

pg 28 soln

ML

Q-1

Q1) What is an artificial neuron?

An artificial neuron is a mathematical function that models biological neuron in an artificial neural network (ANN). It is a tiny part of a computer system that takes i/p, does some calculations, & gives an o/p, similar to how a brain cell works.

(Diagram) - Done

Q2) Describe the multipoint crossover for a GA problem. Done

Q3) Differentiate the fuzzy sets for the triangular & trapezoidal membership functions.

Definition Done. Fuzzy sets use MFs to define the degree of belonging for elements within the set.

Key differences of fuzzy sets on the basis of triangular & trapezoidal MF:

Feature	Tri <sup>o</sup>	Trape <sup>o</sup>
No. of Parameters	3 (a, b, c)	4 (a, b, c, d)
Shape	Single peak (triangle)	Flat-top (trapezoid)
Complexity	Simpler, fewer parameters	Slightly more complex
Application	Sharp, single-point focus	Broader range of fuzzy membership.



↳ compare the justification & defuzzification of a fuzzy inference system.

### Fuzzification

- i) It converts crisp i/p vals. into fuzzy sets (crisp & hard, fuzzy = soft)
- ii) In FIS, it's input interface for fuzzifying the system's input.
- iii) It's direction is Crisp to fuzzy.
- iv) Crisp i/p data is given as i/p.
- ↳ technique used - MFs (fm, trapez, gauss)
- vi) It's key func is enabling reasoning with linguistic variables.

### Defuzzification

- i) It converts fuzzy o/p sets into a crisp o/p val.
- ii) output interface for producing actionable result in FIS.
- iii) It's direction is fuzzy to crisp.
- iv) Fuzzy o/p sets from FIS is given as o/p
- ↳ technique used - Defuzzification methods (Centroid wav, mean of maxima)
- vii) It's key func is facilitating practical decision-making / ctrl.

sketch the basic steps for a GA problem.

### Ques

↳ what is reinforcement training?  
It refers to a method of learning where desired behaviours are encouraged by providing positive feedback / rewards when a specific action is performed correctly. (or) Pmr

### Ans

- 1) Establish a standard fuzzy inference system with proper example. Ques
- 2) Establish a explain the terms, "chromosome", gene, Allele, Locus, Genotype, phenotype" with proper examples for a GA problem. Ques

↳ eg let a string / chromosome is = 101011

- i) chromosome = 101011
- ii) gene = each digit of 101011, eg = 1 or 0
- iii) Allele = (val. of gene) for binary chromo. Possible alleles are 0 or 1
- iv) Locus = ~~the~~ the locus of the 3rd gene (val = 1) is position = 3

101011  
1 2 3 4 5 6

↳ Genotype = the full str 101011

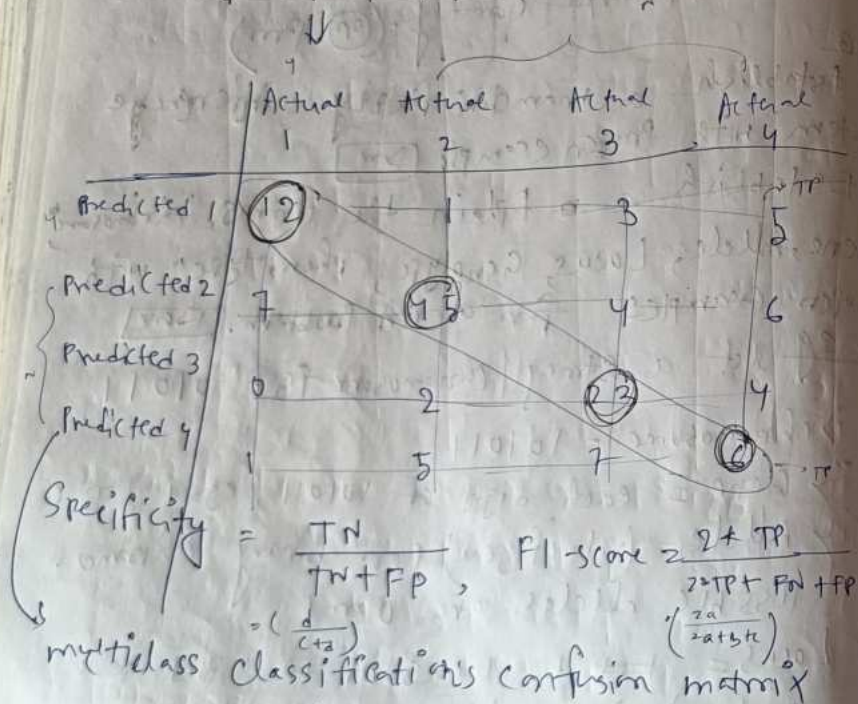
↳ Phenotype =  $\begin{matrix} 1 & 0 & 1 & 0 & 1 & 1 \\ \downarrow & \downarrow & \downarrow & \downarrow & \downarrow & \downarrow \\ A & B & C & & & \end{matrix}$  encoding

↳ Establish a MDC for a 3-class classification problem. Ques (from Data Sci, both algo & mng)



1) calculate specificity & f1-score for the given confusion matrix.

