Module 4: Chapter 6 Bayesian Learning

1. Introduction

- A probabilistic approach to inference
 - It is based on the assumption that the quantities of interest are governed by probability distributions and that optimal decisions can be made by reasoning about these probabilities together with observed data.

- Quantitative approach to weighing the evidence supporting alternative hypotheses
- Important
 - Calculates the explicit probability like naïve Bayes.
 - Naive Bayes classifier competitive, outperforms as a classifier
 - They help Understand learning algorithms that do not explicitly manipulate probabilities

• Bayesian learning methods are relevant to our study of machine learning for two different reasons.

Machine Learning-

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i. Bayesian learning algorithms that calculate explicit probabilities for hypotheses, such as the naive Bayes classifier, are among the most practical approaches to certain types of learning problems.

- ➢ For ex : Michie et al.(1994) provides ar detailed study comparing the naive Bayes classifier to other learning algorithms, including decision tree and neural network algorithms.
- These researchers show that the naive Bayes classifier is competitive with these other learning algorithms in many cases and that in some cases it outperforms these other methods.
- ii. The Bayesian methods are important to our study of machine learning is that they provide a useful perspective for understanding many learning algorithms that do not explicitly manipulate probabilities.

- For ex : We will analyze the algorithms such as the FIND-S and Candidate-Elimination algorithms to determine the conditions under which they output the most probable hypothesis given the training data.
- Bayesian analysis provides an opportunity for the choosing the appropriate alternative error function(cross entropy) in neural network learning algorithms.
- We use a Bayesian perspective to analyze the inductive bias of decision tree learning algorithms that favor short decision trees and examine the closely related *Minimum Description Length principle*.

Feature of Bayesian learning methods include:

- Each observed training example can incrementally decrease or increase the estimated probability that a hypothesis is correct. This provides a more flexible approach to the learning compared to the algorithms that completely eliminate a hypothesis if it is found to be inconsistent with any single example.
- Prior knowledge can be combined with observed data to determine the final probability of a hypothesis. The prior probability is got through (i) a prior probability of each candidate hypothesis (ii) a probability distribution over observed data for each possible hypothesis.
- Bayesian methods can accommodate hypotheses that make probabilistic predictions For ex : hypotheses such as *"this pneumonia patient has a 93% chance of complete recovery"*

- New instances can be classified by combining the predictions of multiple hypotheses, weighted by their probabilities.
- Even in cases where Bayesian methods prove computationally intractable, they can provide a standard of optimal decision making against which other practical methods can be measured.

Difficulty of applying Bayesian method

- Bayesian methods typically require initial knowledge of many probabilities.
- The significant computational cost required to determine the Bayes optimal hypothesis in the general case.

2. BAYES THEOREM

- Determining the **most (best) hypothesis** from some space H, (having initial prior probabilities of various hypotheses) given the observed training data D.
- Calculates the probability of a hypothesis based on its prior probability, the probabilities of observing various data given the hypothesis, and the observed data itself.

Conditional probability



Bowl A has 1 blue and 4 yellow Marbles

Bowl **B** 3 blue and 2 yellow

What is the likelihood of picking a blue marble or a yellow marble from any bowl ?



of outcomes are finite and equally likely the probability of an event happening is

P(event) = # of favourable outcome/#of possible outcome

E x perimental $P(E) = \frac{\# \text{ times } E \text{ occurred}}{\text{ Total } \# \text{ trials}}$

 $P(E) = \frac{\text{number of ways E possible}}{\text{total number of possible outcomes}}$





The probability of picking a blue marble from bowl A =1/5

What is the probability that the picked marble came from **bowl A** given that it is **blue**?







yellow

A Bowl has 2 blue and 3 yellow Marbles Probability of drawing a blue marble is 2/5



Probability of drawing a **second** blue marble is ?

A Bowl has 2 blue and 3 yellow Marbles Probability of drawing a blue marble is 2/5 Probability of drawing a **Second** blue marble is $\frac{1}{4}$



First	Second Blue	Probability of choosing the second given first	
Blue		1/4 =P(B2 B1)	
Blue	yellow	3/4	
Yellow	Blue	2/4	
Yellow	Yellow	2/4	



4 blue 2/5 yellow 3 5 blue y1) P(y2 yellow Probability of first followed by First Second second in a row Blue 1/10 =P(B1) *P(B2|B1) Blue Blue yellow 3/10 =P(B1) *P(Y2|B1) Yellow Blue 3/10 =P(Y1) *P(B2|Y1) =P(Y1)*P(Y2|Y1) Yellow 3/10 Yellow Total =1

Probability of drawing a **first and second** marble in a row with **yellow** is 3/5 * 2/4 = **3/10**



Probability of **second** marble being **yellow** irrespective of the color of the first one is 3/5



e



Notation

- *P(h)* : initial probability that hypothesis *h* holds, before we have observed the training data. *prior probability* of *h* and may reflect any background knowledge we have about the chance that *h* is a correct hypothesis.
- *P(D)* : prior probability that training data D will be observed-no knowledge of h
- P(D|h) : probability of observing data D in which hypothesis h holds.
- *P(h|D)* : probability that *h* holds given the observed training data *D*. posterior probability of *h* reflects the influence of the training data *D*

Bayes theorem:

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 $P(h|D) = \frac{P(D|h)P(h)}{P(D)}$

- The learner considers some set of candidate hypotheses H and is interested in finding the most probable hypothesis $h \in H$ given the observed data D (or at least one of the maximally probable if there are several).
- Any such maximally probable hypothesis is called a *maximum* a *posteriori* (MAP) hypothesis

• We can determine the MAP hypotheses by using Bayes theorem to calculate the posterior probability of each candidate hypothesis.

$h_{MAP} \text{ is a MAP hypothesis provided}$ $h_{MAP} \equiv \underset{h \in H}{\operatorname{argmax}} P(h|D)$ $= \underset{h \in H}{\operatorname{argmax}} \frac{P(D|h)P(h)}{P(D)}$ $= \underset{h \in H}{\operatorname{argmax}} P(D|h)P(h)$

- In some cases, we will assume that every hypothesis in H is equally probable a priori $(P(h_i) = P(h_j)$ for all h_i and h_j in H).
- In this case we can further simplify Equation and need only consider the term *P(D|h)* to find the most probable hypothesis

P(D|h) is often called the *likelihood* of the data *D* given *h*Any hypothesis that maximizes *P(D|h)* is called a *maximum likelihood* (ML) hypothesis, *h_{ML}*.

$$h_{ML} \equiv \operatorname*{argmax}_{h \in H} P(D|h)$$

- Product rule: probability $P(A \land B)$ of a conjunction of two events A and B $P(A \land B) = P(A|B)P(B) = P(B|A)P(A)$
- Sum rule: probability of a disjunction of two events A and B

 $P(A \lor B) = P(A) + P(B) - P(A \land B)$

• Bayes theorem: the posterior probability P(h|D) of h given D

$$P(h|D) = \frac{P(D|h)P(h)}{P(D)}$$

• Theorem of total probability: if events A_1, \ldots, A_n are mutually exclusive with $\sum_{i=1}^n P(A_i) = 1$, then

$$P(B) = \sum_{i=1}^{n} P(B|A_i)P(A_i)$$

TABLE 6.1 Summary of basic probability formulas.

Example:

- To illustrate Bayes rule, consider a medical diagnosis problem in which there are two alternative hypotheses:
- (1) that the patient has a particular form of cancer(2) that the patient does not.

- The available data is from a particular laboratory test with two possible outcomes: + (positive) and -(negative).
- We have prior knowledge that over the entire population of people only .008 have this disease.
- Furthermore, the lab test is only an imperfect indicator of the disease.
- The test returns a correct positive result in only 98% of the cases in which the disease is actually present and a correct negative result in only 97% of the cases in which the disease is not present.
- In other cases, the test returns the opposite result.

	Actual Value (as confirmed by experiment)			
		positives	negatives	
Predicted Value (predicted by the test)	positives	TP True Positive	FP False Positive	
	negatives	FN False Negative	TN True Negative	

$$P(cancer) = .008, \quad P(\neg cancer) = .992$$
$$P(\oplus | cancer) = .98, \quad P(\ominus | cancer) = .02$$
$$P(\oplus | \neg cancer) = .03, \quad P(\ominus | \neg cancer) = .97$$

- Suppose we now observe a new patient for whom the lab test returns a positive result.
- Should we diagnose the patient as having cancer or not?

3. BAYES THEOREM AND CONCEPT LEARNING

- What is the relationship between Bayes theorem and the problem of concept learning?
 - Bayes theorem provides a principled way to calculate the posterior probability of each hypothesis given the training data
 - we can use it as the basis for a straightforward learning algorithm that calculates the probability for each possible hypothesis
 - then outputs the most probable.

3.1 Brute-Force Bayes Concept Learning

- Concept learning problem:
 - assume the learner considers some finite hypothesis space
 H
 - defined over the instance space X,
 - the task is to learn some target concept c : X -> {0,1}.
 - learner is given some sequence of training examples
 ((x₁, d₁)...(x_m, d_m))

• We can design a straightforward concept learning algorithm to output the maximum a posteriori hypothesis, based on Bayes theorem.

BRUTE-FORCE MAP LEARNING algorithm

1. For each hypothesis h in H, calculate the posterior probability $P(h|D) = \frac{P(D|h)P(h)}{P(D)}$

2. Output the hypothesis h_{MAP} with the highest posterior probability $h_{MAP} = \underset{h \in H}{\operatorname{argmax}} P(h|D)$
• what values are to be used for P(h) and for P(D|h)?

- choose the probability distributions *P(h)* and *P(D|h)* in any way you wish, to describe our prior knowledge about the learning task.
- Here let us choose them to be consistent with the following assumptions:

1. The training data **D** is noise free (i.e., $d_i = c(x_i)$).

2. The target concept \boldsymbol{c} is contained in the hypothesis space H

3. We have no a priori reason to believe that any hypothesis is more probable than any other.

- Given these assumptions, what values should we specify for *P(h)*?
 - Given no prior knowledge that one hypothesis is more likely than another, it is reasonable to assign the same prior probability to every hypothesis *h* in *H*.
 - because we assume the target concept is contained in H we should require that these prior probabilities sum to 1.

