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Department of Computer Science & Engineering

Course: Artificial Intelligence Machine Learning (18CS71) CSE HIT Nidasoshi Module'4: Bayesian Learning

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Bayesian Classification: Why? - Features

- Each observed training example can incrementally decrease or increase the estimated probability that a hypothesis is correct. This provides a more flexible approach to learning than 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. In Bayesian learning, prior knowledge is provided by asserting
 - 1. a prior probability for each candidate hypothesis, and

2. a probability distribution over observed data for each possible hypothesis. Dr. Mahesh G. Huddar, Department of CSE, HSIT, Nidasoshi

Bayesian Classification: Why? - Features

- Bayesian methods can accommodate hypotheses that make probabilistic predictions (e.g., 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.

Practical Difficulty

- Bayesian methods typically require initial knowledge of many probabilities. When these probabilities are not known in advance they are often estimated based on background knowledge, previously available data, and assumptions about the form of the underlying distributions.
- Computational cost required to determine the Bayes optimal hypothesis in the general case.

- Suppose a XYZ company needs to predict the service required by the incoming customer.
- If there are only two services offered R and M.
- Then the value to be predicted is whether the next customer will be for R or M.
- The number of classes k are 2 that is, k=2.

- The first step is to compute prior probability.
- Suppose the data gathered for the last one year showed that during that

period there were 2500 customers for R and 1500 customers for M.

- Thus, prior probability for the next customer to be for R is 2500/4000 or 5/8.
- Prior probability for the next customer to be for M is 1500/4000 or 3/8.
- Based on this information alone, the next customer would likely be for R.

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- Another way to predict the service requirement by the next customer is to look at the most recent data.
- One can look at the last few (choose a number) customers, to predict the next

customer. CSE, HIT, Nidasoshi

- Suppose the last five customers were for the services ... R, M, R, M, M order.
- Thus, the data shows the recent probability of R is 2/5 and that of M is 3/5.
- Based on just this information, the next customer will likely to be for M.

- Thomas Bayes suggested that the prior probability should be informed by the more recent data.
- Naive-Bayes posterior probability for a class is computed by multiplying the prior probability and the recent probability.
- NB posterior probability P(R) is (5/8 x 2/5) = 10/40.
- Similarly, the NB probability P(M) is $(3/8 \times 3/5) = 9/40$.
- Since P(R) is greater than P(M), it follows that there is a greater probability of the next customer to be for R.
- Thus the expected class label assigned to the next customer would be R.

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- Suppose, however the next customer coming in was for M service.
- The last five customer sequence now becomes M, R, M, M, M.
- Thus, the recent data shows the probability for R to be 1/5 and that of M to be 4/5.
- Now the NB probability for R is (5/8 x 1/5) = 5/40.
- Similarly, the NB probability for M is $(3/8 \times 4/5) = 12/40$.
- Since P(M) is greater than P(R), it follows that there is a greater probability of the next customer to be for M.
- Thus the expected class label assigned to the next customer is M. Dr. Mahesh G. Huddar, Department of CSE, HSIT, Nidasoshi

- In machine learning we are often interested in determining the best hypothesis from some space H, given the observed training data D.
- One way to specify what we mean by the best hypothesis is to say that we demand the most probable hypothesis, given the data D plus any initial knowledge about the prior probabilities of the various hypotheses in H.
- Bayes theorem provides a direct method for calculating such probabilities.
- More precisely, 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.

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- To define Bayes theorem precisely, let us first introduce a little notation.
- We shall write P(h) to denote the initial probability that hypothesis h holds, before we have observed the training data. P(h) is often called the priorprobability of h and may reflect any background knowledge we have about the chance that h is a correct hypothesis.
- If we have no such prior knowledge, then we might simply assign the same prior probability to each candidate hypothesis.

- **P(D) to denote the prior probability that training data D will be observed (i.e.,** the probability of D given no knowledge about which hypothesis holds).
- Next, we will write P(D|h) to denote the probability of observing data D given some world in which hypothesis h holds. More generally, we write P(x|y) to denote the probability of x given y.
 In machine learning problems we are interested in the probability P (h|D) that h
- In machine learning problems we are interested in the probability P (h|D) that h holds given the observed training data D.
- P (h|D) is called the posterior probability of h, because it reflects our confidence that h holds after we have seen the training data D.
- Notice the posterior probability P(h|d) reflects the influence of the training data
 D, in contrast to the prior probability P(h), which is independent of 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).

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

- The learner considers some set of candidate hypotheses H and is interested in finding the most probable hypothesis h ∈ 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.

• More precisely, we will say that h_{MAP} is a MAP hypothesis provided



 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(hi) = P(hj) for all hi and hj in H).

$CSE, h_{ML} = \underset{h \in H}{\operatorname{argmax}} P(D|h)$ shi

Example - Does patient have cancer or not?

- 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).
- 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.
- Furthermore we know that, 0.008 of the entire population have cancer.
- P(cancer) = 0.008 $P(\exists cancer) = 0.992$
- P(+|cancer) = 0.98 P(-|cancer) = 0.02
- P(+||cancer) = 0.03 P(-||cancer) = 0.97

Example - Does patient have cancer or not?

- 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?

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

P(cancer|+) = P(+|cancer) * P(cancer) = 0.98 * 0.008 = 0.0078P(|cancer|+) = P(+||cancer|) * P(|cancer|) = 0.03 * 0.992 = 0.0298

$$h_{MAP} = \exists cancer$$

Example - Does patient have cancer or not?

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

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

P(cancer|-) = P(-|cancer|) * P(cancer) = 0.02 * 0.008 = 0.00016P(|cancer|-) = P(-||cancer|) * P(|cancer|) = 0.97 * 0.992 = 0.96224 $h_{MAP} = |cancer|$

- Assume the learner considers some finite hypothesis space H defined over the instance space X.
- Here the task is to learn some target concept $c : X \rightarrow \{0,1\}$.
- As usual, we assume that the learner is given some sequence of training examples (<x1, d1>, <x2, d2>, <x3, d3>, . . <xm, dm>,) where xi is some instance from X and where di is the target value of xi.
- To simplify the discussion in this section, we assume the sequence of instances (x1...xm) is held fixed, so that the training data D can be written simply as the sequence of target values D = (d1...Dm)

BRUTE-FORCE MAP LEARNING algorithm

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- 1. For each hypothesis h in H, calculate the posterior probability
- $CSE, HIT, Nidas P(h|D) = \frac{P(D|h)P(h)}{das B}$ 2. Output the hypothesis h_{MAP} with the highest posterior probability

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

BRUTE-FORCE MAP LEARNING algorithm

• This algorithm may require significant computation, because it applies Bayes

theorem to each hypothesis in H to calculate P(h|D).

- While this is impractical for large hypothesis spaces,
- The algorithm is still of interest because it provides a standard against which we may judge the performance of other concept learning algorithms.

- BRUTE-FORCE MAP LEARNING algorithm
- Brute Force MAP learning algorithm must specify values for P(h) and P(D|h).
- P(h) and P(D|h) must be chosen to be consistent with the assumptions:

1. The training data Dis noise free. T, Nidasoshi

2. The target concept 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**.
- 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

- What choice shall we make for P(D|h)?
- P(D|h) is the probability of observing the target values D = <d1...dm> for the fixed set of instances <X1...Xm>.
- Since we assume noise-free training data, the probability of observing classification di given h is just 1 if di = h(xi) and 0 if di != h(xi).
- Therefore, $P(D|h) = \begin{cases} 1 \text{ if } d_i = h(x_i) \text{ for all } d_i \text{ in } D \\ 0 \text{ otherwise} \end{cases}$
- In other words, the probability of data **D given hypothesis h is 1 if D is consistent**
- with h, and 0 otherwise.

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- 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 agorithm.
- 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.

$$CSE, P(h|D) =, \frac{P(D|h)P(h)}{P(D)} \text{soshi}$$

 First consider the case where h is inconsistent with the training data D. We know that 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.

 Now consider the case where h is consistent with D. we know that P(Dlh) to be 1 when h is consistent with D, we have





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

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

- Many learning approaches such as neural network learning, linear regression, and polynomial curve fitting try to learn a continuous-valued target function.
- 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.
- The significance of this result is that it provides a Bayesian justification (under certain assumptions) for many neural network and other curve fitting methods that attempt to minimize the sum of squared errors over the training data.

- Learner L considers an instance space X and a hypothesis space H consisting of some class of real-valued functions defined over X.
- The problem faced by L is to learn an unknown target function f 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
- Each training example is a pair of the form (x_i, d_i) 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.

• In order to find the maximum likelihood hypothesis, we start with our earlier

definition but using lower case **p** to refer to the probability density function.

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

- We assume a fixed set of training instances (x₁...x_m) and therefore consider the data
 D to be the corresponding sequence of target values D = (d₁...d_m).
- Here we can write p(D|h) as the product of the various $p(d_i|h)$

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



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

$$= \operatorname*{argmax}_{h \in H} \prod_{i=1}^{m} \frac{1}{\sqrt{2\pi\sigma^{2}}} e^{-\frac{1}{2\sigma^{2}}(d_{i}-h(x_{i}))^{2}}$$

- 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 In p also maximizes p.*

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

• First term is constant, discard it.

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

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

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

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

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

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- The maximum likelihood hypothesis h_{ML} is the one that minimizes the sum of the squared errors between observed training values d_i and 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.
- Similar derivations can be performed starting with other assumed noise distributions, producing different results.
- Why is it reasonable to choose the Normal distribution to characterize noise?
 - One reason, is that it allows for a mathematically straightforward analysis.
 - A second reason is that the smooth, bell-shaped distribution is a good approximation to many types of noise in physical systems.
- Minimizing the sum of squared errors is a common approach in many neural network, curve fitting, and other approaches to approximating real-valued functions.
- The Minimum Description Length principle is motivated by interpreting the definition of h_{Map} in the light of basic concepts from information theory.
- Consider definition of h_{Map}

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

• 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

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

- To explain this, let us introduce 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 pi.**
- We are interested here in the most compact code; that is, 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.
 CSE, HIT, Nidasoshi
- 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 -log₂pi bits to encode message i.
- We will refer to the number of bits required to encode message i using code C as the description length of message i with respect to C, which we denote by L_c(i).

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- Let us interpret h_{MAP} Equation in light of the above result from coding theory.
- log₂P(h) is the description length of h under the optimal encoding for the hypothesis space H.
- In other words, this is the size of the description of hypothesis h using this optimal representation.
 CSE, HIT, Nidasoshi
- 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, $L_{CD|h}(Dlh) = -\log_2 P(Dlh)$, where $C_{D|h}$ is the optimal code for describing data D assuming that both the sender and receiver know the hypothesis h.

• Therefore we can rewrite Equation 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.

$$h_{MAP} = \underset{h}{\operatorname{argmin}} L_{C_{H}}(h) + L_{C_{DM}}(Dh) SOSN$$

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

- Minimum Description Length principle:
- Choose *h_{MDL}* where

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

CSE, $h \in H$, Nidasoshi

• 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_{DIh} then $h_{MDL} = h_{MAP}$

NAIVE BAYES CLASSIFIER

 The Bayesian approach to classifying the new instance is to assign the most probable target value, v_{MAP} given the attribute values <a1, a2...an> 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

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

Naive Bayes classifier:

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

NAIVE BAYES CLASSIFIER

- One highly practical Bayesian learning method is the naive Bayes learner, often called the *naive Bayes classijier*.
- 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 <al, a2...an>.
- The learner is asked to predict the target value, or classification, for this new

Instance. Dr. Mahesh G. Huddar, Department of CSE, HSIT, Nidasoshi

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

- Here there are 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 to classify the following novel instance:
 (Outlook = sunny, Temperature = cool, Humidity = high, Wind = strong)

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

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

 $P(Humidity = high|v_j)P(Wind = strong|v_j)$

Outlook	Р	Ν		Humidity	Ρ	Ν
sunny	2/9	3/5		high	3/9	4/5
overcast	4/9	0		normal	6/9	1/5
rain CC	3/9	2/5	NI	daeo	shi	
Tempreature	∟,			Windy		
hot	2/9	2/5		true	3/9	3/5
m ild	4/9	2/5		false	6/9	2/5
cool	3/9	1/5				

$$P(PlayT ennis = yes) = 9/14 = .64$$
$$P(PlayT ennis = no) = 5/14 = .36$$

$$P(Wind = strong | PlayTennis = yes) = 3/9 = .33$$

$$P(Wind = strong | PlayTennis = no) = 3/5 = 60$$

 $v_{NB} = P(yes) P(sunny|yes) P(cool|yes) P(high|yes) P(strong|yes) = .0053$

 $v_{NB} = P(no) P(sunny|no) P(cool|no) P(high|no) P(strong|no) = .0206$

 Thus, the naive Bayes classifier assigns the target value *PlayTennis = no* to this new instance

- Estimate conditional probabilities of each attributes {color, legs, height, smelly} for the species classes: {M, H} using the data given in the table.
- Using these probabilities estimate the probability values for the new instance (Color=Green, legs=2, Height=Tall, and Smelly=No).

No	Color	Legs	Height	Smelly	Species
1	White	3	Short	Yes	M
2	Green	2	Tall	No	М
3	Green	3	Short	Yes	М
4	White	3	Short	Yes	М
5	Green	2	Short	No	Н
6	White	2	Tall	No	Н
7	White	2	Tall	No	Н
8	White	2	Short	Yes	Н

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No	Color	Legs	Height	Smelly	Species
1	White	3	Short	Yes	М
2	Green	2	Tall	No	М
3	Green	3	Short	Yes	М
4	White	3	Short	Yes	М
5	Green	2	Short	No	Н
6	White	2	Tall	No	Н
7	White	2	Tall	No	Н
8	White	2	Short	Yes	Н
CSE, H					

NAIVE BAYES CLASSIFIER EXAMPLE - 2

$$P(M) = \frac{4}{8} = 0.5 \quad P(H) = \frac{4}{8} = 0.5$$

Color	М	Н
White	2/4	3/4
Green	2/4	1/4

Legs	М	Н
2	1/4	4/4
3	3/4	0/4

New Instance

(Color=Green, legs=2, Height=Tall, and Smelly=No)

Height	М	Н
Tall	3/4	2/4
Short	1/4	2/4

Smelly	М	Н
Yes	3/4	1/4
No	1/4	3/4

NAIVE BAYES CLASSIFIER - EXAMPLE - 2

$$P(M) = \frac{4}{8} = 0.5 \quad P(H) = \frac{4}{8} = 0.5$$

Color	М	Н	Legs	М	н	Height	М	Н	Smelly	М	н
White	2/4	3/4	2	1/4	4/4	Tall	3/4	2/4	Yes	3/4	1/4
Green	2/4	1/4	3	3/4	0/4	Short	1/4	2/4	No	1/4	3/4

p(M|New Instance) = p(M) * p(Color = Green|M) * p(Legs = 2|M) * p(Height = tall|M) * p(Smelly = no |M)

 $p(M|New\ Instance) = 0.5 * \frac{2}{4} * \frac{1}{4} * \frac{3}{4} * \frac{1}{4} = 0.0117$

p(H|New Instance) = p(H) * p(Color = Green|H) * p(Legs = 2|H) * p(Height = tall|H) * p(Smelly = no |H)

$$p(H|New\ Instance) = 0.5 * \frac{1}{4} * \frac{4}{4} * \frac{2}{4} * \frac{3}{4} = 0.047$$

p(H|New Instance) > p(M|New Instance)

Hence the new instance belongs to Speces H

NAIVE BAYES CLASSIFIER - AN EXAMPLE

Example No.	Color	Туре	Origin	Stolen?
1	Red	Sports	Domestic	Yes
2	Red	Sports	Domestic	No
3	Red	Sports	Domestic	Yes
$4 \cap S$	Yellow	Sports	Domestic	C No
5	Yellow	Sports	Imported	Yes
6	Yellow	SUV	Imported	No
7	Yellow	SUV	Imported	Yes
8	Yellow	SUV	Domestic	No
9	Red	SUV	Imported	No
10	Red	Sports	Imported	Yes

Example No.	Color	Туре	Origin	Stolen?
1	Red	Sports	Domestic	Yes
2	Red	Sports	Domestic	No
3	Red	Sports	Domestic	Yes
4	Yellow	Sports	Domestic	No
5	Yellow	Sports	Imported	Yes
6	Yellow	SUV	Imported	No
7	Yellow	SUV	Imported	Yes
8	Yellow	SUV	Domestic	No
9	Red	SUV	Imported	No
10	Red	Sports	Imported	Yes

New Instance = (Red, SUV, Domestic), No

NAIVE BAYES CLASSIFIER EXAMPLE - 3

p(Yes) =	$\frac{5}{10} = 0.5$
p(No) =	$\frac{5}{10} = 0.5$

Color	Yes	No
Red	3/5	2/5
Yellow	2/5	3/5

Ν	Туре	Yes	No
	Sports	4/5	2/5
	SUV	1/5	3/5

]	Origin	Yes	No
	Domestic	2/5	3/5
	Imported	3/5	2/5

P(Yes|New Instance) = p(Yes) * P(Color = Red|Yes) * P(Type = SUV|Yes) * P(Origin = Domestic|Yes)

 $P(Yes|New \,Instance) = \frac{5}{10} * \frac{3}{5} * \frac{1}{5} * \frac{2}{5} = \frac{3}{125} = 0.024$

P(No|New Instance) = p(No) * P(Color = Red|No) * P(Type = SUV|No) * P(Origin = Domestic|No)

 $P(No|New Instance) = \frac{5}{10} * \frac{2}{5} * \frac{3}{5} * \frac{3}{5} = \frac{9}{125} = 0.072$ P(No|New Instance) > P(Yes|New Instance)

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NAIVE BAYES CLASSIFIER - AN EXAMPLE: LEARNING TO CLASSIFY TEXT

Consider a football game between two rival teams, say team A and team B. Suppose team A wins 65% of the time and team B wins the remaining matches. Among the games won by team A, only 35% of them comes from playing at team B's football field. On the other hand, 75% of the victories for team B are obtained while playing at home.

- 1. If team B is to host the next match between the two teams, what is the probability that it will emerge as the winner?
- emerge as the winner?
 If team B is to host the next match between the two teams, who will emerge as the winner?
 Solution:
- Probability that team A wins is $P(Y_A) = 0.65$.

- Y Winning football match
- X Hosting football match

Probability that team B wins is $P(Y_B) = 1 - P(Y_A) = 0.35$

Probability that team B hosted the match it had won is $P(X_B | Y_B) = 0.75$.

Probability that team B hosted the match won by team A is $P(X_B|Y_A) = 0.35$.

NAIVE BAYES CLASSIFIER - AN EXAMPLE: LEARNING TO CLASSIFY TEXT

The above question can be solved by computing $P(Y_B|X_B)$, which is the conditional probability

that team B wins the next match it hosts. Using the Bayes theorem, we obtain:

$$P(YB|XB) = \frac{P(X_B|Y_B) \times P(Y_B)}{P(X_B)}$$

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$$= \frac{P(X_B|Y_B) \times P(Y_B)}{P(X_B|Y_B) P(Y_B) + P(X_B|Y_A) P(Y_A)}$$

$$= \frac{0.75 \times 0.35}{(0.75 \times 0.35 + 0.35 \times 0.65)} = 0.5357$$

NAIVE BAYES CLASSIFIER EXAMPLE – 4

1. If team B is to host the next match between the two teams, what is the probability that it will emerge as the winner?

Solution:

$$P(Y_B|X_B) = \frac{P(X_B|Y_B) \times P(Y_B)}{P(X_B)} \quad \text{HIT,}$$
$$= \frac{P(X_B|Y_B) \times P(Y_B)}{P(X_B|Y_A) P(Y_A) + P(X_B|Y_B) P(Y_B)}$$
$$= \frac{0.75 \times 0.35}{(0.35 \times 0.65 + 0.75 \times 0.35)}$$

Probability that team A wins is $P(Y_A) = 0.65$.

Probability that team B wins is $P(Y_B) = 1 - P(Y_A) =$

0.35 Probability that team B hosted the match it had won is $P(X_B|Y_B) = 0.75$. Probability that team B hosted the match won by

team A is $P(X_B | Y_A) = 0.35$.

= 0.5357

NAIVE BAYES CLASSIFIER EXAMPLE – 4

2. If team B is to host the next match between the two teams, who will emerge as the winner?

Solution:

Ρ

$$(Y_A | X_B) = \frac{P(X_B | Y_A) \times P(Y_A)}{P(X_B)} \text{ HIT,}$$
$$= \frac{P(X_B | Y_A) \times P(Y_A)}{P(X_B | Y_A) P(Y_A) + P(X_B | Y_B) P(Y_B)}$$
$$= \frac{0.35 \times 0.65}{(0.35 \times 0.65 + 0.75 \times 0.35)}$$

Probability that team A wins is $P(Y_A) = 0.65$.

Probability that team B wins is $P(Y_B) = 1 - P(Y_A) =$

0.35 Probability that team B hosted the match it had won is $P(X_B|Y_B) = 0.75$. Probability that team B hosted the match won by

team A is $P(X_B | Y_A) = 0.35$.

= 0.4642

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- Bayesian networks (BN) are probabilistic graphical models that are based on directed acyclic graphs.
- They provide a tool to deal with two problems: uncertainty and complexity.
- Hence, they provide a compact representation of joint probability distributions using a combination of graph theory and probability theory.
- The graph structure specifies statistical dependencies among the variables and the local probabilistic models specify how these variables are combined.

- The naive Bayes classifier makes significant use of the assumption that the values of the attributes **a1...an** are conditionally independent given the target value v.
- This assumption dramatically reduces the complexity of learning the target

function

- CSE, HIT, Nidasoshi
 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
- Bayesian belief networks allow stating conditional independence assumptions

that apply to subsets of the variables

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- Consider an arbitrary set of random variables Y1... Yn, where each variable Yi can take on the set of possible values V(Yi).
- The joint space of the set of variables Y to be the cross product V(Y1) x V(Y2) x... V(Yn).
- In other words, each item in the joint space corresponds to one of the possible assignments of values to the tuple of variables (Y1 ... Yn). The probability distribution over this joint' space is called the joint probability distribution.
- The joint probability distribution specifies the probability for each of the possible variable bindings for the tuple (Y1 . . . Yn).
- A Bayesian belief network describes the joint probability distribution for a set of variables.

- 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; that is, 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 $x_i \in V(X), y_j \in V(Y)$, and $z_k \in V(Z)$. Solution

P(X|Y,Z) = P(X|Z)

• We say that the set of variables X1 . . . Xi is conditionally independent of the set of variables Y1 . . . Ym given the set of variables Z1 . . . Zn, if

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

BAYESIAN BELIEF NETWORKS - REPRESENTATION

• The joint probability for any desired assignment of values **(y1, . . . , yn)** to the tuple of network variables **(Y1 . . . Yn)** can be computed by the formula

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

CSE, HIT, Nidasoshi

- where Parents(Yi) denotes the set of immediate predecessors of Yi in the network.
- Note the values of P(yi | Parents(Yi)) are precisely the values stored in the conditional probability table associated with node Yi.



Dr. Mahesh (

- You have a new burglar alarm installed at home.
- It is fairly reliable at detecting burglary, but also sometimes responds to minor earthquakes.
- You have two neighbors, John and Merry, who promised to call you at work when they hear the alarm. SEHIT, Nidasoshi
 John always calls when he hears the alarm, but sometimes confuses telephone
- John always calls when he hears the alarm, but sometimes confuses telephone ringing with the alarm and calls too.
- Merry likes loud music and sometimes misses the alarm.
- Given the evidence of who has or has not called, we would like to estimate the probability of a burglary.

BAYESIAN BELIEF NETWORKS - EXAMPLE





Solution:

 $P(j \land m \land a \land \neg b \land \neg e) = P(j \mid a) P(m \mid a) P(a \mid \neg b, \neg e) P(\neg b) P(\neg e)$

 $= 0.90 \times 0.70 \times 0.001 \times 0.999 \times 0.998$



+ $P(j|-a){P(-a|b,e)*P(b,e)+P(-a|-b,e)*P(-b,e)+P(-a|b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b,-e)*P(-a|-b$

P(¬b, ¬e)}

= 0.90 * 0.00252 + 0.05 * 0.9974 = 0.0521

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- Suppose, we are given for the evidence variables E₁,...,E_m, their values e₁,...,e_m, and we want to predict whether the query variable X has the value x or not.
- For this we compute and compare the following:

3. What is the probability that there is a burglary given that John and Merry calls?

 $P(b \mid j, m)$

 $= \alpha P(b, j, m)$

$$= \alpha \sum_{a} \sum_{e} P(b, j, m, a, e)$$

= $\alpha \left(P(b, j, m, a, e) + P(b, j, m, \neg a, e) + P(b, j, m, a, \neg e) + P(b, j, m, \neg a, \neg e) \right)$

$$\begin{split} &P(\neg b \mid j,m) \\ &= \alpha P(\neg b, j,m) \\ &= \alpha \sum_{a} \sum_{e} P(\neg b, j,m,a,e) \\ &= \alpha \Big(P(\neg b, j,m,a,e) + P(\neg b, j,m,\neg a,e) + P(\neg b, j,m,a,\neg e) + P(\neg b, j,m,\neg a,\neg e) \Big) \end{split}$$

3. What is the probability that there is a burglary given that John and Merry calls?

 $P(b \mid j,m) = \alpha P(b) \sum_{a} P(j|a) P(m|a) \sum_{e} P(a|b,e) P(e)$

 $= \alpha P(b) \sum_{a} P(j|a) P(m|a) \left\{ P(a|b,e)P(e) + P(a|b,\neg e)P(\neg e) \right\}$ = $\alpha P(b) \left[P(j|a)P(m|a) \left\{ P(a|b,e)P(e) + P(a|b,\neg e)P(\neg e) \right\} \right]$

 $+ P(j|\neg a)P(m|\neg a) \left\{ P(\neg a|b,e)P(e) + P(\neg a|b,\neg e)P(\neg e) \right\} \right]$

 $=\alpha*.001*(.9*.7*(.95*.002+.94*.998)+.05*.01*(.05*.002+.71*.998))$

 $= \alpha * .00059$
BAYESIAN BELIEF NETWORKS - EXAMPLE

3. What is the probability that there is a burglary given that John and Merry calls?

$$\begin{split} P(\neg b \mid j,m) &= \alpha P(\neg b) \sum_{a} P(j|a) P(m|a) \sum_{e} P(a|\neg b,e) P(e) \\ &= \alpha P(\neg b) \sum_{a} P(j|a) P(m|a) \left\{ \begin{array}{c} P(a|\neg b,e) P(e) + P(a|\neg b,\neg e) P(\neg e) \right\} \\ &= \alpha P(\neg b) \left[P(j|a) P(m|a) \left\{ \begin{array}{c} P(a|\neg b,e) P(e) + P(a|\neg b,\neg e) P(\neg e) \right\} \\ &+ P(j|\neg a) P(m|\neg a) \left\{ \begin{array}{c} P(\neg a|\neg b,e) P(e) + P(\neg a|\neg b,\neg e) P(\neg e) \right\} \right] \\ &= \alpha * .999^* (.9^*.7^* (.29^*.002 + .001^*.998) + .05^*.01^* (.71^*.002 + .999^*.998)) \end{split}$$

 $= \alpha * .0015$

BAYESIAN BELIEF NETWORKS - EXAMPLE

3. What is the probability that there is a burglary given that John and Merry calls?

 $\alpha = 1/(.00059 + .0015)$

$$CSE, HIT, NidasoshiP(b | j,m) = 478.5 * .00059$$

= 0.28
 $P(\neg b | j,m) = 478.5 * .0015$
= 0.72

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- kMeans algorithm is an unsupervised learning algorithm
- Given a data set of items, with certain features, and values for these

features, the algorithm will categorize the items into k groups or clusters of

similarity. CSE, HIT, Nidasoshi

• To calculate the similarity, we can use the Euclidean distance,

Manhattan distance, Hamming distance, Cosine distance as measurement.



Here is the pseudocode for implementing a K-means algorithm.

Input: Algorithm K-Means (K number of clusters, D list of data points)

1. Choose K number of random data points as initial centroids (cluster centers). CSE, HIT, Nidasoshi

2. Repeat till cluster centers stabilize:

a. Allocate each point in D to the nearest of Kth centroids.

b. Compute centroid for the cluster using all points in the cluster.



Advantages and Disadvantages of K-Means Algorithm

Advantages of K-Means Algorithm

- 1. K-means algorithm is simple, easy to understand, and easy to implement.
- 2. It is also efficient, in which the time taken to cluster K-means rises linearly with the number of data points.
- 3. No other clustering algorithm performs better than K-means.

Disadvantages of K-Means Algorithm

- 1. The user needs to specify an initial value of K.
- 2. The process of finding the clusters may not converge.
- 3. It is not suitable for discovering clusters that are not hyper ellipsoids or hyper spheres).







