

ASSIGNMENT-II

DSR

1. What is Predictive modeling? Discuss about evaluation of Predictive models?

Ans Predictive modelling is the process of using known results to create, process and validate a model that can be used to forecast future outcome using statistics
 => It is a tool used in predictive Analytics, a data mining technique that is concerned with forecasting probabilities and trends.

=> examples of specific types of forecasting that benefit business are demand forecasting

headcount planning

clean Analysis

competition Analysis

financial risks

Types of predictive modeling

* Regression

* clustering

* Neural networks

- * classification
- * Timeseries model
- * forecasting

Q. What is linear regression & list out the critical assumptions of linear regression?

Ans Linear regression is the supervised machine learning model in which the model finds the best fit linear line between the independent and dependent variables

Critical Assumptions

We can use to understand the relationship b/w two variables x and y
 $y = ax + b$ ①

① Linear relationship

There exists a linear relationship b/w the independent variable x and the dependent variable y .

② Independence

③ Homoscedasticity

④ Normality.

③ Why logistic regression is used for the classification
explain model building strategies for logistic regression

Ans Logistic regression is used as a classification
technique. It uses logistic function to model
the dependent variable

⇒ The dependent variable is dichotomous
in nature

⇒ As a result, this technique is used while dealing
with binary data

Eg:- customers chosen
spam email / website

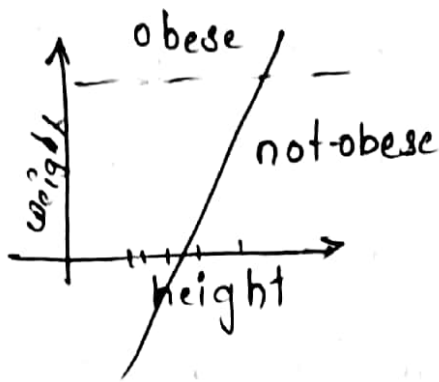
⇒ purposeful selection of variables includes the following

Steps:

- 1) univariable analysis
- 2) multivariable model comparison
- 3) linearity assumption
- 4) interactions among covariate

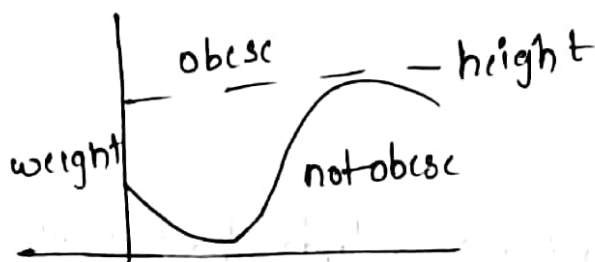
A) Write in detail about linear and
logistic regression

Ans Linear regression is used to predict the
continuous dependent variable using a given
set of independent variables



Logistic Regression is used to predict the categorical dependent variables using a given set of independent variables

$$\log\left(\frac{y}{1-y}\right) = b_0 + b_1x_1 + b_2x_2 + b_3x_3 + \dots + b_nx_n$$



- 5) List out the various control structures supported by R programming language.

Ans control statements are expressions used to control the execution and flow of the program based on the condition provided in the statement

In R programming

- * if-condition
- * if-else
- * for-loop
- * nested-loop

6. Briefly describe the data structures in R-programming language?

Ans A data structure is a particular way of organizing data in a computer so that it can be used effectively
⇒ The most essential data structure used in R includes:

- Vectors
- Lists
- Dataframes
- Matrices
- Arrays
- Factors

* Vectors

A vector is ordered collection of basic type of given length.

```
x = c(1, 3, 5, 7, 8)
```

```
print(x)
```

```
o/p: [1] 1 3 5 7 8
```

Lists

A list is a generic object consisting of an ordered collection of objects

```
l1 = c(1, 2, 3, 4)
```

- * While loop
- * repeat and break statement
- * Return statement
- * next statement
- * if-condition

Syntax

```
if (expression)
{
    statements;
}
```

* for loop

Syntax

```
for (value in vectors)
{
    statements
}
```

* nested loop

```
m ← matrix(2:15, 2)
```

```
for (a in seq(nrow(m)))
```

```
{
  for (c in seq(ncol(m)))
```

```
    {
      print(m[a, c])
```

```
    }
```

```
[1] 2
[1] 4
[1] 6
```

```
[1] 8
[1] 10
[1] 12
```


7. Define Object. List the methods for measuring Distance b/w objects?

Ans Objects

Object are the instance of the class. Also everything and to know more look at datatypes in R.
→ method for measuring distance b/w objects

Euclidean distance

The most common distance is euclidean distance
$$edist(x, y) \leftarrow \sqrt{(x[1] - y[1])^2 + (x[2] - y[2])^2 + \dots}$$