Predicted \ Actual	Actual			
	1	2	3	4
1	12	7	0	1
2	45	4	2	5
3	23	4	6	1
4	5	7	13	6



Converting it into C.M. for binary classification (binary C.M.)

Step 1

	TP	TN	FP	FN
1	12	95	7	0
2	45	1	2	4
3	23	4	6	1
4	5	7	13	6

Step 1

	TP	TN	FP	FN
1	12	95 + 4 + 2 + 1 = 102	7 + 0 + 1 = 8	1 + 3 + 5 = 9
2	45	12 + 3 + 5 + 1 + 4 + 6 = 31	1 + 2 + 5 = 8	1 + 4 + 6 = 11
3	23	12 + 1 + 5 + 7 + 4 + 6 = 35	3 + 4 + 7 = 14	0 + 2 + 4 = 6
4	6	12 + 1 + 3 + 5 + 7 + 4 + 10 + 2 + 23 = 57	5 + 6 + 4 = 15	1 + 5 + 7 = 13

FP = Col. sum - TP  
FN = Row. sum - TP

Specificity of 1 =  $\frac{102}{102 + 8} = 92\%$

Specificity of 2 =  $\frac{31}{31 + 8} = 79\%$

Specificity of 3 =  $\frac{35}{35 + 14} = 71\%$

Specificity of 4 =  $\frac{57}{57 + 15} = 79\%$

F1 Score of 1 =  $\frac{2 * 12}{2 * 12 + 9 + 0} = 57\%$

F1 Score of 2 =  $\frac{2 * 45}{2 * 45 + 11 + 4} = 83\%$

F1 Score of 3 =  $\frac{2 * 23}{2 * 23 + 6 + 1} = 70\%$

F1 Score of 4 =  $\frac{2 * 6}{2 * 6 + 13 + 6} = 30\%$



2) maximize the function,  $f(x) = 4x^2 + 9x + 1$ , where  $x = 9, 11, 13, 15$  with chromosome size = 4 such that if selection operation (Rank selection), if uniform crossover, (ii) UP to 2 iterations.

selection (Rank selection) → initial

String no.	Population	n	f(x) (fitness val)	Prob-Count	Expected-Count	Actual-Count
1	010010	9	406 (3rd)	0.21	0.84	1
2	010110	11	584 (2nd)	0.3	1.21	1
3	011010	13	794 (1st)	0.41	1.64	2
4	001010	5	146 (4th)	0.08	0.3	0

$$f(x) = 4x^2 + 9x + 1 \Rightarrow 4 \times 81 + 9 \times 9 + 1 = 406$$

$$\Rightarrow 4 \times 121 + 9 \times 11 + 1 = 584$$

$$\Rightarrow 4 \times 169 + 9 \times 13 + 1 = 794$$

$$\Rightarrow 4 \times 25 + 9 \times 5 + 1 = 146$$

$$\Sigma f(x) = 1930 \text{ (total fitness)}$$

$$\text{avg fitness} = \frac{\Sigma f(x)}{h} = 482.5$$

$$P.C = \frac{f(x)}{\Sigma f(x)}$$

$$E.C = \frac{f(x)}{\text{avg fitness}}$$

$$\text{Avg fitness}$$

$$A.C = \text{round}(E.C)$$

<del><math>S_1 = 10010</math></del>	$S_1 = 011010$
<del><math>S_2 = 10110</math></del>	$S_2 = 011010$
<del><math>S_3 = 11010</math></del>	$S_3 = 010110$
<del><math>S_4 = 11010</math></del>	$S_4 = 010010$

ii) Crossover (uniform)

$S_1 = 011010$	$S_3 = 011010$
$S_2 = 011010$	$S_4 = 011010$
$S_1' = 011010$	$S_3' = 011010$
$S_2' = 011010$	$S_4' = 011010$

iii) Color the table again for best val of n.

it > 1 iteration

off str.	popu	n	f(x)	P.C	E.C	A.C
$S_1'$	011010	13	794 (1st)	0.31	1.23	1
$S_2'$	011010	13	794 (1st)	0.31	1.23	1
$S_3'$	010010	9	406 (3rd)	0.16	0.63	1
$S_4'$	010110	11	584 (2nd)	0.27	0.91	1

$$\Sigma f(x) = 2578$$

$$\text{avg } f(x) = 644.5$$

ii) crossover

$S_1' = 011010$	$S_1' = 011010$	$S_4' = 010010$
$S_2' = 011010$	$S_2' = 011010$	$S_3' = 010110$
$S_3' = 010010$	$S_1' = 011010$	$S_3' = 010110$
$S_4' = 010110$	$S_2' = 011010$	$S_4' = 010010$

compute table for best val of n

off str.	popu	n	f(x)	P.C	E.C	A.C
$S_{11}'$	011010	13	794	0.31	1.23	1
$S_{21}'$	011010	13	794	0.31	1.23	1
$S_{31}'$	010110	11	584	0.27	0.91	1
$S_{41}'$	010010	9	406	0.16	0.63	1

$$\Sigma f(x) = 2578$$

$$\text{avg } f(x) = 644.5$$

(same as prev)



$\therefore$  the best val. of  $x$  after 2 iteration = 13  
(chromosome = 01101) with fitness val = 7.24

Ans-c

12) considers the fuzzy set  $Small = \{0/0 + 0.2/1 + 0.3 + 0/4\}$  & negative =  $\{0/1 + 0.7/2 + 1/3 + 0.7/4 + 0/5\}$  & the following fuzzy rule: "Rule 1: If  $x$  is small &  $y$  is negative then  $z$  is low". Find the firing strength of Rule 1 when  $x=3$  &  $y=2$  where fuzzy "AND" operation is the minimum operator. What is ELITISM?

$Small = \{0/0 + 0/2 + 1/3 + 0/4\}$

~~Ans-c~~ negative =  $\{0/1 + 0.7/2 + 1/3 + 0.7/4 + 0/5\}$

Rule 1: "If  $x$  is small &  $y$  is negative then  $z$  is low."

Given,  $x=3$ ,  $y=2$  & fuzzy "AND" operation is the minimum operator.

$x=3$ , compare.

i) Finding the membership value of  $x$  &  $y$  in the respective fuzzy sets

$Small(3) = 1$ ,  $negative(2) = 0.7$

ii) Applying AND operator =

$Small(3)$  AND  $negative(2)$

=  $\min\{Small(3), negative(2)\}$

=  $\min\{1, 0.7\}$

= 0.7

$\therefore$  firing strength is = 0.7

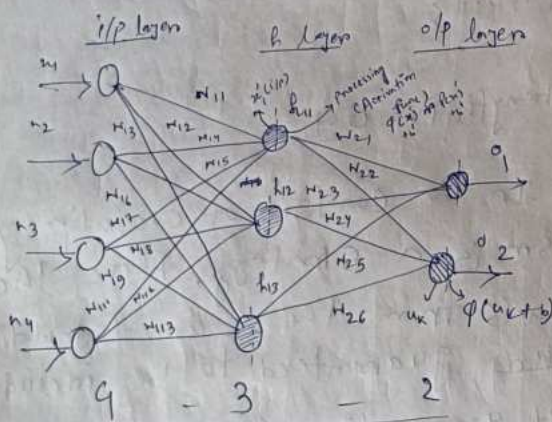
ELITISM - In the context of evolutionary algo., elitism is a strategy, where the best individuals from the current population are guaranteed to be included in the next generation. This ensures that the best solutions found so far, are not lost & can contribute to the search for even better solutions in future generations.

13) What is Gradient-Descent? Draw a very clear u-3-2 ANN architecture with explaining all its components. What is a self-organizing feature map?

Gradient-Descent - It is an iterative optimization algo used to find the minimum of a function. It works by repeatedly adjusting the parameters of a model in the direction of the negative gradient, that indicates the direction of the steepest descent.



4-3-2 ANN



$x_1, x_2, x_3, x_4 =$  i/p feature vectors

$o_1, o_2 =$  output class

$h_1, h_2, h_3 =$  hidden nodes

$W =$  weights

$\text{shaded circle} = u_k$  Processing node

$b =$  bias

$\phi(\cdot) =$  Activation func  $[\phi(x)]$   
 $[x = \text{input}]$

$$x = \text{input } u_k + b$$

$$= \sum_{j=1}^n W_{kj} \cdot x_j + b \quad \text{or} \quad \sum_{i=1, j=1}^{h, n} W_{ij} \cdot x_{ij} + b$$

[eg]  $h_{11} \rightarrow x_1' = u_k + b$   
 $= W_{11} \cdot x_1 + W_{12} \cdot x_2 + W_{13} \cdot x_3 + W_{14} \cdot x_4 + b$  (let  $b=0$ )

$h_{11} = \phi(x_1') = \tanh(x_1')$  [say A.F =  $\tanh(x)$ ]  
 $= \tanh(W_{11} \cdot x_1 + W_{12} \cdot x_2 + \dots + W_{14} \cdot x_4)$   
 $= \text{outcome}$

[eg]  $o_1 \rightarrow x_1'' = u_k + b$  (let  $b=0$ )

$= W_{21} \cdot h_{11} + W_{22} \cdot h_{12} + W_{23} \cdot h_{13} + b$   
 $o_1 = \phi(x_1'') = \tanh(x_1'')$   
 $= \tanh(W_{21} \cdot h_{11} + \dots + W_{23} \cdot h_{13})$   
 $= \text{outcome}$

I/p layer - The 1st layer of ANN, consisting of  $n$  number of nodes. Each node is a feature. I/p layer receives i/p data.

Hidden layer - The 2nd layer, consists of  $n$  nodes / neurons. It processes i/p data from i/p layer.

o/p layer - The final layer, consists of  $n$  nodes. It produces the final output of the ANN. (Note: the o/p is actual o/p, not predicted o/p)

Weights - It is the connections b/w nodes in diff. layers. The values of  $w$  always lie b/w  $[0, 1]$  ( $0 \leq w \leq 1$ ). The  $w$  vals. are adjusted during training.

Activation func - They introduce non-linearity into the ANN, allowing it to learn complex patterns.

Bias - A.F. contains some error to stop it. A small val. is added to the processed i/p ( $u_k$ ), that is called bias.



of soft or self-organizing feature map.

It is also known as Kohonen map. It is a type of neural net that learns to represent the i/p data in a lower-dimensional space, while preserving the topological relationships b/w the data points.

14/ what is clustering? what are the main parameters for a good clustering technique? what are conventional & fuzzy set theories? Define uniform crossover & single point crossover in GA. <sup>(a)</sup> Done, <sup>(c)</sup> doe, <sup>(d)</sup> done

Main Parameters for good clustering

- i) No. of clusters (too few clusters = overgeneralization, too many = fragmentation)
  - ii) Distance metric (Euclidean, Manhattan etc)
  - iii) clustering algo (choosing the right clustering algo. is very imp)
- (depends on data & desired outcome)

conventional set theory - Here, an element either belongs to a set or it doesn't. This is binary classification.

Fuzzy set theory - It allows for partial membership, where an element can belong to a set to a certain degree. This is more flexible & can handle uncertainty & ambiguity better.

15/ what is PCA & why is it important? Describe each step of PCA by considering a proper example. Give some Real-time applications of neural networks.

<sup>(a)</sup> Done <sup>(b)</sup> Done

Real-time applications of NN

- i) NLP or natural language processing
- ii) Healthcare
- iii) Speech recognition
- iv) Facial recognition
- v) Computers vision & DSP (Digital Image Processing)
- vi) Finance
- vii) Social media
- viii) Chatbots
- ix) Deep learning
- x) Fraud detection

What is Naive Bayes so 'naive'? (a) You came to know that your model is suffering from low bias & high variance. which algo should you use to fix it? why? (c) what do you understand about Type I & Type II errors? (d) what is non linear classification of supervised learning? Explain with an ex.



Q10]

a) Naïve Bayes is called 'naïve' as it makes a strong assumption that all features are conditionally independent given the class label. This means it assumes that the presence or absence of a particular feature doesn't influence the presence or absence of another feature, that is often not true in real-world scenarios.

b) To tackle low bias & high variance, we should use (a regularization method or ensemble method like) RF or Random Forest or Gradient Boosting.

As these methods help to reduce overfitting, that is the cause of high variance, while still maintaining low bias.

c) Non-linear classification is a type of supervised learning where the decision boundary b/w classes is not a ~~straight~~ straight line. For eg, a SVM with a non-linear kernel such as RBF (Radial basis function), that can create complex decision boundaries to

separate data points belonging to diff. classes.

1b) Describe ~~some~~ generative & discriminative ml. techniques. Suppose a genetic algorithm uses chromosomes of the form  $x = abcdefgh$  with a fixed length of eight genes. Each gene can be any digit b/w 0 & 9. Let the fitness of individual  $x$  be calculated as:  $f(x) = (a+b) * (c+d) + (e+f) - (g+h)$ . Let the initial population consist of 4 individuals with the following chromosomes:  $x_1 = 72413532$ ;  $x_2 = 97121601$ ;  $x_3 = 53221285$ ;  $x_4 = 71852494$ . Use the following (i) Evaluate the fitness of each individual, (ii) Cross the fittest 2 individuals using one-point crossover at the middle point, (iii) Evaluate the fitness of the new population with the best 4 chromosomes (2 old & 2 new) (iv) Perform (ii) & (iii) upto 3 iterations.

Generative ml. techniques - It focus on learning the underlying patterns of data to create new data instances that resemble the original data set. It often uses the unsupervised learning for training.

Discriminative ml. techniques - It aims to distinguish b/w diff. categories of data. Primarily used for classification tasks. ~~like~~ It often uses supervised learning for the training.



$$f(h) = (a+b) * (c+d) + (e+f) - (g+h)$$

$$n_1 = 42413532$$

$$n_2 = 97121601$$

$$n_3 = 53221285$$

$$n_4 = 71852494$$

$$f(n_1) = (7+2) * (4+1) + (3+5) - (3+2) = 48$$

$$f(n_2) = (9+7) * (1+2) + (1+6) - (0+1) = 54$$

$$f(n_3) = (5+3) * (2+2) + (1+2) - (8+5) = 22$$

$$f(n_4) = (7+1) * (8+5) + (2+4) - (9+4) = 94$$

$n_4$	$\rightarrow$	Rank
$n_2$	$\rightarrow$	1
$n_1$	$\rightarrow$	2
$n_3$	$\rightarrow$	3
	$\rightarrow$	4

$$n_4: 7185 | 2494$$

$$n_2: 9712 | 1601$$

$$n_1: 71851601$$

$$n_2: 97122494$$

$$n_1: 71851601 \text{ (new)}$$

$$n_2: 97122494$$

$$n_3: 53221285$$

$$n_4: 53221285$$

$$53221285 \text{ (old)}$$

$$f(n_1') = (7+1) * (8+5) + (1+6) - (0+1) = 110$$

$$f(n_2') = (9+7) * (1+2) + (2+4) - (9+4) = 41$$

$$f(n_3') = (7+2) * (4+1) + (3+5) - (3+2) = 48$$

$$f(n_4') = (5+3) * (2+2) + (1+2) - (8+5) = 22$$

$$i+1$$

$$n_1' \rightarrow \text{Rank}$$

$$n_3' \rightarrow 2$$

$$n_2' \rightarrow 3$$

$$n_4' \rightarrow 4$$

$$n_1': 7185 | 1601$$

$$n_3': 7241 | 3532$$

$$\downarrow$$

$$n_1'': 71853532 \text{ (new)}$$

$$n_2'': 72411601$$

$$n_3'': 97122494$$

$$n_4'': 53221285 \text{ (old)}$$

$$f(n_1'') = (7+1) * (8+5) + (3+5) - (3+2) = 107$$

$$f(n_2'') = (7+2) * (4+1) + (1+6) - (0+1) = 51$$

$$f(n_3'') = (9+7) * (1+2) + (2+4) - (9+4) = 41$$

$$f(n_4'') = (5+3) * (2+2) + (1+2) - (8+5) = 22$$

$$i+2$$

$$n_1'' \rightarrow R$$

$$n_2'' \rightarrow 2$$

$$n_3'' \rightarrow 3$$

$$n_4'' \rightarrow 4$$

$$n_1''': 7185 | 3532$$

$$n_2''': 7241 | 1601$$

$$\downarrow$$

$$n_1''': 71851601 \text{ (new)}$$

$$n_2''': 72413532$$

$$n_3''': 97122494$$

$$n_4''': 53221285 \text{ (old)}$$



$$f(h_1^{III}) = (7+1) * (8+5) + (1+6) - (10+1) = 110$$

$$f(h_2^{III}) = (7+2) * (4+1) + (3+5) - (3+2) = 48$$

$$f(h_3^{III}) = (5+7) * (1+2) + (2+4) - (5+4) = 41$$

$$f(h_4^{III}) = (5+3) * (2+2) + (1+2) - (8+5) = 22$$

it=3

$$h_1^{III} \rightarrow 1$$

$$h_2^{III} \rightarrow 2$$

$$h_3^{III} \rightarrow 3$$

$$h_4^{III} \rightarrow 4$$

$$h_1^{III}: 7185 \mid 1601$$

$$h_2^{III}: 7241 \mid 3532$$

↓

$$h_1^{IV}: 71853532 + h_3^{IV}: 57122454$$

$$h_2^{IV}: 7241601 \quad (\text{new}) \quad h_4^{IV}: 53221285$$

(old)

$$f(h_1^{IV}) = (7+1) * (8+5) + (3+5) - (3+2) = 107$$

$$f(h_2^{IV}) = (7+2) * (4+1) + (1+6) - (10+1) = 51$$

$$f(h_3^{IV}) = (5+7) * (1+2) + (2+4) - (5+4) = 41$$

$$f(h_4^{IV}) = (5+3) * (2+2) + (1+2) - (8+5) = 22$$



Data Preprocessing  
(Process of transforming raw data into a clean data set)

ii) Data imputation (missing value imputation)  
[we try to estimate the values of the missing tuples, we can apply in various statistical computation]

A diagram showing a 3x3 grid of squares. An arrow points to the top-right square, which is labeled  $(p)$ . A checkmark is drawn below the label.