• Together these constraints imply that we should choose

$$P(h) = \frac{1}{|H|}$$
 for all h in H

- What choice shall we make for *P***(D|h)**?
 - *P(D|h)* is the probability of observing the target values D = (d₁...d_m) for the fixed set of instances (x₁...x_m), given a world in which hypothesis *h* holds (i.e., given a world in which *h* is the correct description of the target concept c).

Since we assume noise-free training data, the probability of observing classification d_i, given h is

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• 1 if d_i = h(x_i) and
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• 0 if $d_i \neq h(x_i)$

$$P(D|h) = \begin{cases} 1 \text{ if } d_i = h(x_i) \text{ for all } d_i \text{ in } D \\ 0 \text{ otherwise} \end{cases}$$

In other words, the probability of data **D** given hypothesis **h** is 1 if **D** is consistent with **h**, and 0 otherwise.

Given these choices for *P(h)* and for *P(D|h)* we now have a fully-defined problem for the above BRUTE-FORCE MAP LEARNING algorithm.

• Step1 :

Let us consider the first step of this algorithm, which uses Bayes theorem to compute the posterior probability *P(h|D)* of each hypothesis *h* given the observed training data *D*.

Recalling Bayes theorem, we have

$$P(h|D) = \frac{P(D|h)P(h)}{P(D)}$$

• Case 1 : *h* is inconsistent with the training data *D*.

$$P(h|D) = \frac{0 \cdot P(h)}{P(D)} = 0 \text{ if } h \text{ is inconsistent with } D$$

The posterior probability of a hypothesis inconsistent with *D* is zero.

• Case 2: h is consistent with *D*.

$$P(h|D) = \frac{1 \cdot \frac{1}{|H|}}{P(D)}$$
$$= \frac{1 \cdot \frac{1}{|H|}}{\frac{|VS_{H,D}|}{|H|}}$$
$$= \frac{1}{|VS_{H,D}|} \text{ if } h \text{ is consistent with } D$$

To summarize, Bayes theorem implies that the posterior probability P(h|D)under our assumed P(h) and P(D|h) is

$$P(h|D) = \begin{cases} \frac{1}{|VS_{H,D}|} & \text{if } h \text{ is consistent with } D\\ 0 & \text{otherwise} \end{cases}$$
(6.5)



FIGURE 6.1

Evolution of posterior probabilities P(h|D) with increasing training data. (a) Uniform priors assign equal probability to each hypothesis. As training data increases first to D1 (b), then to D1 \wedge D2 (c), the posterior probability of inconsistent hypotheses becomes zero, while posterior probabilities increase for hypotheses remaining in the version space.

- The above analysis implies that under our choice for P(h) and P(D|h), every *consistent* hypothesis has posterior probability (1 /|VS_{H,D}|), and every inconsistent hypothesis has posterior probability **O**.
- Every consistent hypothesis is, therefore, **a MAP hypothesis**.

3.2 MAP Hypotheses and Consistent Learners

• Every hypothesis consistent with *D* is a MAP hypothesis

consistent learners:

- We will say that a learning algorithm is a *consistent learner* provided it outputs a hypothesis that commits zero errors over the training examples.
- Given the above analysis, we can conclude that *every consistent learner outputs a MAP hypothesis, if we assume*
 - a uniform prior probability distribution over H (i.e., P(h_i) = P(h_j) for all *i*, *j*), and
 - deterministic, noise free training data (i.e., P(D|h) = 1 if D and h are consistent, and 0 otherwise).

4. MAXIMUM LIKELIHOOD AND LEAST-SQUARED ERROR HYPOTHESES

• Let's we consider the problem of learning a continuous-valued target function.

• A straightforward Bayesian analysis will show that under certain assumptions any learning algorithm that minimizes the squared error between the output hypothesis predictions and the training data will output a maximum likelihood hypothesis.

- Consider the following problem setting:
- Learner **L** considers an instance space **X** and a hypothesis space **H** consisting of some class of real-valued functions defined over X (i.e., each h in H is a function of the form $h : X \rightarrow R$, where R represents the set of real numbers).

- The problem faced by L is to learn an unknown target function f : X → R drawn from H.
- A set of *m* training examples is provided, where the target value of each example is corrupted by random noise drawn according to a Normal probability distribution.

- More precisely, each training example is a pair of the form (x_i, d_i) where
 d_i = f (x_i) + e_i
 - f(x_i): the noise-free value of the target function
 e_i: is a random variable representing the noise.

- It is assumed that the values of the *e_i* are drawn independently and that they are distributed according to a Normal distribution with zero mean.
- The task of the learner is to output a maximum likelihood hypothesis, or, equivalently, a MAP hypothesis assuming all hypotheses are equally probable a priori.

2 basic concepts

- probability densities
- Normal distributions.

- First, in order to discuss probabilities over continuous variables such as e, we must introduce probability densities.
- The reason, roughly, is that we wish for the total probability over all possible values of the random variable to sum to one.

- In the case of continuous variables we cannot achieve this by assigning a finite probability to each of the infinite set of possible values for the random variable.
- Instead, we speak of a probability density for continuous variables such as *e* and require that the integral of this probability density over all possible values be one.

In general we will use lower case p to refer to the probability density function, to distinguish it from a finite probability P (which we will sometimes refer to as a probability mass). The probability density $p(x_0)$ is the limit as ϵ goes to zero, of $\frac{1}{\epsilon}$ times the probability that x will take on a value in the interval $[x_0, x_0 + \epsilon)$.

Probability density function:

$$p(x_0) \equiv \lim_{\epsilon \to 0} \frac{1}{\epsilon} P(x_0 \le x < x_0 + \epsilon)$$

- Second, we stated that the random noise variable e is generated by a Normal probability distribution.
- A Normal distribution is a smooth, bell-shaped distribution that can be completely characterized by its mean μ and its standard deviation σ .

- Given this background we now return to the main issue:
 - showing that the least-squared error hypothesis is, in fact, the maximum likelihood hypothesis within our problem setting.
- We will show this by deriving the maximum likelihood hypothesis, but using lower case *p* to refer to the probability density

$h_{ML} = \operatorname*{argmax}_{h \in H} p(D|h)$

- we assume a fixed set of training instances $(x_1, x_2 \dots x_m)$ and
- Therefore consider the data D to be the corresponding sequence of target values $D = (d_1, d_2 \dots d_m)$.
- Here $d_i = f(x_i) + e_i$.

- Assuming the training examples are mutually independent given *h*,
- we can write P(D|h) as the product of the various $p(d_i|h)$

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \prod_{i=1}^{m} p(d_i | h)$$

- Given that the noise e_i obeys a Normal distribution with zero mean and unknown variance σ^2 , each d_i must also obey a Normal distribution with variance σ^2 centered around the true target value $f(x_i)$ rather than zero.
- Therefore $p(d_i \mid h)$ can **be** written as a Normal distribution with variance σ^2 and mean $\mu = f(x_i)$

$$a_{ML} = \operatorname*{argmax}_{h \in H} \prod_{i=1}^{m} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2\sigma^2}(d_i - \mu)^2}$$
$$= \operatorname*{argmax}_{h \in H} \prod_{i=1}^{m} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2\sigma^2}(d_i - h(x_i))^2}$$

We now apply a transformation that is common in maximum likelihood calculations: Rather than maximizing the above complicated expression we shall choose to maximize its (less complicated) logarithm. This is justified because $\ln p$ is a monotonic function of p. Therefore maximizing $\ln p$ also maximizes p.

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \sum_{i=1}^{m} \ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} (d_i - h(x_i))^2$$

The first term in this expression is a constant independent of h, and can therefore be discarded, yielding

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \sum_{i=1}^{m} -\frac{1}{2\sigma^2} (d_i - h(x_i))^2$$

Maximizing this negative quantity is equivalent to minimizing the corresponding positive quantity.

$$h_{ML} = \operatorname*{argmin}_{h \in H} \sum_{i=1}^{m} \frac{1}{2\sigma^2} (d_i - h(x_i))^2$$

Finally, we can again discard constants that are independent of h.

$$h_{ML} = \underset{h \in H}{\operatorname{argmin}} \sum_{i=1}^{m} (d_i - h(x_i))^2$$
(6.6)

• Thus, Equation shows that the maximum likelihood hypothesis h_{ML} is the one that minimizes the sum of the squared errors between the observed training values d_i and the hypothesis predictions $h(x_i)$.

• Limitations of this problem setting.

The above analysis considers noise only in the *target value* of the training example and does not consider noise in the *attributes describing the instances themselves*.



Machine Learning-Module 4
Chapter 6: Bayesian Learning

Introduction

- Bayesian reasoning provides a probabilistic approach to inference.
- Bayesian learning methods are relevant to our study of machine learning for two different reasons.
 - i. Bayesian learning algorithms that calculate explicit probabilities for hypotheses, such as the naive Bayes classifier, are among the most practical approaches to certain types of learning problems.

- ➢ For ex : Michie et al.(1994) provide a detailed study comparing the naive Bayes classifier to other learning algorithms, including decision tree and neural network algorithms.
- These researchers show that the naive Bayes classifier is competitive with these other learning algorithms in many cases and that in some cases it outperforms these other methods.
- ii. The Bayesian methods are important to our study of machine learning is that they provide a useful perspective for understanding many learning algorithms that do not explicitly manipulate probabilities.

- ➢ For ex : We will analyze the algorithms such as the FIND-S and Candidate-Elimination algorithms to determine the conditions under which they output the most probable hypothesis given the training data.
- Bayesian analysis provides an opportunity for the choosing the appropriate alternative error function(cross entropy) in neural network learning algorithms.
- We use a Bayesian perspective to analyze the inductive bias of decision tree learning algorithms that favor short decision trees and examine the closely related *Minimum Description Length principle*.

Features of Bayesian Learning methods

- Each observed training example can incrementally decrease or increase the estimated probability that a hypothesis is correct. This provides a more flexible approach to the learning compared to the algorithms that completely eliminate a hypothesis if it is found to be inconsistent with any single example.
- Prior knowledge can be combined with observed data to determine the final probability of a hypothesis. The prior probability is got through (i) a prior probability of each candidate hypothesis (ii) a probability distribution over observed data for each possible hypothesis.
- Bayesian methods can accommodate hypotheses that make probabilistic predictions For ex : hypotheses such as *"this pneumonia patient has a 93% chance of complete recovery"*

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- New instances can be classified by combining the predictions of multiple hypotheses, weighted by their probabilities.
- Even in cases where Bayesian methods prove computationally intractable, they can provide a standard of optimal decision making against which other practical methods can be measured.

