

2 Marks

1. Ans Frequent pattern mining: A Road map:

- Frequent pattern mining can be classified in various ways, based on the following criteria
- i, Based on the completeness of patterns to be mined.
 - ii, Based on the levels of abstraction involved in the rule set.
 - iii, Based on the no. of data dimensions involved in the rules.
 - iv, Based on the types of values handled in the rules.
 - v, Based on the kinds of rules to be mined.
 - vi, Based on the kinds of patterns to be mined.

2. Ans Apriori Steps:-

1. It is an influential algorithm for mining frequent itemset for Boolean association rules, this uses prior knowledge of itemset properties.
2. It follows iterative approach known as "level-wise" search where 'k' itemsets are used to explore (k+1) itemset.
3. First, the set of frequent itemset are found denoted by 'L₁', which is used to find 'L₂' which in turn use to find 'L₃' and so on.
4. Apriori property says that "All non-empty subset of a frequent itemset must also be frequent", it is based on the observations that if an itemset does not satisfy.

step 5: if an item 'A' is added to the itemset 'I' then resulting itemset $(I \cup A)$ cannot occur more frequently than 'I' i.e., $(I \cup A)$ is not found frequent.

Therefore probability of $(I \cup A) < \text{min_sup}$

step 6: A two step process is followed in generating candidate and frequent itemsets such as join and prune actions.

Algorithm

C_k : candidate itemset of size k and L_k : Frequent itemset of size k

$L_1 = \{\text{frequent item}\}$;

for $(k=1; L_k \neq \phi; k++)$ do begin

C_{k+1} = candidate generated from L_k ;

for each transaction t in database do

increment the count of all candidates in C_{k+1} that are contained in t

L_{k+1} = candidates in C_{k+1} with min-support.

end

return $L = \bigcup_k L_k$;

3AM FP-growth :-

It is an Algorithm is an efficient and scalable method for mining the complete set of frequent pattern by pattern fragment growth, using an extend prefix Tree structure for storing compressed and crucial information about frequent pattern named frequent pattern tree.

4Au a) correlation is a term that is a measure of the strength of a linear relationship b/w two quantitative variables.

Ex: positive correlation may be that the more exercise. The more calories you will burn.

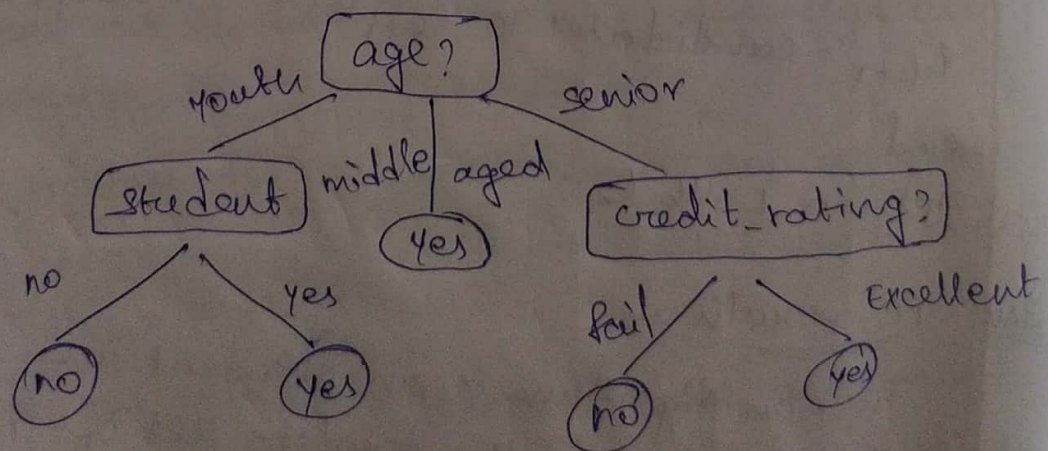
b) outline chi-square test with example

The chi-square test of association evaluates relationships b/w categorical variables.