* Manhattan distance

Manhattan distance measure distance in the no. of horizontal and vertical units it takes to get from one point to the other

$$mdist \leftarrow \text{sum}(\text{abs}(x[1] - y[1]) + \text{abs}(x[2] - y[2]) + \dots)$$

* Cosine Similarity

cosine similarity is a common similarity metric in text analysis

$$\text{dot}(x, y) \leftarrow \text{sum}(x[1] \times y[1] + x[2] \times y[2] + \dots)$$

$$\text{cosine}(x, y) \leftarrow \text{dot}(x, y) / \sqrt{\text{dot}(x, x) \times \text{dot}(y, y)}$$

$$\cos(\theta) = \frac{A \cdot B}{|A||B|}$$

Dataframes

Dataframes are generic data objects of R which are used to store tabular form

=> Dataframes are the foremost popular data objects in R programming

=> Each item in a single column must be of same datatype

```
Name = c("Sweety", "Madhu", "Ramani")
```

```
lang = c("R", "python", "C++")
```

```
df = data.frame(Name, lang)
```

```
print(df)
```

Name	lang
Sweety	R
Madhu	python
Ramani	C++

Matrices

A matrix is a rectangular arrangement of numbers in rows and columns

```
A = matrix(c(1, 2, 3, 4, 5, 6, 7, 8, 9))
```

```
nrow = 3, ncol = 3, byrow = TRUE)
```

```
print(A)
```

	[,1]	[,2]	[,3]
[1,]	1	2	3
[2,]	4	5	6
[3,]	7	8	9

8 : Define list and dataframe in R and explain various operations on lists and dataframes with suitable example.

Ans A list is generated using list() function.

Creating lists

```
Geek-list <- list("Geek", TRUE, 123)
```

```
Print(Geek-list)
```

* Naming the elements of a list

Accessing elements of a list

Adding, deleting and updating elements of a list

Merging elements of a list

converting a list to vector

Dataframes

Dataframes are generic data objects of R which are used to store the tabular data

creating a dataframe using vector

Accessing rows and columns

Selecting a subset of the dataframe

Editing dataframes

5 Explain k-Nearest Neighbours Algorithm and its implementation in R programming language.

Ans k-Nearest Neighbors (KNN)

KNN can be defined as a supervised machine learning algorithm that can classify new data point into target class based on neighbouring data points features.

For example: consider a KNN algorithm for a machine that differentiates between apples and mangoes.

To perform this a dataset of apples and mangoes are must be provided as input and then model must be trained in such a way that it detects that fruit based on certain characteristics.

For examples, features of an apple would be red in colour, round in shape etc.

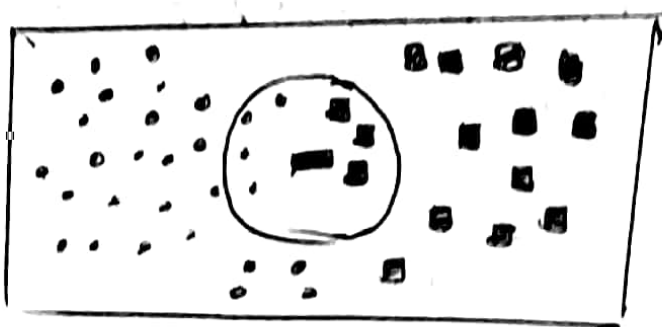
Similarly features of mango would be yellow in colour, oval in shape etc.

An image is provided to the model as input

measures such as Euclidean and Mahalanobis distance are used to classify the closeness b/w data points

When $k=3$ the nearest neighbours are 3 squares and 1 circle dot. When $k=3$ it will be assigned to class B.

But \wedge $k=7$ the neighbours would be 4 dots and 3 squares.



When it is classified the new data point will be assigned to class A.

Implementation of KNN in R programming language.

Consider a problem to analyze the bank data and then build a machine learning model that can predict whether a loan for an applicant can be approved or rejected depending on its socio-economic profile.

The bank credit dataset consists of the information about 1000's of applicants such as name, age, amount balance, loan records etc.

Based upon this data, prediction can be made that whether a loan can be approved or rejected. This

problem can be solved by KNN algorithm by classifying loan application into two classes.

1. Approved
2. Rejected.

Steps to solve this problem as follows -

Step 1: Importing the dataset -

Import DATA the dataset of applicants -

```
loan <- read.csv("C:/users/desktop/DATASET/knn/credits.csv")
```

The structure of data would be

```
str(loan)
```

data frame: 1000 obs. of 2 variables:-

\$ credibility: int 1 1 1 1 1 1 1 - -

\$ Account Balance: int 12 11 11 11 11 11 11

\$ Duration of credits month: int 18 9 12 12 10 8 6 18 24

\$ purpose: int 20 9 0 0 0 1 2 3

\$ credit amount: int 1049 2799 841 122 2171

\$ values: savings: stocks ? int 1 1 2 1 1 1 1 3 - -

\$ Length of current employment: int 2 3 4 3 3 2 4 2 1

\$ instalment percent: int 4 2 2 3 4 1 2 4 1

\$ marital status: int 4 2 3 2 3 2 2 1 1

\$ Guarantors: int 1 1 1 1 1 1 1 - -

\$ Duration in current address: int 4 2 4 2 4 3 4 4 4 -

\$ Age: years: int 21 36 23 39 38 40

\$ No. of dependices: int 1 2 1 1 2 1 1 1

\$ Telephone: int 1 1 1 1 1 - -

Step: Data cleaning

The structure of dataset consists of ²¹ predictable variables that are used to decide whether loan for applicant can be approved or rejected.

Some of the variables might not be useful in predicting the loan. For example - credit concurrent credits telephones etc. can be removed because such type of variable leads to complexity of machine learning model.

`loan_subset = loan[['Credibility', 'Age', 'years', 'Account balance']]`

The dataset now would be

`dataframe (loan_subset)`

'dataframe' 10000s of 8 variables

\$credibility

\$age - years

\$ Marital status

\$ Occupation

\$ Account balance

\$ length of current employment

\$ purpose

int 1 1 1 1 1 1

int 29 30 49

int 2 3 1 1 1

int 3929 4232 - -

int 1 2 2 1 1 4 2 1

int 23 43 32 44 1

int 20 90 000 33 - -

Steps: Data Normalization

Normalization of data set is mandatory to make the output unbiased consider below observations

head(loan_subset)

	credibility	age_years	Maritalstatus	Occupation	Account Balance
1	1	49	4	1	1
2	1	63	3	1	2
3	1	78	2	2	4
4	1	58	1	1	3
5	1	15	1	1	20

credit amount	Length of employment	Purpose
2943	2	2
5678	3	0
9629	4	9
1423	3	0
2428	3	6
1928	2	6

Here credit amount variable has value scale in 1000s and the remaining are in single or two digits. If the data is not normalized outcome is in biased form

$\text{normalize} \leftarrow \text{function}(x)$

$\text{return} (x - \min(x)) / (\max(x) - \min(x))$

Now store the normalized data set in loan_subset and then remove credibility variable because it is

Explain K-Means Algorithm and its implementation in R programming

K-means

The K-means is an iterative clustering algorithm in which objects are moved ~~over~~ among set of clusters until desired set is achieved.

It is most popular and commonly used method. The algorithm is based on the concept of most specified input parameter (k). A set of n objects are divided into k clusters. A high degree of similarity among objects. elements in clusters is obtained.

Working of K-means Algorithm

The number of clusters input
A dataset containing n objects are denoted by k .
A dataset containing n objects denoted by D .

Algorithm's output

k : A set consisting of k clusters.

Procedure:

- Step 1: Initially select k objects randomly from D , as initial cluster centers
- Step 2: Depending upon the distance between the object and the cluster mean, each remaining object assigned to cluster to which it is most similar.

Euclidean distance b/w mean and objects to
classify objects into 4 clusters

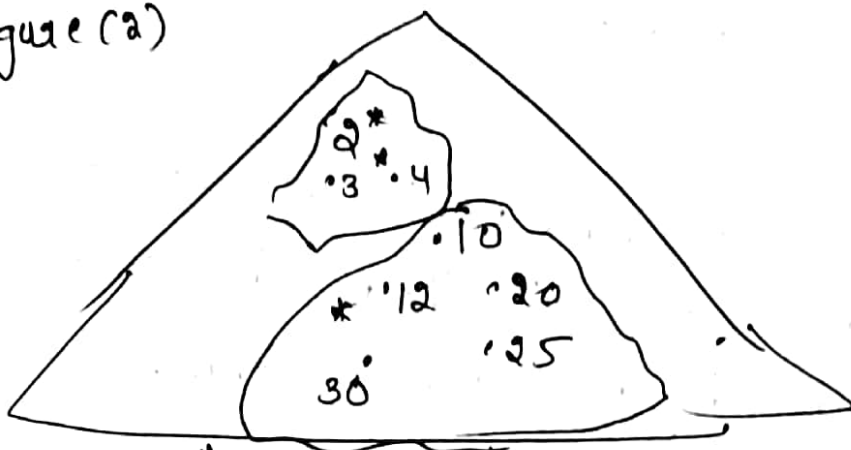


Fig(1) initial partitioning with $m_1 = 2$ and $m_2 = 4$
Now evaluate new mean for the resulting clusters
and again partition is done base of euclidean
distance as follows

$$m_1 = \frac{2+3}{2} = 2.5$$

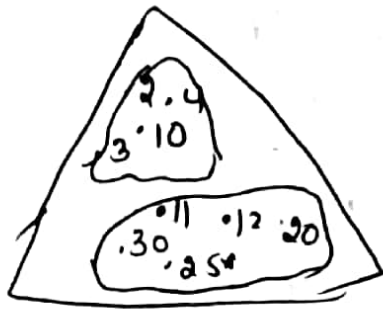
$$m_2 = \frac{4+10+11+12+20+25+30}{7} = 16$$

$m_1 = 2.5$ and $m_2 = 16$ and resulting cluster is shown
in Figure (2)



Fig(2) ^{new} clusters formed with $m_1 = 2.5$ and $m_2 = 16$.

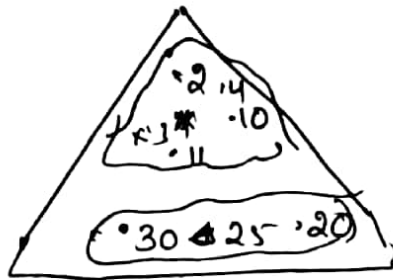
The process iterates for the successive values of the mean
to enhance the partition of clusters



clusters with $m_1 = 3$ and $m_2 = 18$



clusters with $m_1 = 4.75$ and $m_2 = 19.5$



clusters with $m_1 = 7$ and $m_2 = 25$

Thus finally the correct partition of the numbered objects into two clusters is achieved.

K-Means Implementation in R

The K-means algorithm can be implemented by predefining. Here K represents number of clusters to be defined.

Steps to implement K-means are as follows.

1. In the first step K centres are defined and every cluster is assigned to the cluster that has closest centre to it.

2. In the second step the centres are redefined by using the observation of each cluster. The column means are used for defining the centroid.

The above two steps are repeated until the centres are converge. Therefore K-means algorithm said to be iterative.

List out the various performance metrics for classification?

Performance Measures

Performance of a classification model is evaluated once the predictive model is built. It determines whether the model can predict the outcome of new observation test data that is not used in the training the model. The performance of a model can be assessed by comparing the predicted outcome values against known outcome values. The commonly used metrics and methods to assess the performance of predictive classification models are as follows.

1. Average classification accuracy

It represents the proportion of correctly classified observation. The complete classification rate corresponds to the part of observations that are classified correctly. The first step of assessing the performance of a model is to determine the average classification accuracy.

The average classification error rate can be inversely defined as proportion of observation that are classified incorrectly.

$$\text{error rate} = 1 - \text{accuracy}$$

The row classification accuracy and error can be determined by comparing the observed classes in test data and predicted classes by model

$\gamma_{accuracy} \leftarrow \text{mean}(\text{observed class} == \text{predicted class})$

$\gamma_{accuracy}$

[1] 0.08

$\gamma_{error} \leftarrow \text{mean}(\text{observed class} != \text{predicted class})$

γ_{error}

[1] 0.192

The binary classifier in this example makes two types of error

\Rightarrow It can wrongly class assign individual who has diabetes positive to diabetes negative

\Rightarrow It can wrongly assign individual who has diabetes negative to diabetes positive

2. Confusion matrix

The main factor required for performance evaluation of ~~most~~ classification models includes correct or incorrect predictions of no. of test cases by the model. The test records are given in a tabular form known as confusion matrix.

Actual class	Predicted class	
	class y	class x
class y	fy y	fy x
class x	fx y	fx x

The above specified table outlines the binary classification problem and each individual entry is given as $fx y$. Here x represents no of records belongs to class x and y represents no of records belongs to class y .

$fx x$
This entry specifies the no of records belonging to class ' x '. There is no mis prediction in this entry, since the records of class ' x ' are predicted as records from class ' x '.

$fx y$
This entry specifies the no of records belonging to class ' x '. There is mis prediction in this entry since the records of class x are incorrectly predicted as records from class ' y '.

$fy y$
This entry specifies the no of records belonging to class ' y '. There is no mis prediction in this entry.

A) ROC curve

The graphical representation that exhibits the performance of predicted classification model is known as receiver operating characteristics curve.

Applications of ROC curve

ROC curves depicts the tradeoff between proportion of correctly determined positive tuples known as true positive rate and the proportion of negative tuples incorrectly tuples as positive tuples known as negative rate for given model.