Practical Difficulties in applying Bayesian Methods

- Bayesian methods typically require initial knowledge of many probabilities.
- The significant computational cost required to determine the Bayes optimal hypothesis in the general case.

Bayes Theorem

- In machine learning we are interested in determining the best hypothesis from some space *H*, given the observed training data *D*.
- Bayes theorem provides a way to calculate the probability of a hypothesis based on its prior probability, the probabilities of observing various data given the hypothesis, and the observed data itself.
- To define Bayes theorem precisely, let us define the following notations
 - > P(h) denote the initial probability or prior probability that hypothesis h holds.
 - > P(D) denote the prior probability that training data D will be observed.

- > P(D/h) –denote the probability of observing data D given some world in which hypothesis h holds.
- P(h/D) denote the posterior probability of h because it reflects our confidence that h holds after we have seen the training data D.
- Bayes theorem is the cornerstone of Bayesian learning methods because it provides a way to calculate the posterior probability *P(h/D)* from the prior probability *P(h)*, together with *P(D)* and *P(D/h)*

$$P(h/D) = \frac{P(D/h)P(h)}{P(D)} \longrightarrow Eqn \ 6.1$$

P(h/D) increases with *P(h)* and with *P(D/h)* according to Bayes theorem.

- It is also reasonable to see that P(h/D) decreases as P(D) increases, because the more probable it is that D will be observed independent of h, the less evidence D provides in support of h.
- In many learning scenarios, the learner considers some set of candidate hypotheses H and is interested in finding the most probable hypothesis $h \in H$ given the observed data D.
- Any such maximally probable hypothesis is called a *maximum a posteriori (MAP)* hypothesis.
- We can determine the *MAP* hypotheses by using Bayes theorem to calculate the posterior probability of each candidate hypothesis.
- More precisely, we will say that h_{MAP} is a MAP hypothesis provided:

$$h_{MAP} \equiv \underset{h \in H}{\operatorname{argmax}} P(h|D)$$

= $\underset{h \in H}{\operatorname{argmax}} \frac{P(D|h) P(h)}{P(D)}$
= $\underset{h \in H}{\operatorname{argmax}} P(D|h) P(h) \longrightarrow Eqn \ 6.2$

- Notice in the final step above we dropped the term P(D) because it is a constant independent of h.
- In some cases, we will assume that every hypothesis in H is equally probable *a priori* ($P(h_i) = P(h_j)$ for all h_i and h_j in H). In this case we can further simplify Eqn 6.2 and need only consider the term P(D/h) to find the most probable hypothesis.

P(*D*/*h*) is often called the likelihood of the data *D* given *h*, and any hypothesis that maximizes *P*(*D*/*h*) is called a *maximum likelihood* (ML) hypothesis, *h*_{ML}

$$h_{ML} \equiv \underset{h \in H}{\operatorname{argmax}} P(D|h) \longrightarrow Eqn \ 6.3$$

In order to make clear the connection to machine learning problems, we have learnt Bayes theorem above by referring to the data *D* as training examples of some target function and referring to *H* as the space of candidate target functions.

An Example

- To illustrate Bayes rule, consider a medical diagnosis problem in which there are two alternative hypotheses:
 - i. that the patient has a particular form of cancer
 - ii. that the patient does not
- The available data is from a particular laboratory test with two possible outcomes: \bigoplus (positive) and \ominus (negative).
- We have prior knowledge that over the entire population of people only **.008** have this disease.
- The test returns a correct positive result in only **98%** of the cases in which the disease is actually present and a correct negative result in only **97%** of the cases in which the disease is not present.

- In other cases, the test returns the opposite result.
- The above situation can be summarized by the following probabilities:

 $P(cancer) = .008, \qquad P(\neg cancer) = .992$ $P(\oplus | cancer) = .98, \qquad P(\ominus | cancer) = .02$ $P(\oplus | \neg cancer) = .03, \qquad P(\ominus | \neg cancer) = .97$

- Suppose we now observe a new patient for whom the lab test returns a positive result. *Should we diagnose the patient as having cancer or not?*
- The maximum a posteriori hypothesis can be found using Eqn 6.2: $P(cancer/\oplus) = P(\oplus/cancer) P(cancer) = 0.98 * 0.008 = 0.0078$ $P(\neg cancer/\oplus) = P(\oplus/\neg cancer) P(\neg cancer) = 0.03 * 0.992 = 0.0298$

- Thus, $h_{MAP} = -cancer$
- Notice that while the posterior probability of cancer is significantly higher than its prior probability, the most probable hypothesis is still that the patient does not have cancer.
- As this example illustrates, the result of Bayesian inference depends strongly on the prior probabilities, which must be available in order to apply the method directly.
- Basic formulas for calculating probabilities are summarized in Table 6.1.

• Product rule: probability $P(A \land B)$ of a conjunction of two events A and B

 $P(A \wedge B) = P(A|B)P(B) = P(B|A)P(A)$

• Sum rule: probability of a disjunction of two events A and B

 $P(A \lor B) = P(A) + P(B) - P(A \land B)$

• Bayes theorem: the posterior probability P(h|D) of h given D

$$P(h|D) = \frac{P(D|h)P(h)}{P(D)}$$

• Theorem of total probability: if events A_1, \ldots, A_n are mutually exclusive with $\sum_{i=1}^n P(A_i) = 1$, then

$$P(B) = \sum_{i=1}^{n} P(B|A_i) P(A_i)$$

Table 6.1: Summary of basic probability formulas

Bayes Theorem and Concept Learning

- Consider the concept learning problem in which we assume that learner considers some finite hypothesis space *H* defined over the instance space *X*, in which the task is to learn some target concept *c:X→{0,1}*.
- Let us assume that the learner is given some sequence of training examples $\langle x_{l}, d_{l} \rangle \ldots \langle x_{m}, d_{m} \rangle \rangle$ where x_{i} is some instance from X and where d_{i} is the target value of x_{i} (i.e., $d_{i}=c(x_{i})$).
- To understand in very simple way, let us make one more assumption that the sequence of instances $\langle x_l \dots x_m \rangle$ is held fixed, so that the training data D can be written simply as the sequence of target values $D = \langle d_l \dots d_m \rangle$.

• Thus, we can design a straightforward concept learning algorithm to output the *maximum a posteriori* hypothesis, based on Bayes theorem, as follows:

Brute-Force MAP Learning Algorithm

1. For each hypothesis h in H, calculate the posterior probability

$$P(h|D) = \frac{P(D|h)P(h)}{P(D)}$$

2. Output the hypothesis h_{MAP} with the highest posterior probability $h_{MAP} = \underset{h \in H}{\operatorname{argmax}} P(h|D)$

This algorithm may require significant computation, because it applies Bayes theorem to each hypothesis in H to calculate P(h/D).

- In order to specify a learning problem for the Brute-force MAP learning algorithm we must specify what values are to be used for *P(h)* and for *P(D/h)*.
- Let us choose them to be consistent with the following assumptions:
 - i. The training data **D** is noise free (i.e., $d_i = c(x_i)$).
 - ii. The target concept c is contained in the hypothesis space H.
 - iii. We have no *a priori* reason to believe that any hypothesis is more probable than any other.
- Given these assumptions, we specify the value for P(h) in the following way
 - Given no prior knowledge that one hypothesis is more likely than another, it is reasonable to assign the same prior probability to every hypothesis *h* in *H*.

Furthermore, because we assume the target concept is contained in *H* we should require that these prior probabilities sum to 1.

> Together these constraints imply that we should choose

$$P(h) = \frac{1}{|H|}$$
 for all h in H

- The value for P(D/h) can be specified in the following way:
 - > P(D/h) is the probability of observing the target values $D = \langle d_l \\ \dots \\ d_m \rangle$ for the fixed set of instances $\langle x_1 \dots \\ x_m \rangle$ given a world in which hypothesis h holds.
 - > Since we assume noise-free training data, the probability of observing classification d_i given h is just 1 if $d_i = h(x_i)$ and 0 if $d_i \neq h(x_i)$

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➤ Therefore,

$$P(D|h) = \begin{cases} 1 \text{ if } d_i = h(x_i) \text{ for all } d_i \text{ in } D \\ 0 \text{ otherwise} \end{cases} \xrightarrow{} Eqn \ 6.4$$

- Given these choices for P(h) and for P(D/h) we now have a fullydefined problem for the above Brute-Force *MAP* learning algorithm.
- Now, let us consider the first step of this algorithm, which uses Bayes theorem to compute the posterior probability *P(h/D)* of each hypothesis *h* given the observed training data *D*.
- Recalling Bayes theorem, we have

$$P(h|D) = \frac{P(D|h)P(h)}{P(D)}$$

• Case 1: Consider where *h* is inconsistent with the training data *D*

Since Eqn 6.4 defines P(D/h) to be 0 when h is inconsistent with D, we have $P(h|D) = \frac{0 \cdot P(h)}{P(D)} = 0$ if h is inconsistent with D

The posterior probability of a hypothesis inconsistent with D is zero.

- Case 2:Consider the case where *h* is consistent with *D*.
 - Since Eqn 6.4 defines P(D/h) to be 1 when h is consistent with D, we have

$$P(h|D) = \frac{1 \cdot \frac{1}{|H|}}{P(D)}$$
$$= \frac{1 \cdot \frac{1}{|H|}}{\frac{|VS_{H,D}|}{|H|}}$$
$$= \frac{1}{|VS_{H,D}|} \text{ if } h \text{ is consistent with } D$$

where $VS_{H,D}$ is the subset of hypotheses from H that are consistent with D.

Verification of the value $P(D) = \frac{|VSH_D|}{|H|}$ for concept learning

- It is easy to verify that $P(D) = \frac{|VSH_D|}{|H|}$, because the sum over all hypotheses of P(h/D) must be one and because the number of hypotheses from H consistent with D is by definition $|VS_{H,D}|$
- Alternatively, we can derive P(D) from the theorem of total probability and the fact that the hypotheses are mutually exclusive (i.e., $(\forall i \neq j)(P(h_i \land h_j) = 0))$

$$P(D) = \sum_{h_i \in H} P(D|h_i) P(h_i)$$

|H|

$$= \sum_{h_i \in VS_{H,D}} 1 \cdot \frac{1}{|H|} + \sum_{h_i \notin VS_{H,D}} 0 \cdot \frac{1}{|H|}$$
$$= \sum_{h_i \in VS_{H,D}} 1 \cdot \frac{1}{|H|}$$
$$= \frac{|VS_{H,D}|}{|H|}$$

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• To summarize, Bayes theorem implies that the posterior probability P(h/D) under our assumed P(h) and P(D/h) is

$$P(h|D) = \begin{cases} \frac{1}{|VS_{H,D}|} & \text{if } h \text{ is consistent with } D\\ 0 & \text{otherwise} \end{cases} \longrightarrow Eqn \ 6.5$$

where $/VS_{H,D}/$ is the number of hypotheses from H consistent with D.