✓ When Python the m.l model fails to give us the desired result. This problem is called curse of dimensionality. This can be solved by dimensionality reduction or F.E or F.S. [We will remove the very less imp attribute / feature] that will not give any impact to the target attribute. We will try to find out the less imp feature & will remove it, this is called feature selection (F.S). There are many algos for this. We'll try to approximate <sup>max</sup> no. of attributes into <sup>(the original features)</sup> no. of attributes to get desired result [e.g. 10,000 ~ 100], this called feature execution (F.E). There is an algo for it that is called Principal Component analysis (PCA).

iv) Data transformation (data normalization)  
 In ds, diff. attributes have diff. ranges. If on this we apply the Process it will not give us the desired result. So we have to take them all in same range.

It has 2 types - i) min-max

ii) Standard scaler (Z-score)

i) try to transform the data in b/w (0,1)

ii) try to transform the data in b/w (-1,1)

iii) min-max

$X_i$  = original value  
 $X_i'$  = updated value

$$X_i' = \frac{X_i - X_{\min}}{X_{\max} - X_{\min}}$$

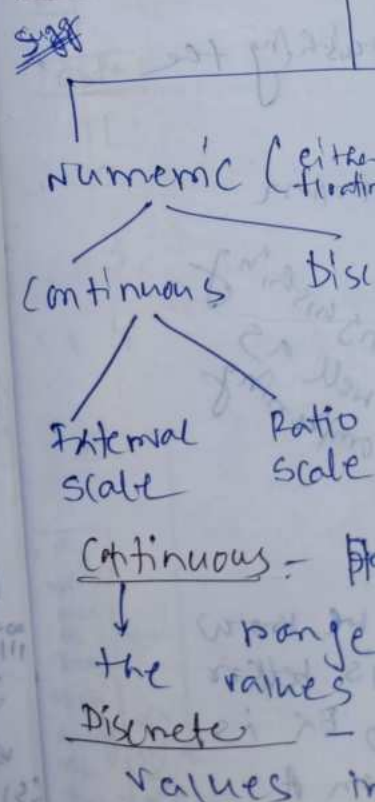
ii) Standard scaler

$$X_i' = \frac{X_i - \mu}{s}$$

Formula for Z-score normalization

We have to convert to the other scale

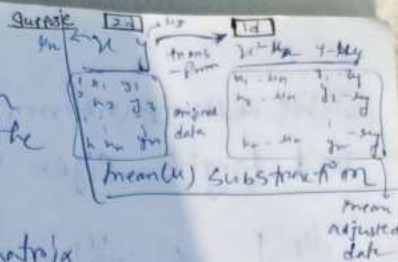
Types of (Data)





# PCA

1. Subtract the mean from each data dim. & find the mean adjusted data.



2. Compute the covariance matrix 'A'.

(this is a type of mat that is used to represent the cov values b/w pairs of elems gives in a random vector)

Covariance mat  
p-dim  
(no. of features)

$$A = \begin{pmatrix} \text{var}(x) & \text{cov}(x, y) \\ \text{cov}(y, x) & \text{var}(y) \end{pmatrix}$$

$$\text{var}(x) = \frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n-1}$$

$$\text{cov}(x, y) = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{n-1}$$

3. Calculate the eigen values & eigen vectors of the covariance matrix 'A'.

if is a symmetric mat.  
if n no. of eigen values & vectors are given, then no. of data set

$$A X = \lambda X$$

$$(A - \lambda I) X = 0$$

if considers,

$A - \lambda = 0$  we will get diff  $\lambda$  values  
 $\lambda_1, \lambda_2, \dots, \lambda_p$

As the data is 2D we will get 2 eigen values -  $\lambda_1, \lambda_2$

4. Arrange the eigen values in descending order from largest to smallest.

5. From eigen values we will find out the eigen vectors.

$$A X = \lambda X$$

$$\begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} u_1 \\ u_2 \end{pmatrix} \text{ eigen vectors}$$

$$\begin{pmatrix} x \\ y \end{pmatrix} = \begin{pmatrix} v_1 \\ v_2 \end{pmatrix} \text{ eigen vectors}$$

As we have 2 eigen vals  $\lambda_1, \lambda_2$  we have to solve this eq for 2 times.