5Au Decision Tree Induction is the learning of decision from class-labeled training tuples. A decision tree

- * flow-chart like Tree structure.
- * Each internal node denotes a test on an attribute
- * each branch represents an outcome of the test
- * Each leaf node holds a class label.
- * The topmost node in a tree is the root node

Ex:

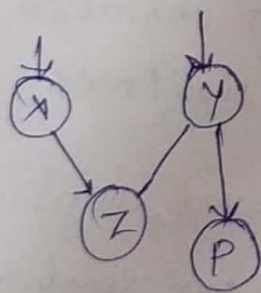


1Aii Bayesian Network:-

This can be deal with dependencies. This specifies joint conditional probability distribution. Bayesian belief network are also known as belief network, Bayesian network, and probabilistic networks. A belief network is defined by two components

- A directed acyclic graph and
- A set of conditional probability tables.

A graphical model of causal relationship. Represents dependency among the variables and gives a specification of joint probability distribution

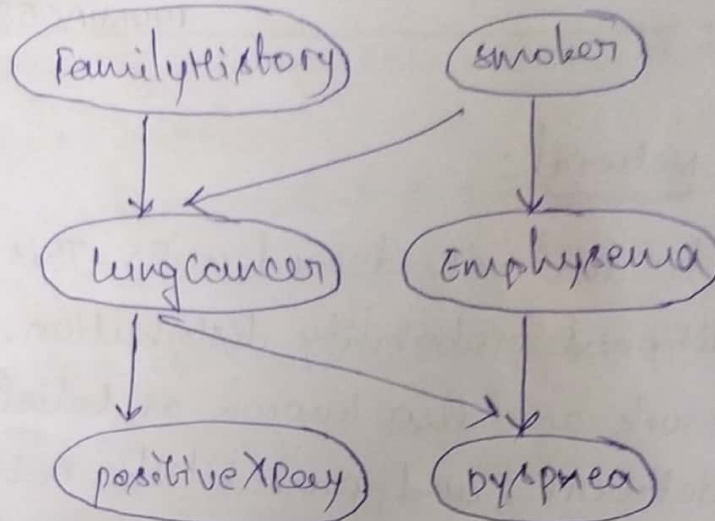


- Nodes : random variables
- Links : dependency
- X & Y are the parents of Z, and Y is the parent of P.
- No dependency b/w Z & P.
- Has no loops or cycles.

Ex:- A Simple Bayesian Network:

- A proposed causal model represented by a directed acyclic graph.
- The conditional probability table for the values of the variable, LungCancer(LC) showing each possible combination of the values of its parents node, FamilyHistory (FH) and Smoker (S).

a)



b)

	FH, S	FH, \sim S	\sim FH, S	\sim FH, \sim S
LC	0.8	0.5	0.7	0.1
-LC	0.2	0.5	0.3	0.9

The conditional probability for each known value of LungCancer is given for each possible combination of values of its parents. For instance, from the upper leftmost and bottom rightmost entries

$$P(\text{LungCancer} = \text{yes} \mid \text{FamilyHistory} = \text{yes}, \text{Smoker} = \text{yes}) = 0.8$$

$$P(\text{LungCancer} = \text{no} \mid \text{FamilyHistory} = \text{no}, \text{Smoker} = \text{no}) = 0.9$$

let $X = (x_1, \dots, x_n)$ be a data tuple described by the variables or attributes y_1, \dots, y_n respectively.

$P(x_1, \dots, x_n)$ is the probability of a particular combination of values of X , and the values for $P(x_i \mid \text{parents}(y_i))$ correspond to the entries in the CPT for y_i .