• The evolution of probabilities associated with hypotheses is depicted schematically in figure 6.1. Initially figure 6.1(a) shows all hypotheses have the same probability. As the training data accumulates (figure 6.1(b) & figure 6.1(c)) the posterior probability for inconsistent hypotheses becomes zero while the total probability summing to one is shared equally among the remaining consistent hypotheses.

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Figure 6.1: Evolution of posterior probabilities P(h/D) with increasing training data

MAP Hypotheses and Consistent Learners

- Given the above analysis, every consistent learner outputs a MAP hypothesis, if we assume a uniform prior probability distribution over H (i.e., P(h_i) = P(h_j) for all i, j), and if we assume deterministic, noise free training data (i.e., P(D/h) = 1 if D and h are consistent, and 0 otherwise).
- For ex :
 - Consider the Find-S concept learning algorithm. The Find-S searches the hypothesis space *H* from specific to general hypotheses, outputting a maximally specific consistent hypothesis.
 - Example Because FIND-S outputs a consistent hypothesis, we know that it will output a *MAP* hypothesis under the probability distributions P(h) and P(D/h) defined.

- Actually, FIND-S does not explicitly manipulate probabilities at all-it simply outputs a maximally specific member of the version space.
- However, by identifying distributions for P(h) and P(D/h) under which its output hypotheses will be MAP hypotheses, we have a useful way of characterizing the behavior of FIND-S.
- Are there other probability distributions for *P(h)* and *P(D/h)* under which FIND-S outputs MAP hypotheses?
 - ➢ Yes. Because FIND-S outputs a maximally specific hypothesis from the version space, its output hypothesis will be a MAP hypothesis relative to any prior probability distribution that favors more specific hypotheses.

- ➤ More precisely, suppose H is any probability distribution P(h) over H that assigns P(h₁) ≥ P(h₂) if h_l is more specific than h₂.
- > Then it can be shown hat FIND-S outputs a MAP hypothesis assuming the prior distribution \mathcal{H} and the same distribution P(D/h)
- To summarize, the Bayesian framework allows one way to characterize the behavior of learning algorithms (e.g., FIND-S), even when the learning algorithm does not explicitly manipulate probabilities.

Definitions of various Probability Terms

Random Variable: A random variable, usually written X, is a variable whose possible values are numerical outcomes of a random phenomenon.

There are two types of random variables, *discrete* and *continuous*.

For ex : Random variable can be defined for a coin flip as follows $X = \begin{cases} 1 & if it is head \\ 0 & if it is tail \end{cases}$

Discrete Random Variable : The variables which can take distinct/separate values are called discrete random variables.

For ex : Flipping a fair coin, rolling a dice

Continuous Random Variable : The variables which can take any values in a range are called continuous random variables.

For ex : Height and Weight of the person, Mass of an animal

Probability Distribution : It is a mathematical function that provides the probabilities of occurrence of different possible outcomes in an experiment.

Constructing a probability distribution for random variable

- Let us take random variable,
 X = no of heads after 3 flips of a fair coin
- Then the probability distribution table can be written as follows:

Outcomes (No of Heads)	X=0	X=1	X=2	X=3
Probability	1/8	3/8	3/8	1/8

Maximum Likelihood and Least **Squared** Error **Hypotheses**

- Many learning approaches such as neural network learning, linear regression and polynomial curve fitting will face a problem of learning a continuous-valued target function.
- A straightforward Bayesian analysis will show that *under certain* assumptions any learning algorithm that minimizes the squared error between the output hypothesis predictions and the training data will output a maximum likelihood hypothesis.
- Consider the following problem setting. Learner L considers an instance space X and a hypothesis space H consisting of some class of real-valued functions defined over X. (i.e., each h in H is a function of the form $h: X \to \mathbb{R}$, where \mathbb{R} represents the set of real numbers). 11-11-2019

- The problem faced by *L* is to learn an unknown target function *f*: $X \rightarrow \mathbb{R}$ drawn from *H*.
- A set of *m* training examples is provided, where the target value of each example is corrupted by random noise drawn according to a Normal probability distribution.
- More precisely, each training example is a pair of the form $\langle x_i, d_i \rangle$ where $d_i = f(x_i) + e_i$. Here $f(x_i)$ is the noise-free value of the target function and e_i is a random variable representing the noise.
- It is assumed that the values of the e_i are drawn independently and that they are distributed according to a Normal distribution with zero mean.
- The task of the learner is to output a maximum likelihood hypothesis, or, equivalently, a MAP hypothesis assuming all hypotheses are equally probable *a* priori. 33

- As a simple example of such a problem is learning a linear function, though our analysis applies to learning arbitrary real-valued functions.
- The figure 6.2 illustrates a linear target function f depicted by the solid line, and a set of noisy training examples of this target function.
- The dashed line corresponds to the hypothesis h_{ML} with least-squared training error, hence the maximum likelihood hypothesis.
- The maximum likelihood hypothesis is not necessarily identical to the correct hypothesis, *f*, because it is inferred from only a limited sample of noisy training data.



Figure 6.2 :Learning a real-valued function. The target function f corresponds to the solid line. The training examples (x_i, d_i) are assumed to have Normally distributed noise e_i with zero mean added to the true target value $f(x_i)$. The dashed line corresponds to the linear function that minimizes the sum of squared errors.

Review of Basic Concepts from Probability Theory

Probability Density Function

- In order to know probabilities over continuous variables such as *e*, we must know about probability densities.
- The reason, roughly, is that we wish for the total probability over all possible values of the random variable to sum to one.

Definition : A probability density function (PDF), or density of a continuous random variable, is a function that describes the relative likelihood for this random variable to take on a given value.

• In the case of continuous variables we cannot achieve this by assigning a finite probability to each of the infinite set of possible values for the random variable.

- Instead, we take a probability density for continuous variables such as *e* and require that the integral of this probability density over all possible values be one.
- In general we will use lower case p to refer to the probability density function, to distinguish it from a finite probability P.
- The probability density $p(x_0)$ is the limit as ϵ goes to zero, of $\frac{1}{\epsilon}$ times the probability that x will take on a value in the interval $[x_0, x_0 + \epsilon)$
- The probability density function is

$$p(x_0) \equiv \lim_{\epsilon \to 0} \frac{1}{\epsilon} P(x_0 \le x < x_0 + \epsilon)$$
Normal Distribution

- A Normal distribution is a smooth, bell-shaped distribution that can be completely characterized by its mean μ and its standard deviation σ .
- A Normal distribution (also called a Gaussian distribution) is defined by the probability density function

$$p(x) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2}(\frac{x-\mu}{\sigma})^2}$$

A Normal distribution is fully determined by two parameters in the above formula: μ and σ

• If the random variable *X* follows a normal distribution, then:

- The probability that X will fall into the interval (a,b) is given by $\int_{a}^{b} p(x) dx$
- > The expected, or mean value of *X*, *E*[*X*], is $E[X] = \mu$
- The variance of *X*, *Var*(*X*), is $Var(X) = \sigma^2$
- \succ The standard deviation of *X*, $\sigma_{X_{1}}$ is

 $\sigma_x = \sigma$

Prove that Least Squared Hypothesis is Maximum Likelihood Hypothesis

• We will show this by deriving the maximum likelihood hypothesis starting with our earlier definition *Eqn 6.3* but using lower case *p* to refer to the probability density

$$h_{ML} = \operatorname*{argmax}_{h \in H} p(D|h)$$

- We assume a fixed set of training instances $\langle x_1 \dots x_m \rangle$ and therefore consider the data D to be the corresponding sequence of target values $D = \langle d_1 \dots d_m \rangle$. Here $d_i = f(x_i) + e_i$.
- Assuming the training examples are mutually independent given h, we can write P(D/h) as the product of the various $p(d_i/h)$

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \prod_{i=1}^{m} p(d_i | h)$$

- Given that the noise e_i obeys a Normal distribution with zero mean and unknown variance σ^2 , each d_i must also obey a Normal distribution with variance σ^2 centered around the true target value $f(x_i)$ rather than zero.
- Therefore $p(d_i/h)$ can be written as a Normal distribution with variance σ^2 and mean $\mu = f(x_i)$.
- Let us write the formula for this Normal distribution to describe $p(d_i/h)$, beginning with the general formula for a Normal distribution and then substituting appropriate μ and σ^2

• Because we are writing the expression for the probability of d_i given that h is the correct description of the target function f, we will also substitute $p=f(x_i)=h(x_i)$, yielding

$$h_{ML} = \operatorname*{argmax}_{h \in H} \prod_{i=1}^{m} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2\sigma^2}(d_i - \mu)^2}$$
$$= \operatorname*{argmax}_{h \in H} \prod_{i=1}^{m} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2\sigma^2}(d_i - h(x_i))^2}$$

- We now apply a transformation that is common in maximum likelihood calculations. Rather than maximizing the above complicated expression we shall choose to maximize its (less complicated) logarithm.
- This is justified because *ln p* is a monotonic function of *p*. Therefore maximizing *ln p* also maximizes *p*.

$$h_{ML} = \operatorname*{argmax}_{h \in H} \sum_{i=1}^{m} \ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} (d_i - h(x_i))^2$$

• The first term in this expression is a constant independent of h, and can therefore be discarded, yielding

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \sum_{i=1}^{m} -\frac{1}{2\sigma^2} (d_i - h(x_i))^2$$

• Maximizing this negative quantity is equivalent to minimizing the corresponding positive quantity.

$$h_{ML} = \operatorname*{argmin}_{h \in H} \sum_{i=1}^{m} \frac{1}{2\sigma^2} (d_i - h(x_i))^2$$

• Finally, we can again discard constants that are independent of h.

$$h_{ML} = \underset{h \in H}{\operatorname{argmin}} \sum_{i=1}^{m} (d_i - h(x_i))^2 \longrightarrow Eqn \ 6.6$$

- Thus, Eqn 6.6 shows that the maximum likelihood hypothesis h_{ML} the one that minimizes the sum of the squared errors between the observed training values d_i and the hypothesis predictions $h(x_i)$
- This holds under the assumption that the observed training values d_i are generated by adding random noise to the true target value, where this random noise is drawn independently for each example from a Normal distribution with zero mean.
- As the above derivation makes clear, the squared error term $(d_i h(x_i))^2$ directly from the exponent in the definition of the Normal distribution.

- Notice the structure of the above derivation involves selecting the hypothesis that maximizes the logarithm of the likelihood (ln p(D/h)) in order to determine the most probable hypothesis.
- This approach of working with the log likelihood is common to many Bayesian analyses, because it is often more mathematically tractable than working directly with the likelihood.
- In all cases, the maximum likelihood hypothesis might not be the MAP hypothesis, but if one assumes uniform prior probabilities over the hypotheses then it is.
- Minimizing the sum of squared errors is a common approach in many neural network, curve fitting, and other approaches to approximating real-valued functions.

Reason to choose Normal distribution to characterize noise

- i. It allows for a mathematically straightforward analysis.
- ii. The smooth, bell-shaped distribution is a good approximation to many types of noise in physical systems.

Maximum Likelihood Hypotheses for Predicting Probabilities

- Here we will derive criterion for a setting that is common in neural network learning: learning to predict probabilities.
- Consider the setting in which we wish to learn a nondeterministic (probabilistic) function $f: X \rightarrow \{0,1\}$, which has two discrete output values.
- For ex: the instance space X might represent medical patients in terms of their symptoms, and the target function f(x) might be 1 if the patient survives the disease and 0 if not.
- In this case we might well expect f to be probabilistic. For ex: among a collection of patients exhibiting the same set of observable symptoms, we might find that 92% survive, and 8% do not. Machine Learning-15CS73 47

- This unpredictability could arise from our inability to observe all the important distinguishing features of the patients, or from some genuinely probabilistic mechanism in the evolution of the disease.
- The effect is that we have a target function f(x) whose output is a probabilistic function of the input.
- Given this problem setting, we might wish to learn a neural network (or other real-valued function approximator) whose output is the probability that f(x)=1.
- In other words, we seek to learn the target function, $f': X \rightarrow [0,1]$, such that f'(x) = P(f(x) = 1).
- In order to learn f', we can train a neural network directly from the observed training examples of f, and derive a maximum likelihood hypothesis for f'.

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- To find a maximum likelihood hypothesis for f' we must first obtain an expression for P(D/h).
- Let us assume the training data D is of the form $D = \{ <x_b, d_l > ... < <x_m, d_m > \}$, where d_i is the observed 0 or 1 value for $f(x_i)$.
- Thus treating both x_i and d_i as random variables, and assuming that each training example is drawn independently, we can write P(D/h) as

$$P(D|h) = \prod_{i=1}^{m} P(x_i, d_i|h) \longrightarrow Eqn \ 6.7$$

• It is reasonable to assume, that the probability of encountering any particular instance x_i is independent of the hypothesis h. For ex: the probability that our training set contains a particular patient x_i is independent of our hypothesis about survival rates.

• When *x* is independent of *h* we can rewrite the *Eqn 6.7* using the product rule of probability as

$$P(D|h) = \prod_{i=1}^{m} P(x_i, d_i|h) = \prod_{i=1}^{m} P(d_i|h, x_i) P(x_i) \longrightarrow Eqn \ 6.8$$

• The probability of $P(d_i/h, x_i)$ of observing $d_i=1$ for a single instance x_i , given a world in which hypothesis h holds is $h(x_i)$ i.e., $P(d_i=1/h, x_i) = h(x_i)$ and in general

$$P(d_i|h, x_i) = \begin{cases} h(x_i) & \text{if } d_i = 1 \\ (1 - h(x_i)) & \text{if } d_i = 0 \end{cases} \longrightarrow Eqn \ 6.9$$

In order to substitute *Eqn 6.9* into *Eqn 6.8*, let us re-express *Eqn 6.9* in a more mathematically manipulable form, as

$$P(d_i|h, x_i) = h(x_i)^{d_i} (1 - h(x_i))^{1 - d_i} \longrightarrow Eqn \ 6.10$$

It is easy to verify that the expressions in Eqn 6.9 and Eqn 6.10 are equivalent. We can use Eqn 6.10 to substitute for P(d_i/h,x_i) in Eqn 6.8 to obtain

$$P(D|h) = \prod_{i=1}^{m} h(x_i)^{d_i} (1 - h(x_i))^{1 - d_i} P(x_i) \longrightarrow Eqn \ 6.1.$$

• Now we write an expression for the maximum likelihood hypothesis

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \prod_{i=1}^{m} h(x_i)^{d_i} (1 - h(x_i))^{1 - d_i} P(x_i)$$

• The last term is a constant independent of h, so it can be dropped

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \prod_{i=1}^{m} h(x_i)^{d_i} (1 - h(x_i))^{1 - d_i} \longrightarrow Eqn \ 6.12$$

• As in earlier cases, we will find it easier to work with the log of the likelihood, yielding

$$h_{ML} = \underset{h \in H}{\operatorname{argmax}} \sum_{i=1}^{m} d_i \ln h(x_i) + (1 - d_i) \ln(1 - h(x_i)) \longrightarrow Eqn \ 6.13$$

- *Eqn 6.13* describes the quantity that must be maximized in order to obtain the maximum likelihood hypothesis in our current problem setting.
- This result is analogous to our earlier result showing that minimizing the sum of squared errors produces the maximum likelihood hypothesis in the earlier problem setting.

Gradient Search to Maximize Likelihood in a Neural Net

- Let us take G(h,D) to denote the quantity of Maximum Likelihood hypotheses for the probabilistic target function.
- Our objective here is to derive a weight-training rule for neural network learning that seeks to maximize G(h,D) using gradient ascent.
- The gradient of *G*(*h*,*D*) is given by the vector of partial derivatives of *G*(*h*,*D*) with respect to the various network weights that define the hypothesis *h* represented by the learned network.
- In this case, the partial derivative of G(h,D) with respect to weight w_{ik} from input k to unit j is

$$\frac{\partial G(h, D)}{\partial w_{jk}} = \sum_{i=1}^{m} \frac{\partial G(h, D)}{\partial h(x_i)} \frac{\partial h(x_i)}{\partial w_{jk}}$$
$$= \sum_{i=1}^{m} \frac{\partial (d_i \ln h(x_i) + (1 - d_i) \ln(1 - h(x_i)))}{\partial h(x_i)} \frac{\partial h(x_i)}{\partial w_{jk}}$$
$$= \sum_{i=1}^{m} \frac{d_i - h(x_i)}{h(x_i)(1 - h(x_i))} \frac{\partial h(x_i)}{\partial w_{jk}} Eqn \ 6.14$$

• Suppose our neural network is constructed from a single layer of sigmoid units then we have

$$\frac{\partial h(x_i)}{\partial w_{jk}} = \sigma'(x_i)x_{ijk} = h(x_i)(1 - h(x_i))x_{ijk}$$

where x_{ijk} is the k^{th} input to unit j for the i^{th} training example, and $\sigma'(x)$ is the derivative of the sigmoid squashing function

• Finally, substituting this expression into *Eqn 6.14*, we obtain a simple expression for the derivatives that constitute the gradient

$$\frac{\partial G(h, D)}{\partial w_{jk}} = \sum_{i=1}^{m} (d_i - h(x_i)) x_{ijk}$$

• Because we seek to maximize rather than minimize P(D/h), we perform gradient ascent rather than gradient descent search. On each iteration of the search the weight vector is adjusted in the direction of the gradient, using the weight update rule

$$w_{jk} \leftarrow w_{jk} + \Delta w_{jk}$$

where,

$$\Delta w_{jk} = \eta \sum_{i=1}^{m} (d_i - h(x_i)) x_{ijk}$$

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→ Eqn 6.15

where η is a small positive constant that determines the step size of the gradient ascent search.

• Comparing this weight-update rule to the weight-update rule used by the Backpropagation algorithm we can get

$$w_{jk} \leftarrow w_{jk} + \Delta w_{jk}$$

where,

$$\Delta w_{jk} = \eta \sum_{i=1}^{m} h(x_i)(1 - h(x_i))(d_i - h(x_i)) x_{ijk}$$

Notice this is similar to the rule given in *Eqn 6.15* except for the extra term $h(x)(l - h(x_i))$, which is the derivative of the sigmoid function.

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Minimum Description Length Principle

- The Minimum Description Length principle is motivated by interpreting the definition of h_{MAP} in the light of basic concepts from information theory.
- Consider the definition of h_{MAP}

$$h_{MAP} = \operatorname*{argmax}_{h \in H} P(D|h) P(h)$$

which can be equivalently expressed in terms of maximizing the *log2*

$$h_{MAP} = \underset{h \in H}{\operatorname{argmax}} \log_2 P(D|h) + \log_2 P(h)$$

or alternatively, minimizing the negative of this quantity

$$h_{MAP} = \underset{h \in H}{\operatorname{argmin}} - \log_2 P(D|h) - \log_2 P(h) \longrightarrow Eqn \ 6.16$$

- *Eqn 6.16* can be interpreted as a statement that short hypotheses are preferred, assuming a particular representation scheme for encoding hypotheses and data.
- To explain this, let us take a basic result from information theory: Consider the problem of designing a code to transmit messages drawn at random, where the probability of encountering message i is p_{i} .
- We are interested here in the most compact code i.e., we are interested in the code that minimizes the expected number of bits we must transmit in order to encode a message drawn at random.

- Clearly, to minimize the expected code length we should assign shorter codes to messages that are more probable.
- Shannon and Weaver (1949) showed that the optimal code(i.e., the code that minimizes the expected message length) assigns $-log_2p_i$ bits to encode message *i*.
- The number of bits required to encode message *i* using code *C* will be referred as the *description length of message i with respect to C*, which is denoted as $L_c(i)$.
- Let us interpret *Eqn 6.16* in the perspective of the above result from coding theory
 - > $-log_2 P(h)$ is the description length of h under the optimal encoding for the hypothesis space H.

In our notation, $L_{C_H}(h) = -log_2 P(h)$, where C_H is the optimal code for hypothesis space H.