			Distance to			Cluster
	Χ	Υ	(1, 5)	(4, 1)	(8, 4)	Number
<u>Iteration - 1</u> C1 - Seed Point1 – (1, 5) C2 - Seed Point2 – (4, 1) C3 - Seed Point3 – (8, 4)	2	4	1.41	3.61	6.00	C1
	2	6	1.41	5.39	6.32	C1
	5	6	4.12	5.10	3.61	C3
	4	- 7	3.61	26.00 S	5.00	C1
$D = \sqrt{((x_2 - x_1)^2 + (y_2 - y_1)^2)}$	8	3	7.28	4.47	1.00	C3
	6	6	5.10	5.39	2.83	C3
	5	2	5.00	1.41	3.61	C2
C1 – Centroid – (2.66, 5.66) C2 – Centroid – (4.5, 3) C3 – Centroid – (6, 5)	5	7	4.47	6.08	4.24	C3
	6	3	5.39	2.83	2.24	C3
	4	4	3.16	3.00	4.00	C2

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			Distance to			Cluster
	Χ	Y	(2.66, 5.66)	(4.5, 3)	(6, 5)	Number
$\frac{\text{Iteration - 2}}{\text{C1 - Centroid} - (2.66, 5.66)}$ $C2 - Centroid - (4.5, 3)$ $C3 - Centroid - (6, 5)$ $C1 - Centroid - (2.66, 5.66)$ $C2 - Centroid - (5, 3)$ $C3 - Centroid - (6, 5.5)$	2	4	1.79	2.69	4.12	C1
	2	6	0.74	3.91	4.12	C1
	5	6	2.36	3.04	1.41	C3
	_4	- 7 -	1.90	c 4.03h	2.83	C1
	8	3	5.97	3.5	2.83	C3
	6	6	3.36	3.35	1	C3
	5	2	4.34	1.12	3.16	C2
	5	7	2.70	4.03	2.24	C3
	6	3	4.27	1.5	2	C2
	4	4	2.13	1.12	2.24	C2

			Distance to			Cluster
	Χ	Υ	(2.66, 5.66)	(5, 3)	(6, 5.5)	Number
Iteration - 3	2	4	1.79	3.16	4.27	C1
C1 - Centroid - (2.66, 5.66) $C2 - Centroid - (5, 3)$ $C3 - Centroid - (6, 5.5)$ $C1 - Centroid - (2.66, 5.66)$ $C2 - Centroid - (5.75, 3)$ $C3 - Centroid - (5.33, 6.33)$	2	6	0.74	4.24	4.03	C1
	5	6	2.36	3.00	1.12	C3
	4	_7_	1.90	c 412h	2.50	C1
	8	3	5.97	3.00	3.20	C2
	6	6	3.36	3.16	0.50	C3
	5	2	4.34	1.00	3.64	C2
	5	7	2.70	4.00	1.80	C3
	6	3	4.27	1.00	2.50	C2
	4	4	2.13	1.41	2.50	C2

			C	Cluster		
	Χ	Υ	(2.66, 5.66)	(5.75, 3)	(5.33, 6.33)	Number
<u>Iteration - 4</u>	2	4	1.79	3.88	4.06	C1
C1 – Centroid – (2.66, 5.66) C2 – Centroid – (5.75, 3) C3 – Centroid – (5.33, 6.33)	2	6	0.74	4.80	3.35	C1
	5	6	2.36	3.09	0.47	C3
	-4	-7	1.90	a4375	1.49	C3
C1 – Centroid – (2, 5) C2 – Centroid – (5.75, 3) C3 – Centroid – (5, 6.5)	8	3	5.97	2.25	4.27	C2
	6	6	3.36	3.01	0.75	C3
	5	2	4.34	1.25	4.34	C2
	5	7	2.70	4.07	0.75	C3
	6	3	4.27	0.25	3.40	C2
	4	4	2.13	2.02	2.68	C2

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			0	Cluster		
	Χ	Υ	(2, 5)	(5.75, 3)	(5, 6.5)	Number
<u>Iteration - 5</u> C1 – Centroid – (2, 5) C2 – Centroid – (5.75, 3) C3 – Centroid – (5, 6.5)	2	4	1.00	3.88	3.91	C1
	2	6	1.00	4.80	3.04	C1
	5	6	3.16	3.09	0.50	C3
	4	-7	2.83	a 437S	1.12	C3
No movement of data Points Hence these are the final positions	8	3	6.32	2.25	4.61	C2
	6	6	4.12	3.01	1.12	C3
	5	2	4.24	1.25	4.50	C2
	5	7	3.61	4.07	0.50	C3
	6	3	4.47	0.25	3.64	C2
	4	4	2.24	2.02	2.69	C2

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MAXIMUM LIKELIHOOD AND LEAST-SQUARED ERROR HYPOTHESES



- In the real-world applications of machine learning, it is very common that there are many relevant features available for learning but only a small subset of them are observable.
- So, for the variables which are sometimes observable and sometimes not, then we can use the instances when that variable is visible is observed for the purpose of learning and then predict its value in the instances when it is not observable.
- The *Expectation-Maximization algorithm* can be used for the latent variables (variables that are not directly observable and are actually inferred from the values of the other observed variables) in order to predict their values with the condition that the general form of probability distribution governing those latent variables is known to us.
- This algorithm is actually at the base of many unsupervised clustering algorithms in the field of machine learning.