$$A X = \lambda_1 X$$

$$A X = \lambda_2 X$$

we have to find out the value of x & y by using this

$$A = \begin{pmatrix} 1 & 1 \\ 1 & 1 \end{pmatrix} \text{ 2x2 mat.}$$



6. we will transform the eigen vectors into unit length eigen vectors.

$$\begin{pmatrix} u_1' \\ u_2' \end{pmatrix} = \begin{pmatrix} \frac{u_1}{\sqrt{u_1^2 + u_2^2}} \\ \frac{u_2}{\sqrt{u_1^2 + u_2^2}} \end{pmatrix}$$

7. derive the new dataset.

$$\boxed{\text{new dataset} = \text{Row Feature vector} \times \text{Row Data Adjacent}} \\ (\text{dim} = 2 \times N)$$

Feature vector =  $\begin{pmatrix} u_1' & u_2' \\ u_1' & u_2' \\ \vdots & \vdots \\ u_p' & u_p' \end{pmatrix}$   $(P \times P = \text{dim})$   
 [P Eigen values, P eigen vectors]

$$\begin{pmatrix} u_1' & u_2' \\ u_1' & u_2' \\ \vdots & \vdots \\ u_p' & u_p' \end{pmatrix} = X \quad \text{From 2d}$$

Row Feature vector =  $(\text{Feature vector})^T \rightarrow \text{transpose}$   
 $(P \times P)$

Row Data Adj =  $(\text{mean Adj Data})^T \rightarrow \text{transpose}$   
 $2 \times N$

$$\begin{pmatrix} u_1' & u_2' \\ u_1' & u_2' \\ \vdots & \vdots \\ u_p' & u_p' \end{pmatrix} = X \quad \text{From 2d, } 2 \times 2 \text{ Feature vector}$$

Eigen value of a matrix - It is a special set of scalar values that is associated with the set of linear eqns most probably in the matrix eqns.

Eigen vectors of a matrix - It is a vector that is associated with a set of linear eqns. These are also known as latent vectors, Proper vectors or characteristic vectors.

PCA (Dimensionality Reduction)

Let's we have 2D data & how

x	y
2.5	2.4
0.5	0.7
2.2	2.9
1.9	2.2
3.1	3
2.3	2.7
2	1.6
1	1.1
1.5	1.6
1.1	0.9

$$\bar{x} = \frac{1.81}{10} \\ \therefore \text{Mean} = \frac{18.1}{10} = 1.81$$

$$A = \begin{pmatrix} C_{11} & C_{12} \\ C_{21} & C_{22} \end{pmatrix} \quad \text{Covariance matrix}$$

$$\text{Var}(x) =$$

$$\text{Var}(y) =$$



weight, scatter  
data But + discrete  
ml algo (ml algo).  
of those are given, but not of  
logical levels/ops.

(prob)

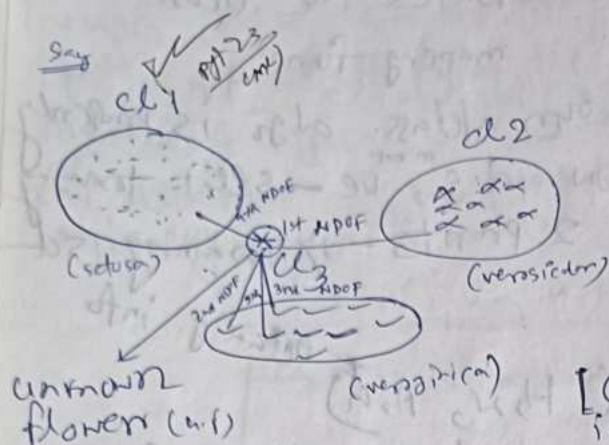
values / continuous values.  
ervised learning

Compare  
to see the accuracy  
of our mapping func / class. algo  
[known,  $Y_A \rightarrow$  Actual  
info]

KNN (K-nearest neighbour classification algo)

$K$  = The no. of neighbour info that we use to  
build our classification model.  
→ [A supervised m.l. algo that uses a distance-based approach to classify / predict  
the grouping of a data point] ||





[cl = class]

let  $k=5$

[N.D.O.F.  
nearest dist  
of flower]

$[c_i = 1, 2, 3]$   $[cl = c]$   
 $i=1-3$

As the NDOF towards  $cl_3$  is the most, so we'll consider that unknown obj falls into  $cl_3$ .

(demonstration)

- Cal. the dis b/w the u.f. & all other flowers
- see, the ~~shortest dist~~ b/w u.f. &  $k$  no. of min (nearest neighbours), here 5 n.n.
- Among the NDOF, we can see,  $\{3, 3, 3, 3, 3\}$  (n.n.)

i.e. the majority of the nearest flowers belong to  $cl_3$ , so that u.f. belongs to 3rd category /  $cl_3$ .

$$\begin{cases} c_1 = \frac{1}{5} = 0.2 \\ c_2 = \frac{1}{5} = 0.2, c_3 = \frac{3}{5} = 0.6 \end{cases}$$

- If we have a tie among the no. of n.n. b/w the classes we can increase the value of  $k$  & see the same (way 1)

eg -  $\{3, 2, 3, 2, 1\}$   
 $d_1, d_2, d_3, d_4, d_5$

Way 2:  $d_1 + d_3$  compare  $d_2 + d_3$   
if  $(d_1 + d_3) < (d_2 + d_3)$ , then  $cl_3 \leftarrow u.f.$   
if  $(d_1 + d_3) > (d_2 + d_3)$ , then  $cl_2 \leftarrow u.f.$

Q23 K-NN Algo

- 1) Store a
- 2) For each
- 3) search
- input obj
- calculate
- For each
- for each
- of obj
- going to

5) The class

Eg-1

Dataset

$S_i$	
1	
2	
3	
4	
5	
6	
7	
8	

[d = distance - cc]



# KNN Algo

- 1/ Store all the input data in the training set.
- 2/ For each object in the test set
  - a) search for  $K$  nearest neighbours to the input obj's / patterns using any distance <sup>(obj's)</sup> <sub>metric</sub> (e.g. euclidean etc)
  - b) For classification we'll compute the confidence for each class as  $C_i/K$  where  $C_i$  is the no. of obj's among the  $K$  nearest neighbours belong to class label 'i'
  - c) The classification for the i/p obj is the class with the highest confidence.

Eg 1) Dataset

$S_i$	CGPA	Assim	Proj m	Result
1	9.2	8.5	8	Pass
2	8	8.0	7	Pass
3	8.5	8.1	8	Pass
4	6	4.5	5	Fail
5	6.5	5.0	4	Fail
6	8.9	7.2	7	Pass
7	5.8	3.8	5	Fail
8	8.9	9.1	9	Pass

(eg of binary classification)  
Class label  
↓  
attribute

Training  
d.s  
(available d.s)



- 1) model designing
- category of machine learning model
- i) Unsupervised learning [clusters] as similar as possible
  - ii) Supervised learning [clustering] as similar as possible
  - iii) Reinforcement learning

iris data - setosa, versicolour, virginica

Petal length, Sepal length & width

50 rows (total 150 rows)

Unsupervised learning - we don't know which obj falls into which category. eg - In above eg data there will be clusters of 3 types of flowers but we can't distinguish that which cluster is of which flower.

Supervised learning - Here we basically train our model, & if there is a unknown obj in the model it can correctly predict that in which category it falls into.

Reinforcement learning - Semisupervised learning. Here step by step we try to improve our ml model to get accurate result. If the model isn't giving accurate result then



Penalize it otherwise give it reward (Save it).

Eg- self driving

Iris data - The iris data are a data frame of 150 measurements of iris petal & sepal length & width, with 50 measurements for each species of 'setosa', 'versicolor' & 'virginica'.

Unsupervised learning - It is a type of m.l. that learns from data without human supervision. Unlike supervised learning, unsupervised m.l. models are given unlabeled data & allowed to discover patterns & insights without any explicit guidance / instruction.

Supervised learning - It is a category of m.l. that uses labeled datasets to train algorithms to predict outcomes & recognize patterns. Unlike unsupervised l., s.l. algos are given labeled training to learn the relationship b/w the i/p & o/rs.

Reinforcement learning - It is a m.l. training method based on rewarding desired behaviours & punishing undesired ones. In general, a reinforcement learning agent - the entity being trained - is able to perceive & interpret its environment, take actions & learn through trial & error.



car 4-3-2 ANN arc  
meters for a good

solution



## 2. Draw a very clear 4-3-2 ANN architecture with explaining all its components.

Component	Explanation
Input Layer	The first layer of the ANN, consisting of 4 neurons. It receives the input data.
Hidden Layer 1	The second layer, consisting of 3 neurons. It processes the input data from the input layer.
Hidden Layer 2	The third layer, consisting of 2 neurons. It further processes the data from the previous layer.
Output Layer	The final layer, consisting of 2 neurons. It produces the final output of the ANN.
Weights	The connections between neurons in different layers. They represent the strength

Are these results  
useful?

Yes

No





(c) What do you understand about Type I & Type II errors?

**Type I error (False Positive):**

This occurs when you reject a true null hypothesis. In other words, you predict something to be true when it's actually false.

**Type II error (False Negative):**

This occurs when you fail to reject a false null hypothesis. In other words, you predict something to be false when it's actually true.