- → -log₂P(D/h) is the description length of the training data D given hypothesis h, under its optimal encoding. In our notation, $L_{C_{D/h}}(D/h) = -log_2P(D/h), \text{ where } C_{D/h} \text{ is the optimal code for describing data D assuming that both the sender and receiver know the hypothesis h.}$
- Therefore we can rewrite Eqn 6.16 to show that h_{MAP} is the hypothesis h that minimizes the sum given by the description length of the hypothesis plus the description length of the data given the hypothesis.

$$h_{MAP} = \underset{h}{\operatorname{argmin}} L_{C_H}(h) + L_{C_{D|h}}(D|h)$$

where C_H and $C_{D/h}$ are the optimal encodings for H and for D given h, respectively.

- The Minimum Description Length (MDL) principle recommends choosing the hypothesis that minimizes the sum of these two description lengths.
- To apply this principle in practice we must choose specific encodings or representations appropriate for the given learning task.
- Assuming we use the codes C_1 and C_2 to represent the hypothesis and the data given the hypothesis, we can state the MDL principle as

$h_{MDL} = \underset{h \in H}{\operatorname{argmin}} L_{C_1}(h) + L_{C_2}(D|h) \xrightarrow{} Eqn \ 6.17$

• The above analysis shows that if we choose C_1 to be the optimal encoding of hypotheses C_H , and if we choose C_2 to be the optimal encoding $C_{D/h}$ then $h_{MDL} = h_{MAP}$.

• Conclusions from MDL Principle

- Does MDL principle prove once and for all that short hypotheses are best?
- No. We have only shown that if a representation of hypotheses is chosen so that the size of hypothesis h is $-log_2P(h)$, and if a representation for exceptions is chosen so that the encoding length of D given h is equal to $-log_2P(D/h)$, then the MDL principle produces MAP hypotheses.

Naïve Bayes Classifier

- One highly practical Bayesian learning method is the naive Bayes learner, often called the Naive Bayes classifier.
- The Naive Bayes algorithm is a method that uses the **probabilities** of each attribute belonging to each class to make a prediction.
- In some domains its performance has been shown to be comparable to that of neural network and decision tree learning.

- The naive Bayes classifier applies to learning tasks where each instance *x* is described by a conjunction of attribute values and where the target function *f*(*x*) can take on any value from some finite set *V*.
- A set of training examples of the target function is provided, and a new instance is presented, described by the tuple of attribute values $\langle a_b, a_{2}...a_n \rangle$.

- The learner is asked to predict the target value, or classification, for this new instance.
- The Bayesian approach to classifying the new instance is to assign the most probable target value, v_{MAP} , given the attribute values $\langle a_1, a_2, ..., a_n \rangle$ that describe the instance

$$v_{MAP} = \operatorname*{argmax}_{v_j \in V} P(v_j | a_1, a_2 \dots a_n)$$

• We can use Bayes theorem to rewrite this expression as

$$v_{MAP} = \operatorname*{argmax}_{v_j \in V} \frac{P(a_1, a_2 \dots a_n | v_j) P(v_j)}{P(a_1, a_2 \dots a_n)}$$

=
$$\operatorname*{argmax}_{v_j \in V} P(a_1, a_2 \dots a_n | v_j) P(v_j) \longrightarrow Eqn \ 6.19$$

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- Now we could attempt to estimate the two terms in *Eqn 6.19* based on the training data.
- It is easy to estimate each of the $P(v_j)$ simply by counting the frequency with which each target value v_j occurs in the training data.
- However, estimating the different $P(a_b, a_2..., a_n/v_j)$ terms in this fashion is not feasible unless we have a very, very large set of training data.
- The problem is that the number of these terms is equal to the number of possible instances times the number of possible target values.
- Therefore, we need to see every instance in the instance space many times in order to obtain reliable estimates.

Assumption

The naive Bayes classifier is based on the simplifying assumption that *the attribute values are conditionally independent given the target value*.

• From this assumption it is possible to say that given the target value of the instance, the probability of observing the conjunction $a_1, a_2...a_n$ is just the product of probabilities for the individual attributes:

$$P(a_1, a_2, \dots, a_n/v_j) = \prod_i P(ai/vj)$$

Substituting this into **Eqn 6.19**, we have the approach used by the Naive Bayes classifier.

 $v_{NB} = \underset{v_j \in V}{\operatorname{argmax}} P(v_j) \prod_i P(a_i | v_j)$

→ Eqn 6.20

where v_{NB} denotes the target value output by the Naïve Bayes classifier.

- Thus, in a in a Naive Bayes classifier the number of distinct $P(a_i/v_j)$ terms that must estimated from the training data is just the number of distinct attribute values times the number of distinct target values- a much smaller number compared to estimating $P(a_1,a_2,...,a_n/v_j)$.
- To summarize, the naive Bayes learning method involves a learning step in which the various $P(v_j)$ and $P(a_i/v_j)$ terms are estimated, based on their frequencies over the training data.

- The set of these estimates corresponds to the learned hypothesis. This hypothesis is then used to classify each new instance by applying the rule in *Eqn 6.20*.
- Whenever the naive Bayes assumption of conditional independence is satisfied, this naive Bayes classification v_{NB} is identical to the MAP classification.
- One interesting difference between the naive Bayes learning method and other learning methods we have considered is that there is no explicit search through the space of possible hypotheses.

An Illustrative Example

- Let us apply the naive Bayes classifier to a concept learning problem we considered during our discussion of decision tree learning: classifying days according to whether someone will play tennis(*PlayTennis*).
- Table 3.2 provides a set of 14 training examples of the target concept *PlayTennis*, where each day is described by the attributes *Outlook*, *Temperature*, *Humidity*, and *Wind*.
- Here we use the naive Bayes classifier and the training data from this Table 3.2 to classify the following novel instance:

<Outlook = sunny, Temperature = cool, Humidity = high, Wind = strong>

Day	Outlook	Temperature	Humidity	Wind	PlayTennis
D1	Sunny	Hot	High	Weak	No
D2	Sunny	Hot	High	Strong	No
D3	Overcast	Hot	High	Weak	Yes
D4	Rain	Mild	High	Weak	Yes
D5	Rain	Cool	Normal	Weak	Yes
D6	Rain	Cool	Normal	Strong	No
D7	Overcast	Cool	Normal	Strong	Yes
D8	Sunny	Mild	High	Weak	No
D9	Sunny	Cool	Normal	Weak	Yes
D10	Rain	Mild	Normal	Weak	Yes
D11	Sunny	Mild	Normal	Strong	Yes
D12	Overcast	Mild	High	Strong	Yes
D13	Overcast	Hot	Normal	Weak	Yes
D14	Rain	Mild	High	Strong	No

Table 3.2: Training examples for the target concept *PlayTennis*
- Our task is to predict the target value (yes or no) of the target concept *PlayTennis* for this new instance.
- Instantiating *Eqn 6.20* to fit the current task, the target value VNB is given by

$$v_{NB} = \underset{v_j \in \{yes, no\}}{\operatorname{argmax}} P(v_j) \prod_i P(a_i | v_j)$$

$$= \underset{v_j \in \{yes, no\}}{\operatorname{argmax}} P(v_j) \qquad P(Outlook = sunny | v_j) P(Temperature = cool | v_j)$$

$$P(Humidity = high|v_j)P(Wind = strong|v_j) \quad (6.21)$$

Estimating probabilities of attribute values and target value from training data

Probabilities	of Target	Value(Play'	Fennis)
			/

PlayTennis		P(Yes)/P(No)	
Yes	9	9/14	
No	5	5/14	
Total	14		

Probabilities of Outlook Attribute values

	Yes	No	P(Yes)	P(No)
Sunny	2	3	2/9	3/5
Overcast	4	0	4/9	0/5
Rain	3	2	3/9	2/5
Total	9	5		

Probabilities of Temperature Attribute Values

	Yes	No	P(Yes)	P(No)
Hot	2	2	2/9	4/5
Mild	4	2	4/9	2/5
Cool	3	1	3/9	1/5
Total	9	5		

Probabilities of Humidity Attribute Values

	Yes	No	P(Yes)	P(No)
Normal	6	1	6/9	1/5
High	3	4	3/9	4/5
Total	9	5		

Probabilities of Wind Attribute Values

	Yes	No	P(Yes)	P(No)
Strong	3	3	3/9	3/5
Weak	6	2	6/9	2/5
Total	9	5		

• Using these probability estimates and similar estimates for the remaining attribute values, we calculate v_{NB} according to Eqn 6.21 as follows:

For
$$v_j = Yes$$

 $V_{NB} = P(Yes) P(Sunny/Yes) P(Cool/Yes) P(High/Yes) P(Strong/Yes)$
 $= 9/14 * 2/9 * 3/9 * 3/9 * 3/9$
 $= 0.00529$

For $v_j = No$ $V_{NB} = P(No) P(Sunny/No) P(Cool/No) P(High/No) P(Strong/No)$ = 5/14 * 3/5 * 1/5 * 4/5 * 3/5= 0.020571

• Thus, the naive Bayes classifier assigns the target value *PlayTennis* = *No* to this new instance, based on the probability estimates learned from the training data.

RID	Age	Income	Student	Credit_Rating	Buys_Computer
1	Youth	High	No	Fair	No
2	Youth	High	No	Excellent	No
3	Middle_aged	High	No	Fair	Yes
4	Senior	Medium	No	Fair	Yes
5	Senior	Low	Yes	Fair	Yes
6	Senior	Low	Yes	Excellent	No
7	Middle_aged	Low	Yes	Excellent	Yes
8	Youth	Medium	No	Fair	No
9	Youth	Low	Yes	Fair	Yes
10	Senior	Medium	Yes	Fair	Yes
11	Youth	Medium	Yes	Excellent	Yes
12	Middle_aged	Medium	No	Excellent	Yes
13	Middle_aged	High	Yes	Fair	Yes
14	Senior	Medium	No	Excellent	No

 Table 3.2: Training examples for the target concept Buys_Computer

Estimating Probabilities

- Till now, we have estimated probabilities by the fraction of times the event is observed to occur over the total number of opportunities.
- For ex : we estimated P(Wind = strong/Play Tennis = no) by the fraction \$\frac{n_c}{n}\$ where \$n = 5\$ is the total number of training examples for which PlayTennis = no, and \$n=3\$ is the number of these for which Wind = strong.
- While this observed fraction provides a good estimate of the probability in many cases, it provides poor estimates when n_c is very small.
- To see the difficulty, for time being let us imagine that the value of P(Wind = strong | PlayTennis = no) is .08 and that we have a sample containing only 5 examples for which *PlayTennis = no*. Machine Learning-15C573 83

- Then the most probable value for n_c is 0 which raises two difficulties:
 - *i.* $\frac{n_c}{n}$ produces a biased underestimate of the probability.
 - ii. when this probability estimate is zero, this probability term will dominate the Bayes classifier if the future query contains *Wind = strong*.
- To avoid this difficulty we can adopt a Bayesian approach to estimating the probability, using the *m-estimate* defined as follows:

$$\frac{n_c + mp}{n + m} \longrightarrow Eqn \ 6.22$$

- Here, *n_c* and *n* are defined as before, *p* is our prior estimate of the probability we wish to determine and *m* is a constant called the *equivalent sample size* which determines how heavily to weight *p* relative to the observed data.
- A typical method for choosing *p* in the absence of other information is to assume uniform priors, i.e., if an attribute has *k* possible values we set $p = \frac{1}{k}$.
- For ex: in estimating P(Wind = Strong | PlayTennis = no) we note the attribute *Wind* has two possible values, so uniform priors would correspond to choosing p = 0.5.
- Note that in Eqn 6.22 if m is zero, then m-estimate is equivalent to the simple fraction $\frac{n_c}{n}$.

- If both *n* and *m* are nonzero, then the observed fraction $\frac{n_c}{n}$ and prior *p* will be combined according to the weight *m*.
- The reason *m* is called the *equivalent sample size* is that *Eqn 6.22* can be interpreted as augmenting the *n* actual observations by an additional *m* virtual samples distributed according to *p*.

Bayesian Belief Networks

- The naive Bayes classifier makes significant use of the assumption that the values of the attributes $a_1 \, \ldots \, a_n$ are conditionally independent given the target value v.
- This assumption dramatically reduces the complexity of learning the target function.
- When it is met, the naive Bayes classifier outputs the optimal Bayes classification. However, in many cases this conditional independence assumption is clearly overly restrictive.
- A Bayesian belief network describes the probability distribution governing a set of variables by specifying a set of conditional independence assumptions along with a set of conditional probabilities.

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- In contrast to the *naive Bayes classifier*, which assumes that all the variables are conditionally independent given the value of the target variable, *Bayesian belief networks* allow stating conditional independence assumptions that apply to subsets of the variables.
- Thus, Bayesian belief networks provide an intermediate approach that is less constraining than the global assumption of conditional independence made by the naive Bayes classifier.
- Bayesian belief networks more tractable than avoiding conditional independence assumptions altogether.
- In general, a Bayesian belief network describes the probability distribution over a set of variables. Consider an arbitrary set of random variables $Y_1 \ldots Y_n$, where each variable Y_i can take on the set of possible values $V(Y_i)$.

- We define the joint space of the set of variables Y to be the cross product $V(Y_l) \times V(Y_2) \times \ldots V(Y_n)$. Each item in the joint space corresponds to one of the possible assignments of values to the tuple of variables $\langle Y_l, \ldots, Y_n \rangle$.
- The probability distribution over this joint space is called the *joint probability distribution*.
- A Bayesian belief network describes the joint probability distribution for a set of variables.

Conditional Independence

• Let X, Y, and Z be three discrete-valued random variables. We say that X is conditionally independent of Y given Z if the probability distribution governing X is independent of the value of Y given a value for Z i.e., if

$$(\forall x_i, y_j, z_k) P(X = x_i | Y = y_j, Z = z_k) = P(X = x_i | Z = z_k)$$

where $x_i \in V(X)$, $y_i \in V(Y)$, and $z_k \in V(Z)$.

- We commonly write the above expression in abbreviated form as
 P(X / Y, Z) = P(X / Z). This definition of conditional independence can be extended to sets of variables as well.
- We say that the set of variables $X_1 \ldots X_l$ is conditionally independent of the set of variables $Y_l \ldots Y_m$ given the set of variables $Z_1 \ldots Z_n$ if Machine Learning-15CS73 90

 $P(X_1 \ldots X_l | Y_1 \ldots Y_m, Z_1 \ldots Z_n) = P(X_1 \ldots X_l | Z_1 \ldots Z_n)$

• The correspondence can be drawn between this definition and our use of conditional ,independence in the definition of the naive Bayes classifier.

Bayesian Belief Network Representation

- A Bayesian belief network (Bayesian network for short) represents the joint probability distribution for a set of variables.
- For example, the Bayesian network in *figure 6.3.1* represents the joint probability distribution over the boolean variables *Storm*, *Lightning*, *Thunder*, *ForestFire*, *Campfire*, *and BusTourGroup*.
- In general, a Bayesian network represents the joint probability distribution by specifying a set of conditional independence assumptions (represented by a directed acyclic graph), together with sets of local conditional probabilities.
- *Each variable in the joint space is represented by a node* in the Bayesian network

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- *Each variable in the joint space is represented by a node* in the Bayesian network



Figure 6.3.1: Bayesian Belief Network

- For each variable two types of information are specified.
 - i. The network arcs represent the assertion that the variable is conditionally independent of its nondescendants in the network given its immediate predecessors in the network. We say X is a descendant of Y if there is a directed path from Y to X.
 - ii. A conditional probability table is given for each variable, describing the probability distribution for that variable given the values of its immediate predecessors.
- The joint probability for any desired assignment of values $\langle y_1, \ldots, y_n \rangle$ to the tuple of network variables $\langle Y_1 \ldots Y_n \rangle$ can be computed by the formula

$$P(y_1,\ldots,y_n) = \prod_{i=1}^n P(y_i | Parents(Y_i))$$

where $Parents(Y_i)$ denotes the set of immediate predecessors of Y_i in the network.

- Let us illustrate the Bayesian network given in figure 6.3.1 which represents the joint probability distribution of boolean variables *Storm, Lightning, Thunder, ForestFire, Campfire, and BusTourGroup*.
- Consider the node *Campfire*. The network nodes and arcs represent the assertion that *Campfire* is conditionally independent of its nondescendants *Lightning* and *Thunder*, given its immediate parents *Storm* and *BusTourGroup*.
- This means that once we know the value of the variables *Storm* and *BusTourGroup*, the variables *Lightning* and *Thunder* provide no additional information about *Campfire*.

The figure 6.3.2 below shows the conditional probability table associated with the variable *Campfire*.



Figure 6.3.2: The conditional Probability Table for Campfire node

- The top left entry in this table, for ex: expresses the assertion that P(Campfire=True/Storm=True, BusTourGroup=True)=0.4
- Note this table provides only the conditional probabilities of *Campfire* given its parent variables *Storm* and *BusTourGroup*. 11-11-2019

- The set of local conditional probability tables for all the variables, together with the set of conditional independence assumptions described by the network, describe the full joint probability distribution for the network.
- One attractive feature of Bayesian belief networks is that they allow a convenient way to represent causal knowledge such as the fact that *Lightning* causes *Thunder*.

Learning in Bayesian Belief Networks

- Can we devise effective algorithms for learning Bayesian belief networks from training data?
- Several different settings for this learning problem can be considered.
 - i. First, the network structure might be given in advance, or it might have to be inferred from the training data.
 - ii. Second, all the network variables might be directly observable in each training example, or some might be unobservable.
- In the case where the network structure is given in advance and the variables are fully observable in the training examples, learning the conditional probability tables is straightforward.

- We simply estimate the conditional probability table entries just as we would for a naive Bayes classifier.
- In the case where the network structure is given but only some of the variable values are observable in the training data, the learning problem is more difficult.
- This problem is somewhat analogous to learning the weights for the hidden units in an artificial neural network, where the input and output node values are given but the hidden unit values are left unspecified by the training examples.
- Russell et al.(1995) proposed a gradient ascent procedure that learns entries in conditional probability tables.
- This gradient ascent procedure searches through a space of hypotheses that corresponds to the set of all possible entries for the **conditional** probability tables in Learning-15CS73 103

Learning Structure of Bayesian Networks

- Learning Bayesian networks when the network structure is not known in advance is also difficult.
- Cooper and Herskovits (1992) present a Bayesian scoring metric for choosing among alternative networks.
- They also present a heuristic search algorithm called *K2* for learning network structure when the data is fully observable.
- Constraint-based approaches to learning Bayesian network structure have also been developed

The EM Algorithm

- In many practical learning settings, only a subset of the relevant instance features might be observable.
- For ex : in training our or using the Bayesian belief network we might have data where only a subset of the network variables *Storm*, *Lightning*, *Thunder*, *ForestFire*, *Campfire*, and *BusTourGroup* have been observed.
- Many approaches have been proposed to handle the problem of learning in the presence of unobserved variables.
- The EM algorithm (*Dempster et al. 1977*), a widely used approach for learning in the presence of unobserved variables.

- The EM algorithm can be used even for variables whose value is never directly observed, provided the general form of the probability distribution governing these variables is known.
- The EM algorithm has been used to train Bayesian belief networks as well as radial basis function networks.
- The EM algorithm is also the basis for many unsupervised clustering algorithms.

Estimating Means of *k* **Gaussians**

- Consider a problem in which the data D is a set of instances generated by a probability distribution that is a mixture of k distinct Normal distributions.
- This problem setting is illustrated in *figure 6.4* for the case where *k* = 2 and where the instances are the points shown along the *x*-axis.
- Each instance is generated using a two-step process.
 - i. One of the *k* Normal distributions is selected at random.
 - ii. A single random instance x_i is generated according to this selected distribution.

This process is repeated to generate a set of data points as shown in *figure 6.4*

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Figure 6.4 : Instances generated by a mixture of two Normal distributions with identical variance σ

- Let us consider a special case, where the selection of the single Normal distribution at each step is based on choosing each with uniform probability, where each of the k Normal distributions has the same variance σ^2 .
- The learning task is to output a hypothesis $h = \langle \mu_1 \dots \mu_k \rangle$ that describes the means of each of the *k* distributions.
- This task involves finding a maximum likelihood hypothesis for these means; i.e., a hypothesis *h* that maximizes *p*(*D*/*h*).
- It is easy to calculate the maximum likelihood hypothesis for the mean of a single Normal distribution given the observed data instances x_1, x_2, \ldots, x_m drawn from this single distribution

• Restating *Eqn 6.6* using our current notation, we have

$$\mu_{ML} = \underset{\mu}{\operatorname{argmin}} \sum_{i=1}^{m} (x_i - \mu)^2 \longrightarrow Eqn \ 6.27$$

• In this case, the sum of squared errors is minimized by the sample mean

$$\mu_{ML} \doteq \frac{1}{m} \sum_{i=1}^{m} x_i \longrightarrow Eqn \ 6.28$$

Necessity for EM Algorithm

- Our problem involves a mixture of k different Normal distributions, and we cannot observe which instances were generated by which distribution.
- Thus, we have a prototypical example of a problem involving hidden variables.
- In the example of *figure 6.4* we can think of the full description of each instance as the triple $\langle x_i, z_{i1}, z_{i2} \rangle$, where x_i is the observed value of the *i*th instance and where z_{i1} and z_{i2} indicate which of the two Normal distributions was used to generate the value x_i .
- Here x_i is the observed variable in the description of the instance, and z_{i1} and z_{i2} are hidden variables.

- If the values of z_{i1} and z_{i2} were observed, we could use *Eqn 6.27* to solve for the means μ_1 and μ_2 . Because they are not, we will instead use the EM algorithm.
- Applied to our k-means problem the EM algorithm searches for a maximum likelihood hypothesis by repeatedly re-estimating the expected values of the hidden variables z_{ij} given its current hypothesis <µ₁....µ_k> then recalculating the maximum likelihood hypothesis using these expected values for the hidden variables.

Describing an instance of EM algorithm

- Applied to the problem of estimating the two means for figure 6.4, the EM algorithm first initializes the hypothesis to $h = \langle \mu_1, \mu_2 \rangle$, where μ_1 and μ_2 are arbitrary initial values.
- It then iteratively re-estimates *h* by repeating the following two steps until the procedure converges to a stationary value for *h*.

- **Step 1**: Calculate the expected value $E[z_{ij}]$ of each hidden variable z_{ij} , assuming the current hypothesis $h = \langle \mu_1, \mu_2 \rangle$ holds.
- Step 2: Calculate a new maximum likelihood hypothesis $h' = \langle \mu_1 \rangle$; $\mu_2 \rangle$, assuming the value taken on by each hidden variable z_{ij} is its expected value $E[z_{ij}]$ calculated in *Step 1*. Then replace the hypothesis $h = \langle \mu_1, \mu_2 \rangle$ by the new hypothesis $h' = \langle \mu_1, \mu_2 \rangle$ and iterate.

Implementation of steps in practice

• Step 1 must calculate the expected value of each z_{ij} . This $E[z_{ij}]$ is just the probability that instance x_i was generated by the j^{th} Normal distribution
$$E[z_{ij}] = \frac{p(x = x_i | \mu = \mu_j)}{\sum_{n=1}^{2} p(x = x_i | \mu = \mu_n)}$$
$$= \frac{e^{-\frac{1}{2\sigma^2}(x_i - \mu_j)^2}}{\sum_{n=1}^{2} e^{-\frac{1}{2\sigma^2}(x_i - \mu_n)^2}}$$

- Thus the first step is implemented by substituting the current values $\langle \mu_l, \mu_2 \rangle$ and the observed x_i into the above expression.
- In the second step we use the $E[z_{ij}]$ calculated during Step 1 to derive a new maximum likelihood hypothesis $h' = \langle \mu_1, \mu_2 \rangle$. The maximum likelihood hypothesis in this case is given by

$$\mu_j \leftarrow \frac{\sum_{i=1}^m E[z_{ij}] \ x_i}{\sum_{i=1}^m E[z_{ij}]}$$

• Our new expression is just the weighted sample mean for μ_j , with each instance weighted by the expectation $E[z_{ij}]$ that it was generated by the *j*th Normal distribution.

General Statement of EM Algorithm

- The EM algorithm can be applied in many settings where we wish to estimate some set of parameters θ that describe an underlying probability distribution, given only the observed portion of the full data produced by this distribution.
- In the two-means example the parameters of interest were $\theta = \langle \mu_1, \mu_2 \rangle$, and the full data were the triples $\langle x_i, z_{i1}, z_{i2} \rangle$ of which only the x_i were observed.
- In general let $X = \{x_b, \ldots, x_m\}$ denote the observed data in a set of *m* independently drawn instances, let $Z = \{z_1, \ldots, z_m\}$ denote the unobserved data in these same instances, and let Y = X U Z denote the full data.

- The unobserved Z can be treated as a random variable whose probability distribution depends on the unknown parameters θ and on the observed data X.
- Similarly , *Y* is a random variable because it is defined in terms of the random variable *Z*.
- We use h to denote the current hypothesized values of the parameters θ , and h' to denote the revised hypothesis that is estimated on each iteration of the EM algorithm.
- The EM algorithm searches for the maximum likelihood hypothesis *h*' by seeking the *h*' that maximizes *E[ln P(Y / h')]*.

- Let us consider exactly what this expression signifies
 - i. First, P(Y | h') is the likelihood of the full data Y given hypothesis h'. It is reasonable that we wish to find a h' that maximizes some function of this quantity.
 - ii. Second, maximizing the logarithm of this quantity $\ln P(Y | h')$ also maximizes P(Y | h').
 - iii. Third, we introduce the expected value $E[\ln P(Y | h')]$ because the full data Y is itself a random variable.
- Given that the full data *Y* is a combination of the observed data *X* and unobserved data *Z*, we must average over the possible values of the unobserved *Z*, weighting each according to its probability.
- In other words we take the expected value *E[ln P(Y / h')]* over the probability distribution governing the random variable *Y*.

• Let us define a function Q(h'|h) that gives E[ln P(Y | h')] as a function of h', under the assumption that $\theta = h$ and given the observed portion X of the full data Y.

$Q(h'|h) = E[\ln p(Y|h')|h, X]$

• In its general form, the EM algorithm repeats the following two steps until convergence:

Step 1: *Estimation (E) step:*

Calculate Q(h' | h) using the current hypothesis h and the observed data X to estimate the probability distribution over Y.

$Q(h'|h) = E[\ln p(Y|h')|h, X]$

Step 2: Maximization(M) step:

Replace hypothesis h by the hypothesis h' that maximizes this Q function.

$h \leftarrow \operatorname*{argmax}_{h'} Q(h'|h)$

When the function Q is continuous, the EM algorithm converges to a stationary point of the likelihood function P(Y/h').

Derivation of k Means Algorithm

- To illustrate the general EM algorithm, let us use it to derive the algorithm for estimating the means of a mixture of k Normal distributions.
- In the *k*-means objective is to estimate the parameters $\theta = \langle \mu_l ... \mu_k \rangle$ that define the means of the *k* Normal distributions.
- We are given the observed data $X = \{\langle x_i \rangle\}$. The hidden variables $Z = \{\langle z_{i1}, \ldots, z_{ik} \rangle\}$ in this case indicate which of the k Normal distributions was used to generate x_i .
- To apply EM we must derive an expression for Q(h'/h) that applies to our k-means problem.
- First, let us derive an expression for ln p(Y/h')11-11-2019 Machine Learning-15CS73

• The probability $p(y_i / h')$ of a single instance $y_i = \langle x_i, z_{i1}, \dots, z_{ik} \rangle$ if the full data can be written as

$$p(y_i|h') = p(x_i, z_{i1}, \dots, z_{ik}|h') = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2\sigma^2} \sum_{j=1}^k z_{ij} (x_i - \mu'_j)^2}$$

• Given this probability for a single instance $p(y_i \mid h')$, the logarithm of the probability $ln P(Y \mid h')$ for all *m* instances in the data is

$$\ln P(Y|h') = \ln \prod_{i=1}^{m} p(y_i|h')$$

= $\sum_{i=1}^{m} \ln p(y_i|h')$
= $\sum_{i=1}^{m} \left(\ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} \sum_{j=1}^{k} z_{ij} (x_i - \mu'_j)^2 \right)$

Finally we must take the expected value of this *ln P(Y \ h [']*) over the probability distribution governing *Y*. The above expression for *ln P(Y \ h [']*) is a linear function of these *z_{ij}*. In general, for any function *f(z)* that is a linear function of *z*, the following equality holds

$$E[f(z)] = f(E[z])$$

• This general fact about linear functions allows us to write

$$E[\ln P(Y|h')] = E\left[\sum_{i=1}^{m} \left(\ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} \sum_{j=1}^{k} z_{ij} (x_i - \mu'_j)^2\right)\right]$$
$$= \sum_{i=1}^{m} \left(\ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} \sum_{j=1}^{k} E[z_{ij}] (x_i - \mu'_j)^2\right)$$

• To summarize, the function Q(h'/h) for the k means problem is

$$Q(h'|h) = \sum_{i=1}^{m} \left(\ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} \sum_{j=1}^{k} E[z_{ij}](x_i - \mu'_j)^2 \right)$$

where $h' = \langle \mu_1', \ldots, \mu_k' \rangle$ and where $E[z_{ij}]$ is calculated based on the current hypothesis h and observed data X. We know that

$$E[z_{ij}] = \frac{e^{-\frac{1}{2\sigma^2}(x_i - \mu_j)^2}}{\sum_{n=1}^k e^{-\frac{1}{2\sigma^2}(x_i - \mu_n)^2}} \longrightarrow Eqn \ 6.29$$

Thus, the first (estimation) step of the EM algorithm defines the Q function based on the estimated $E[z_{ij}]$ terms.

• The second (maximization) step then finds the values $\mu_1' \dots \mu_k'$ that maximize this Q function. In the current case

$$\operatorname{argmax}_{h'} Q(h'|h) = \operatorname{argmax}_{h'} \sum_{i=1}^{m} \left(\ln \frac{1}{\sqrt{2\pi\sigma^2}} - \frac{1}{2\sigma^2} \sum_{j=1}^{k} E[z_{ij}](x_i - \mu'_j)^2 \right)$$
$$= \operatorname{argmin}_{h'} \sum_{i=1}^{m} \sum_{j=1}^{k} E[z_{ij}](x_i - \mu'_j)^2 \qquad (Eqn \ 6.30)$$

Thus, the maximum likelihood hypothesis here minimizes a weighted sum of squared errors, where the contribution of each instance x_i to the error that defines μ_i 'is weighted by $E[z_{ij}]$

• The quantity given by *Eqn 6.30* is minimized by setting each μ_j to the weighted sample mean

$$\mu_j \leftarrow \frac{\sum_{i=1}^m E[z_{ij}] \quad x_i}{\sum_{i=1}^m E[z_{ij}]} \xrightarrow{} Eqn \ 6.31$$