Let us understand the EM algorithm in detail.

- Initially, a set of initial values of the parameters are considered.
- A set of incomplete observed data is given to the system with the assumption that the observed data comes from a specific model.
- The next step is known as "Expectation" step or *E-step*. In this step, we use the observed data in order to estimate or guess the values of the missing or incomplete data. It is basically used to update the variables.
- The next step is known as "Maximization"-step or *M-step*. In this step, we use the complete data generated in the preceding "Expectation" step in order to update the values of the parameters. It is basically used to update the hypothesis.
- Now, in the fourth step, it is checked whether the values are converging or not, if yes, then stop otherwise repeat *step-2* and *step-3* i.e. "Expectation" – step and "Maximization" – step until the convergence occurs.

Algorithm:

- 1. Given a set of incomplete data, consider a set of starting parameters.
- 2. Expectation step (E step): Using the observed available data of the dataset,

estimate (guess) the values of the missing data.

3. Maximization step (M – step): Complete data generated after the expectation

(E) step is used in order to update the parameters.

4. Repeat step 2 and step 3 until convergence.



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Advantages of EM algorithm –

- It is always guaranteed that likelihood will increase with each iteration.
- The E-step and M-step are often pretty easy for many problems in terms of implementation.
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- Solutions to the M-steps often exist in the closed form.

Disadvantages of EM algorithm –

- It has slow convergence.
- It makes convergence to the local optima only.
- It requires both the probabilities, forward and backward (numerical optimization requires only forward probability).

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- The easiest way to introduce the EM algorithm is via an example.
- 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 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.



- 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.
- For example the parameters of interest were $\theta = (\mu 1, \mu 2)$, and the full data were the triples (*xi, zi1, zi2*) of which only the *xi* were observed.
- In general let X = {x1, ..., xm} denote the observed data in a set of m independently drawn instances, let Z = {z1, ..., zm} denote the unobserved data in these same instances, and let Y = X U Z denote the full data.

• Note 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[In P(Y |h')].
- This expected value is taken over the probability distribution governing Y, which is determined by the unknown parameters θ.
- Let us consider exactly what this expression signifies.
- 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.
- Second, maximizing the logarithm of this quantity In P(Y|h') also maximizes P(Ylh'), as we have discussed on several occasions already.
- Third, we introduce the expected value E[In P(YIh')] because the full data Y is itself a random variable.

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- 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[In P(YIh')] over the probability distribution governing the random variable Y. HIT, NICASOSNI
- The distribution governing **Y** is determined by the completely known values for X, plus the distribution governing Z.

- What is the probability distribution governing Y?
- In general we will not know this distribution because it is determined by the parameters θ that we are trying to estimate.
- Therefore, the EM algorithm uses its current hypothesis h in place of the actual parameters
 θ to estimate the distribution governing Y.
- Let us define a function Q(h'|h) that gives E[In P(Y Ih')] as a function of h', under the assumption that θ = 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.

$Q(h'|h) \leftarrow E[\ln P(Y|h')|h, X]$ CSE, HIT, Nidasoshi

• Step 2: *Maximization (M) step:* Replace hypothesis *h* by the hypothesis *h'* that maximizes this Q function.

$$h \leftarrow \operatorname*{argmax}_{h'} Q(h'|h)$$
- The k-means problem is to estimate the parameters θ = (μ1, μ2) that define the means of the k
 Normal distributions. We are given the observed data X = { < xi > }.
- The hidden variables Z = { < z i1, ..., zik>} in this case indicate which of the k Normal distributions was used to generate xi.
- 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 Inp(Y | h'),
- Note the probability p(yi |h') of a single instance yi = (xi,zi1, ... zik) of the full data can be written,

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

Given this probability for a single instance *p(yi/h')*, the logarithm of the probability In *P(Y/h')* for all *m* instances in the data is,

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

$$CS \sum_{i=1}^{m} \ln p(y_i|h_i) \text{ Nidasoshi}$$

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

 Finally we must take the expected value of this In P(Y|h') over the probability distribution governing Y

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

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

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

- Thus, the first (estimation) step of the EM algorithm defines the Q function based on the estimated E[zij] terms.
- The second (maximization) step then finds the values μ₁', μ_n' that maximize this Q function.
- In the current case

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

• Thus, the maximum likelihood hypothesis here minimizes a weighted sum of squared errors, where the contribution of each instance xi to the error that defines μ_j is weighted by E[zij].