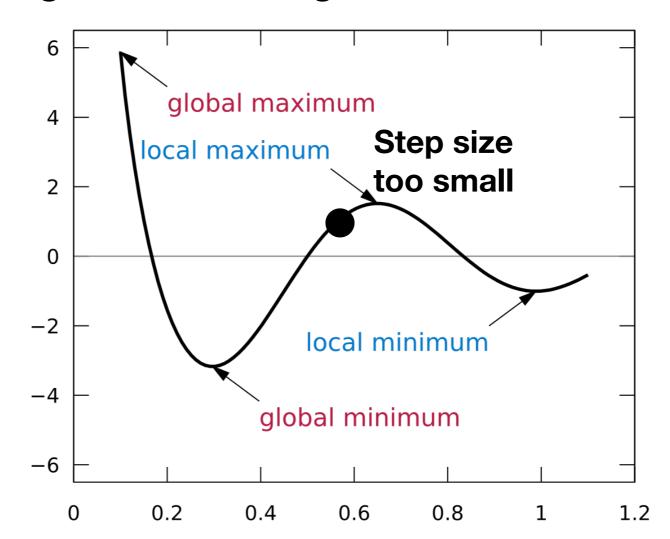
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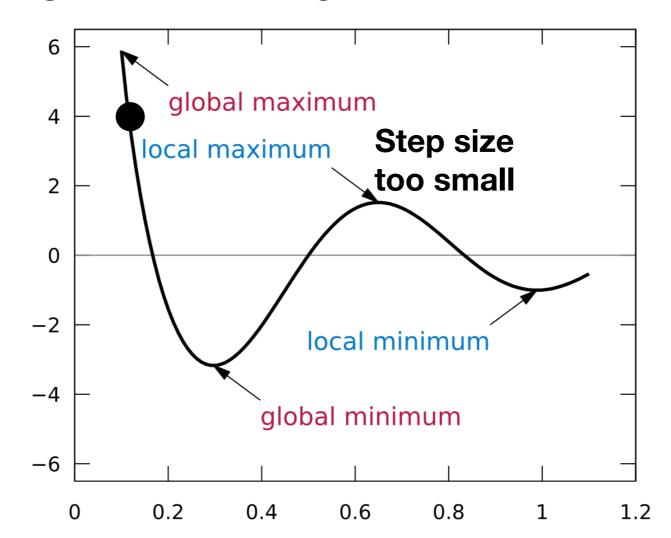
- In general ML and DL models, optimization is usually not so simple, due to:
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- In general ML and DL models, optimization is usually not so simple, due to:
 - 4. Learning rate is too big.



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 - 4. Learning rate is too big.



 In general ML and DL models, optimization is usually not so simple, due to:

4. Learning rate is too big.

(off the chart)

| Good | Go

0.6

0.8

1

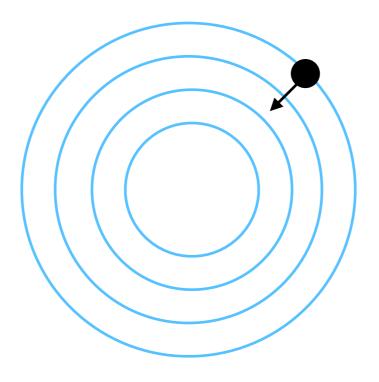
1.2

-6

0.2

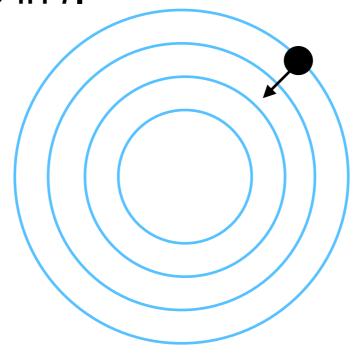
0.4

- With multidimensional weight vectors, badly chosen learning rates can cause more subtle problems.
- Consider the cost f whose level sets are shown below:



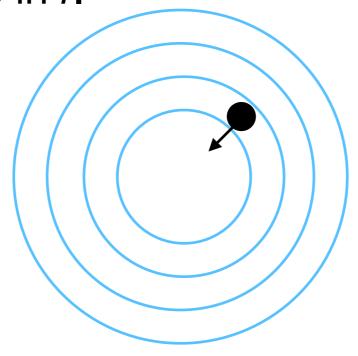
 With multidimensional weight vectors, badly chosen learning rates can cause more subtle problems.

 Gradient descent guides the search along the direction of steepest decrease in f.



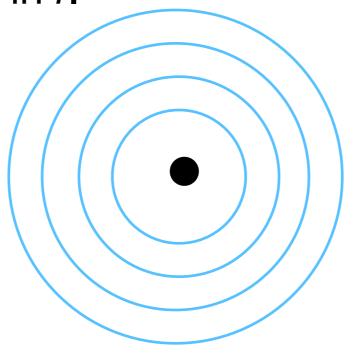
 With multidimensional weight vectors, badly chosen learning rates can cause more subtle problems.

 Gradient descent guides the search along the direction of steepest decrease in f.

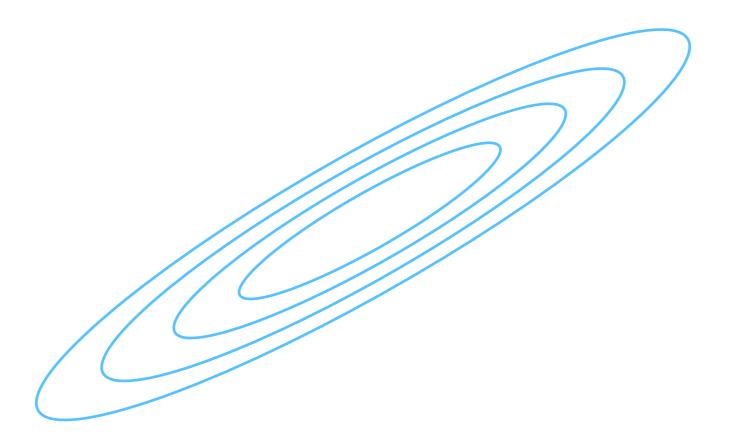


 With multidimensional weight vectors, badly chosen learning rates can cause more subtle problems.

 Gradient descent guides the search along the direction of steepest decrease in f.



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