2 a) classification by Backpropagation

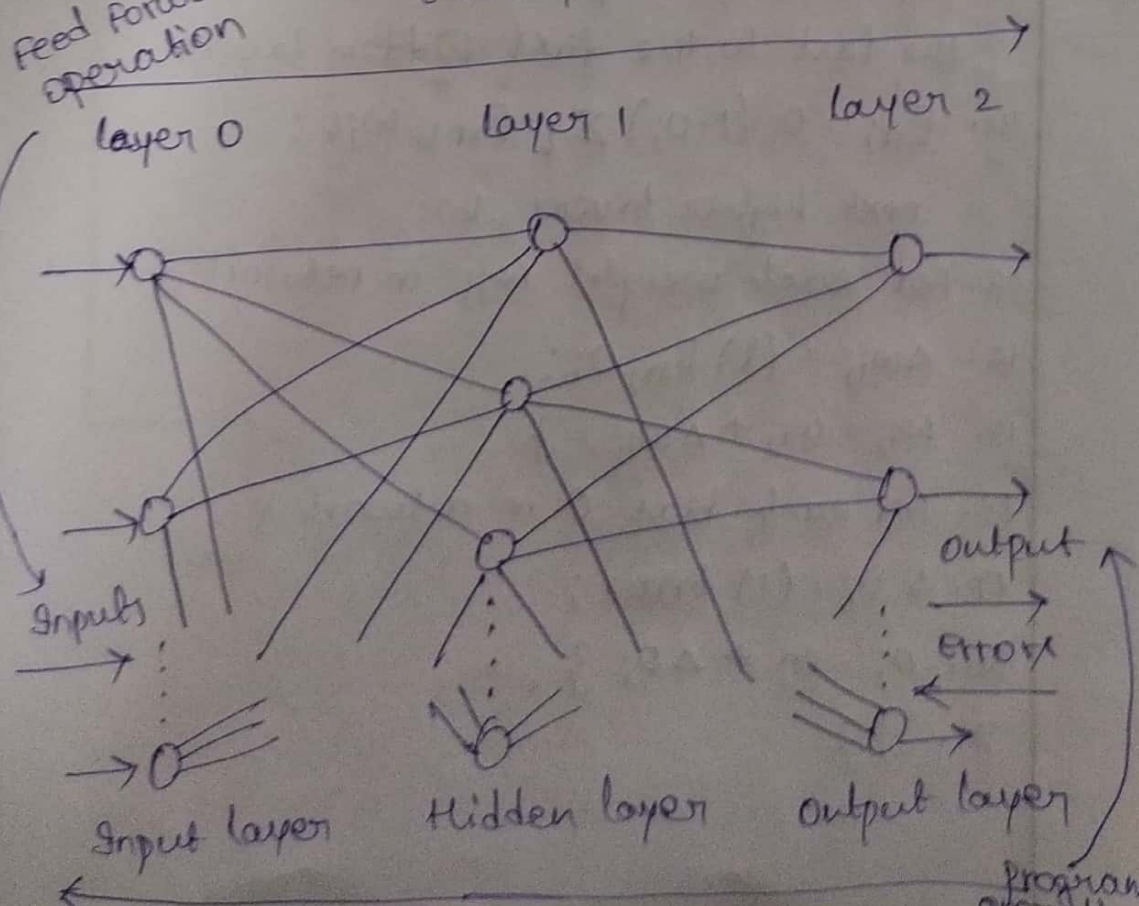
Ans: • Iteratively process a set of training tuples and compare the network prediction with the actual known target value.

- For each training tuple, the weights are modified to minimize the mean squared error plus the networks prediction
- Modifications are made in the backward direction

Steps:

- * Initialize weight and biases in the network
- + propagate the inputs forward.
- + Back propagate the error.
- * Terminating condition

Input vectors → actual value is calculated then Error is calculated.



Method :-

1. Initialize all weights and biases in network;
2. while terminating condition is not satisfied {
3. for each training tuple x in D {
4. // propagate the inputs forward;
5. for each input layer until j {
6. $O_j = I_j$;
7. for each hidden or output layer unit j {
8. $I_j = \sum_i w_{ij} O_i + \theta_j$;
9. $O_j = \frac{1}{1 + e^{-I_j}}$ };
10. // Backpropagate the errors;
11. for each unit j in the output layer
12. $Err_j = O_j (1 - O_j) (T_j - O_j)$;
13. for each unit j in the hidden layers, from the last to the first hidden layers.
14. $Err_j = O_j (1 - O_j) \sum_k Err_k w_{jk}$;
next higher layers, k
15. for each weight w_{ij} in network {
16. $\Delta w_{ij} = (l) Err_j O_i$;
17. $w_{ij} = w_{ij} + \Delta w_{ij}$ };
18. for each bias θ in network {
19. $\Delta \theta_j = (1) Error$;
20. $\theta_j = \theta_j + \Delta \theta_j$ };
21. } }

2b) Support Vector Machine:-

It is an algorithm for classification of both linear and nonlinear data. SVM is an algorithm that works as follows

- It uses a non linear mapping to transform the original training data into a higher dimension.
- within this new dimensions, it searches for the linear optimal separating hyperplane.
- with an appropriate nonlinear mapping to a sufficiently high dimension, data from two classes can always be separated by a hyperplane.
- The SVM finds this hyperplane using support vectors (essential training tuples) and margins.

SVM general philosophy will be as follows:

