<u>DATA MINING</u> (cocsc16)



PRACTICAL FILE

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COE-1

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EXPERIMENT -1

AIM:

Visualize your dataset in Weka and learn about Jitter.

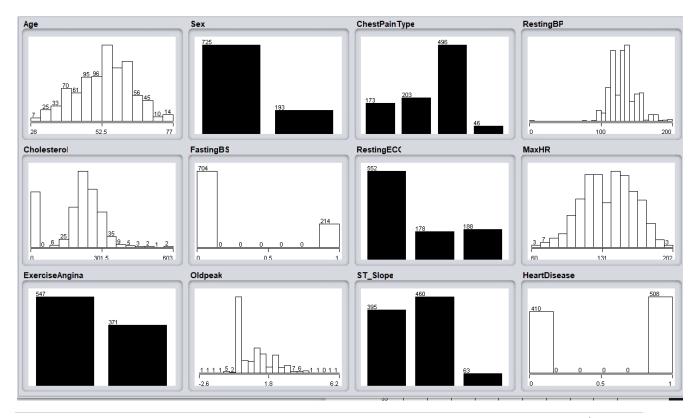
DATASET:

Heart Disease Prediction

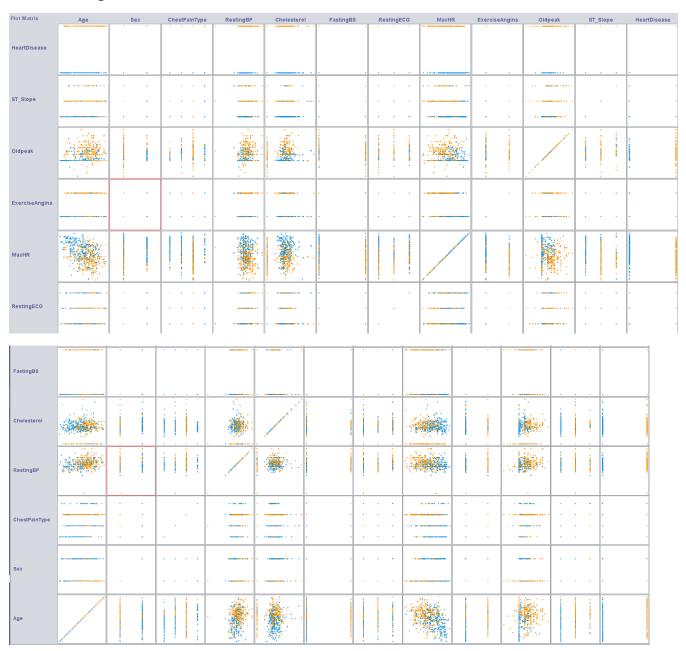
RESULTS:

Loading the dataset Heart Disease Prediction in Weka and Visualizing in the pre-processing step:

Dataset:

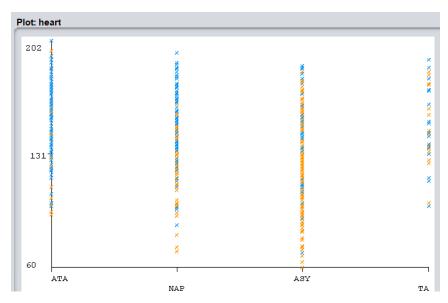


Visualizing the relation between attributes:

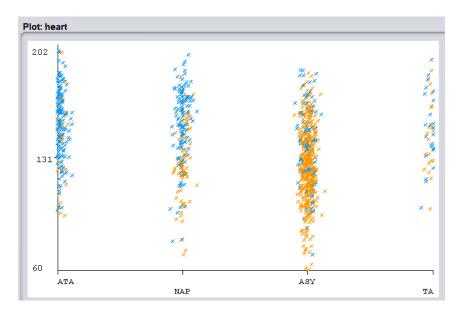


Understanding Jitter:

In a plot between MaxHR and ChestPainType, with no jitter:



With considerable Jitter:



Jitter is a function which adds artificial random noise to the coordinates of the potted points in order to spread the data out so that overlapping points become more visible thus helping in the understanding of dataset.

EXPERIMENT -2

AIM:

Calculate co-relation among attributes

DATASET:

Heart Disease Prediction

RESULTS:

Calculating Person's coefficient between attribute:

```
Search Method:
       Attribute ranking.
Attribute Evaluator (supervised, Class (nominal): 12 HeartDisease):
       Correlation Ranking Filter
Ranked attributes:
 0.5538 11 ST Slope
 0.4943 9 ExerciseAngina
 0.4048 3 ChestPainType
 0.404 10 Oldpeak
 0.4004 8 MaxHR
 0.3054 2 Sex
 0.282 1 Age
 0.2673 6 FastingBS
 0.2327 5 Cholesterol
 0.1076 4 RestingBP
 0.0771 7 RestingECG
Selected attributes: 11,9,3,10,8,2,1,6,5,4,7 : 11
```

By finding the Pearson's correlation and evaluating the worth of an attribute, we can eliminate redundant attributes thus preventing curse of dimensionality.

Since RestingECG and RestingBP are ranked least, removing these attribute from consideration

EXPEIMENT 3

AIM:

Calculate Information Gain of different attributes

DATASET:

Heart Disease Detection

RESULTS:

Calculating Information Gained by each attribute for prediction heart disease.

```
Search Method:
       Attribute ranking.
Attribute Evaluator (supervised, Class (nominal): 10 HeartDisease):
       Information Gain Ranking Filter
Ranked attributes:
0.2993 9 ST Slope
 0.225 3 ChestPainType
 0.19 7 ExerciseAngina
 0.1614 8 Oldpeak
 0.1275 6 MaxHR
 0.0834 4 Cholesterol
 0.0696 1 Age
 0.0685 2 Sex
0.0549
        5 FastingBS
Selected attributes: 9,3,7,8,6,4,1,2,5 : 9
```

CONCLUSION:

ST_Slope continues to provide maximum information for predicting heart disease. While FastingBS, Sex, and Age are low indicators as per information gain, PCA and correlation analysis show a decent score hence are considered as features for prediction of heart disease.

EXPERIMENT 4:

AIM:

Perform PCA on a dataset

DATASET:

Heart Disease Detection

RESULTS:

Performing Principal Component Analysis and Ranking on the basis of variance:

By performing a PCA, a set of combinations of attributes are obtained along with the respective variance in data that they represent. By analysing, we can see ST_slope an ChestPainType capturing maximum amount of variance in data.

EXPERIMENT 5:

AIM:

Training a Decision Tree

DATASET:

Heart Disease Detection

THEORY:

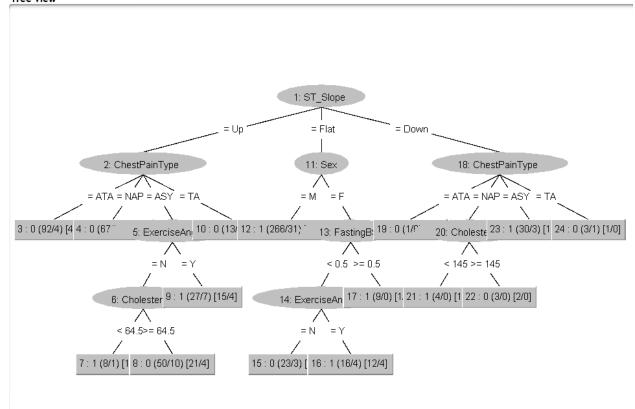
Decision Tree User: REPTree (Reduced Error Pruning Tree) a fast decision tree learner that builds a decision/ regression tree using Information gain as the splitting criterion and prunes it using reduced error pruning algorithm.

Parameters for the Decision tree:



Visualization:

Tree View



RESULTS:

```
=== Summary ===
Correctly Classified Instances 760
Incorrectly Classified Instances 158
                                                               82.7887 %
                                                               17.2113 %
Kappa statistic
                                            0.6492
                                             0.2357
Mean absolute error
Root mean squared error
                                          47.684 %
Relative absolute error
Root relative squared error
                                            72.8761 %
Total Number of Instances
=== Detailed Accuracy By Class ===
                   TP Rate FP Rate Precision Recall F-Measure MCC
                                                                                  ROC Area PRC Area Class
                0.768  0.124  0.833  0.768  0.799  0.651  0.876  0.843  0
0.876  0.232  0.824  0.876  0.849  0.651  0.876  0.854  1
0.828  0.184  0.828  0.828  0.827  0.651  0.876  0.849
Weighted Avg.
=== Confusion Matrix ===
   a b <-- classified as
 315 95 | a = 0
 63 445 | b = 1
```

CONCLUSION:

By using a decision tree classifier, an accuracy of 82.8% was obtained.

EXPERIMENT 6

AIM:

Train a Naïve Bayes Classifier

DATASET:

Heart Disease Detection

PERFORMANCE:

Upon performing the classification through use of naïve bayes classifier: bayes. NaïveBayes the following result was obtained:

```
=== Summary ===
Correctly Classified Instances 795
Incorrectly Classified Instances 123
                                                   86.6013 %
                                                    13.3987 %
                                     0.7282
Kappa statistic
Mean absolute error
                                     0.1566
Root mean squared error
                                     0.338
Relative absolute error
                                    31.6738 %
Root relative squared error
                                    67.9811 %
Total Number of Instances
                                    918
=== Detailed Accuracy By Class ===
               TP Rate FP Rate Precision Recall F-Measure MCC ROC Area PRC Area Class
               0.837 0.110 0.860 0.837 0.848 0.728 0.919 0.911 0
0.890 0.163 0.871 0.890 0.880 0.728 0.919 0.916 1
Weighted Avg. 0.866 0.140 0.866 0.866 0.866 0.728 0.919 0.914
=== Confusion Matrix ===
  a b <-- classified as
 343 67 | a = 0
 56 452 | b = 1
```



An accuracy of 86.6% was obtained. This result is better than the classification accuracy of 82.8% obtained by decision trees.

EXPERIMENT 7

AIM:

Train a Bayesian Belief Network Classifier

DATASET:

Heart Disease Detection

PERFORMANCE:

Classification is performed using the inbuilt bayes.BayesNet

```
=== Summary ===
Incorrectly Classified Instances 139
Kappa statistic
                                                          84.8584 %
                                                          15.1416 %
                                         0.6935
Mean absolute error
                                         0.1682
Root mean squared error
                                         0.3459
Relative absolute error
                                        34.0285 %
Root relative squared error
                                        69.58 %
Total Number of Instances
=== Detailed Accuracy By Class ===
                 TP Rate FP Rate Precision Recall F-Measure MCC ROC Area PRC Area Class
0.827 0.134 0.833 0.827 0.830 0.693 0.915 0.908 0
0.866 0.173 0.861 0.866 0.864 0.693 0.915 0.919 1
Weighted Avg. 0.849 0.156 0.848 0.849 0.849 0.693 0.915 0.914
=== Confusion Matrix ===
      b <-- classified as
 339 71 | a = 0
  68 440 | b = 1
```

RESULT:

Accuracy of 84.8 % was obtained.

EXPERIMENT 8:

AIM:

Train a Linear Regression Model

DATASET:

Heart Disease Detection

PERFORMANCE:

```
Linear Regression Model
HeartDisease =
      0.003 * Age +
      0.1618 * Sex=M +
     0.2357 * ChestPainType=ASY +
     -0.0005 * Cholesterol +
      0.1317 * FastingBS +
     0.1396 * ExerciseAngina=Y +
     0.0499 * Oldpeak +
      0.2196 * ST_Slope=Down,Flat +
     0.164 * ST_Slope=Flat +
     -0.0996
Time taken to build model: 0.09 seconds
=== Cross-validation ===
=== Summary ===
Correlation coefficient
                                        0.7476
Mean absolute error
                                        0.2444
                                        0.3302
Root mean squared error
                                       49.4224 %
Relative absolute error
Root relative squared error
                                      66.3966 %
                                     918
Total Number of Instances
```

RESULT:

Since the task was that of classification, the class labels had a value of 0 or 1, a prediction anywhere in between would count as an error thus such a performance was expected since a regression model was trained for a classification task.

EXPERIMENT 9:

AIM:

Logistic Regression Model

DATASET:

Heart Disease Detection

PERFORMANCE:

```
=== Summary ===
                                                  86.6013 %
Correctly Classified Instances 795
Incorrectly Classified Instances 123
                                                    13.3987 %
                                    0.7277
Kappa statistic
Mean absolute error
                                     0.2007
Root mean squared error
                                     0.3207
                                    40.5949 %
Relative absolute error
Root relative squared error
                                    64.506 %
Total Number of Instances
=== Detailed Accuracy By Class ===
                TP Rate FP Rate Precision Recall F-Measure MCC
                                                                    ROC Area PRC Area Class
               0.827 0.102 0.867 0.827 0.846 0.728 0.924 0.916 0
0.898 0.173 0.865 0.898 0.881 0.728 0.924 0.929 1
Weighted Avg. 0.866 0.142 0.866 0.866 0.866 0.728 0.924 0.923
=== Confusion Matrix ===
  a b <-- classified as
 339 71 | a = 0
 52 456 | b = 1
```

RESULT:

An accuracy of 86.6% was obtained along with a recall of 86.6%.

EXPERIMENT 10:

AIM:

KNN Model

DATASET:

Heart Disease Detection

PERFORMANCE:

```
=== Summary ===

Correctly Classified Instances 735 80.0654 %
Incorrectly Classified Instances 183 19.9346 %

Kappa statistic 0.5979

Mean absolute error 0.2001

Root mean squared error 0.4459

Relative absolute error 40.4746 %

Root relative squared error 89.7013 %

Total Number of Instances 918

=== Detailed Accuracy By Class ===
```

	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	PRC Area	Class
	0.793	0.193	0.768	0.793	0.780	0.598	0.798	0.702	0
	0.807	0.207	0.828	0.807	0.818	0.598	0.798	0.781	1
Weighted Avg.	0.801	0.201	0.802	0.801	0.801	0.598	0.798	0.746	

```
=== Confusion Matrix ===
```

```
a b <-- classified as
325 85 | a = 0
98 410 | b = 1
```

RESULT:

An accuracy of 80% was obtained.

EXPERIMENT 11:

AIM:

K-Means Clustering Model

DATASET:

Heart Disease Detection

PERFORMANCE:

Final cluster centroids:

		Cluster#	
Attribute	Full Data	0	1
	(918.0)	(502.0)	(416.0)
=======================================	========	========	=======
Age	53.5109	56.012	50.4928
Sex	M	M	M
ChestPainType	ASY	ASY	ATA
Cholesterol	198.7996	175.0896	227.4111
FastingBS	0.2331	0.3367	0.1082
MaxHR	136.8094	125.8486	150.0361
ExerciseAngina	N	Y	N
Oldpeak	0.8874	1.3131	0.3736
ST_Slope	Flat	Flat	Up
HeartDisease	1	1	0

```
Time taken to build model (full training data): 0.02 second

=== Model and evaluation on training set ===
```

Clustered Instances

```
0 502 (55%)
1 416 (45%)
```

EXPERIMENT 12:

Q. Knowing the Types of data – ordinal, nominal, ratio, interval.

Nominal

- o Nominal scales are used for labeling variables, without any quantitative value. "Nominal" scales could simply be called "labels."
- o all of the scales shown below are mutually exclusive (no overlap) and none of them have any numerical significance.
- o Properties: distinctness (= != are meaningful)
- o Ex. ID Numbers, eye color, Zip Codes

• Ordinal

- o With ordinal scales, the order of the values is what's important and significant, but the differences between each one is not really known.
- o Ordinal scales are typically measures of non-numeric concepts like satisfaction, happiness, discomfort, etc.
- o distinctness and order, (= != <>are meaningful)
- o Ex. Grades, ranks etc.

Interval

o Interval scales are numeric scales in which we know both the order and the exact differences between the values.

- o For interval attributes, differences between values are meaningful.
- o Properties: distinctness, order and differences, (= != < > + are meaningful)
- o Ex. Calender dates, temperature in Celsius
- Ratio
- o For ratio attributes, both differences and ratio are meaningful.
- o They have an absolute 0.
- o Properties: distinctness, order, differences and ratios (= != < > + * / are meaningful)
- o Ex. Temperature in Kelvin, age, mass.

EXPERIMENT 13:

Q. Find the mean, median, variance, and standard deviation of data.

```
#include <bits/stdc++.h>
using namespace std;
int main() {
    ifstream input file("numeric data.csv");
    if (!input file.is open())
        cout << "ERROR\n";
    vector<vector<string>> data;
    while (input file.good()) {
        string str;
        getline(input file, str, '\n');
        string s = str;
        vector<string> v;
        stringstream ss(s);
        while (ss.good()) {
            string substr;
            getline(ss, substr, ',');
            v.push back(substr);
        data.push back(v);
    }
    input file.close();
    int n = data.size();
    int m = data[0].size();
    cout<<"This input csv file is an "<<n<<"x"<<m<<" dataset\n";</pre>
    for(int j=0;j<m;j++) {</pre>
        double mean = 0;
```

```
for (int i = 0; i < n; i++) {
        mean += stoi(data[i][j]);
    }
    mean = mean * 1.0/n;
    double variance = 0;
    for (int i = 0; i < n; i++) {
        double x = abs(mean - stoi(data[i][j]));
        x = x * x;
        variance += x;
    variance /= (double)n;
    double std deviation = sqrt(variance);
    vector<int> v;
    for (int i = 0; i < n; i++) {
        v.push back(stoi(data[i][j]));
    sort(v.begin(), v.end());
    double median = 0;
    if (n & 1) {
        int idx = (n + 1) / 2;
        idx--;
        median = v[idx];
    }
    else {
        int idx = n/2;
        median = v[idx];
        median += v[idx - 1];
        median /= (double)2;
    cout<<"Measures for "<<j<<" feature : "<<endl;</pre>
    cout << "mean" << " : " << mean << endl;</pre>
    cout << "median" << " : " << median << endl;</pre>
    cout << "variance" << " : " << variance << endl;</pre>
    cout << "standard deviation" << " : " << std deviation << endl;</pre>
return 0;
```

}

Input File:

"numeric_data.csv"

6,11,12,1

125,15,3,5

42,23,63,21

1,63,123,6

4,27,63,3

82,23,95,27

58,74,12,234

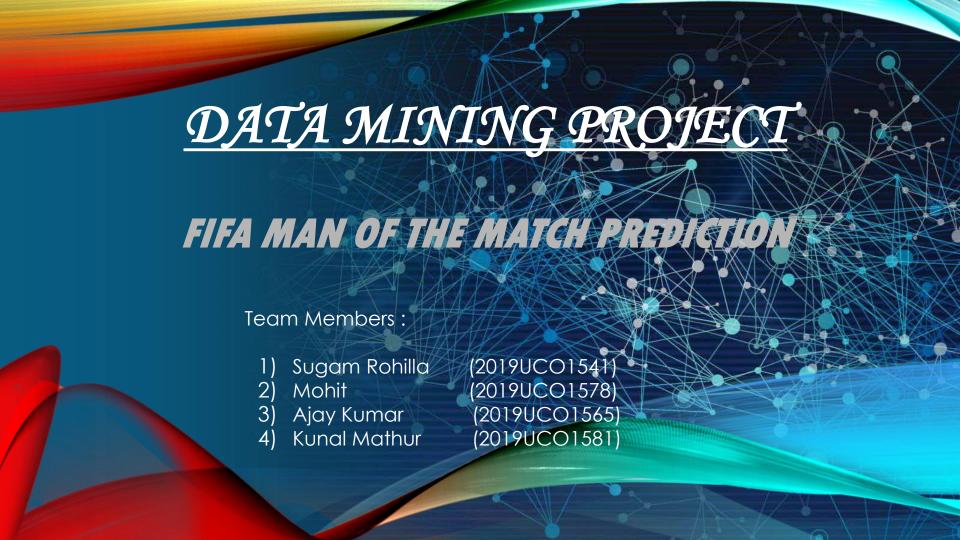
2,12,753,8

1,411,2,12

47,69,6,7

OUTPUT:

This input csv file is an 3x5 dataset Measures for 0 feature : mean : 6 median : 6 variance : 16.6667 standard deviation : 4.08248 Measures for 1 feature : nean : 7 median : 7 variance : 16.6667 standard deviation : 4.08248 Measures for 2 feature : mean : 8 median : 8 variance : 16.6667 standard deviation : 4.08248 Measures for 3 feature : mean: 9 median : 9 variance : 16.6667 standard deviation : 4.08248 Measures for 4 feature : mean : 9.66667 median : 10 variance : 20.2222 standard deviation : 4.49691





- The objective of this project is to predict the Team with Man Of The Match award before the official announcement that will be made right after the match.
- Dataset consists of 26 Predictor Variables (Independent) and one Outcome Variable (Dependent).
- The outcome variable value is either Yes or No indicating whether a person from the team wins the man of the match or not.

FEATURES

	No. 1 AVE CO. 1
Column	Description
Date	match Date
Team	Playing Team
Opponent	Opponent Team
Goal Scored	Number of goals scored by this team
Ball Possession %	Amount of time ball was in control by the team
Attempts	Number of attempts to score goal
On-Target	Number of shots on-target
Off-Target	Number of shots that went off-target
Blocked	Number of attempts blocked by the opponent's team
Corners	Number of corner shots used
Offsides	Number of off-side events
Free Kicks	Number of free-kicks used
Saves	Number of saves by the goal keeper
Pass Accuracy %	Percentage of passes that reached the same team player as aimed
Passes	Total number of passes by the team
Distance Covered (Kms)	Total distance covered by the team members in this game
Fouls Committed	Number of fouls committed by the team members
Yellow Card	Number of Yellow warning received
Yellow & Red	Number of Yellow & Red warning received
Red	Number of Red cards received



Features

Own goals Own goal Time	Number of own goals When did the team score own goal?
Goals in PSO	Number of goals scored in the Penalty shootout
PSO	Was there a penalty shootout (PSO) in this match?
Round	Stage of the match
1st Goal	When did the team score the 1st goal?
Man of the Match	Did this team member win Man of the Match?

<u>Target Feature</u>:- Man of the Match (Did this team member win Man of the Match?.

Exploratory Data Analysis

```
In [30]: data_df = pd.read_csv('fifa.csv')
    data_df.head()
```

Out[30]:

	Date	Team	Opponent	Goal Scored	Ball Possession %	Attempts	On- Target	Off- Target	Blocked	Corners	 Yellow Card	Yellow & Red	Red	Man of the Match	1st Goal	Round	PSO	Goals in PSO	O go
0	14- 06- 2018	Russia	Saudi Arabia	5	40	13	7	3	3	6	 0	0	0	Yes	12.0	Group Stage	No	0	Ζ
1	14- 06- 2018	Saudi Arabia	Russia	0	60	6	0	3	3	2	 0	0	0	No	NaN	Group Stage	No	0	N
2	15- 06- 2018	Egypt	Uruguay	0	43	8	3	3	2	0	 2	0	0	No	NaN	Group Stage	No	0	N
3	15- 06- 2018	Uruguay	Egypt	1	57	14	4	6	4	5	 0	0	0	Yes	89.0	Group Stage	No	0	N
4	15- 06- 2018	Morocco	Iran	0	64	13	3	6	4	5	 1	0	0	No	NaN	Group Stage	No	0	

5 rows × 27 columns



Data preprocessing is the process of transforming raw data into an understandable format. It is also an important step in data mining as we cannot work with raw data. The quality of the data should be checked before applying machine learning or data mining algorithms.

<u> Major Tasks in Data Preprocessing:</u>

- 1. Data Cleaning
- 2. Data integration
- 3. Data reduction
- 4. Data transformation

DATA CLEANING

Data cleaning is the process to remove incorrect data, incomplete data and inaccurate data from the datasets, and it also replaces the missing values.

Removing Duplicates:

```
In [32]: data_df.drop_duplicates()
```

Handling Missing/NULL values:

Data Reduction

This process helps in the reduction of the volume of the data which makes the analysis easier yet produces the same or almost the same result. This reduction also helps to reduce storage space.

DELETING UNWANTED COLUMNS:

```
In [34]: training_features = data_df.drop(['Date', 'Team', 'Opponent', 'Ball Possession %', 'Round', 'Offsides', 'Passes', 'Saves'
    'Distance Covered (Kms)', 'Yellow Card', 'Yellow & Red', 'Red', 'PSO', 'Goals in PSO', 'Own goal Time', '1st Goal',
    'Own goals'], axis = 1)
    training_features.head(3)|
```

Out[34]:

	Goal Scored	Attempts	On-Target	Off-Target	Blocked	Corners	Free Kicks	Pass Accuracy %	Fouls Committed	Man of the Match
0	5	13	7	3	3	6	11	78	22	Yes
1	0	6	0	3	3	2	25	86	10	No
2	0	8	3	3	2	0	7	78	12	No

Replacing Yes & No with 1 & 0 respectively (Man of the match:-

In [7]: training_features['Man of the Match'].replace(['Yes', 'No'], [1, 0], inplace=True)

Out[7]:

	Goal Scored	Attempts	On-Target	Off-Target	Blocked	Corners	Free Kicks	Pass Accuracy %	Fouls Committed	Man of the Match
0	5	13	7	3	3	6	11	78	22	1
1	0	6	0	3	3	2	25	86	10	0

Merging and Merging Attribute :-

Out[38]:

	Goal Scored	Attempts	On- Target	Off- Target	Blocked	Corners	Free Kicks	Pass Accuracy %	Fouls Committed	Man of the Match	Fouls Advantage	Efficient Attempts
0	5	13	7	3	3	6	11	78	22	1	-11	53.8
1	0	6	0	3	3	2	25	86	10	0	15	0.0
2	0	8	3	3	2	0	7	78	12	0	-5	37.5

In [39]: training_features['Fouls Advantage'] = training_features['Free Kicks'] - training_features['Fouls Committed']
training_features.head(3)

Out[39]:

	Goal Scored	Attempts	On- Target	Off- Target	Blocked	Corners	Free Kicks	Pass Accuracy %	Fouls Committed	Man of the Match	Fouls Advantage	Efficient Attempts
0	5	13	7	3	3	6	11	78	22	1	-11	53.8
1	0	6	0	3	3	2	25	86	10	0	15	0.0
2	0	8	3	3	2	0	7	78	12	0	-5	37.5

DELETE UNWANTED COLUMNS AFTER FEATURE ENGINEERING

In [40]: training_features.drop(['On-Target', 'Off-Target', 'Blocked', 'Attempts', 'Free Kicks', 'Fouls Committed'], axis = 1, i
training_features.head(3)

Out[40]:

	Goal Scored	Corners	Pass Accuracy %	Man of the Match	Fouls Advantage	Efficient Attempts
0	5	6	78	1	-11	53.8
1	0	2	86	0	15	0.0
2	0	0	78	0	-5	37.5

Data Visualisation

```
In [12]: corr = training_features.corr()
plt.figure(figsize=(10,8))
    sns.heatmap(corr, vmax=.8, linewidths=0.01, square=False, annot=True, cmap='YlGnBu', linecolor='Black')
plt.title('Correlation between features');
```

- 0.2

- 0.0

		Correlation between features					
Goal Scored	1	0.04	0.14	0.52	0.49	0.013	
Corners	0.04	1	0.33	0.17	-0.13	0.18	
Pass Accuracy %	0.14	0.33	1	0.11	0.03	0.34	
Man of the Match	0.52	0.17	0.11	1	0.24	0.18	
Efficient Attempts	0.49	-0.13	0.03	0.24	1	-0.0098	
Fouls Advantage	0.013	0.18	0.34	0.18	-0.0098	1	
,	Goal Scored	Comers	Pass Accuracy %	Man of the Match	Efficient Attempts	Fouls Advantage	

Visualizing correlation between features using heatmap

Scaling/Normalization

It is basically Data transformation in data preprocessing. It is done to normalize range of independent variables or feature of data.

Out[43]:

	Goal Scored	Corners	Pass Accuracy %	Man of the Match	Fouls Advantage	Efficient Attempts
0	3.194193	0.525857	-0.770604	1	-1.955757	1.624259
1	-1.146112	-1.115843	0.582910	0	2.163711	-2.239935
2	-1.146112	-1.936693	-0.770604	0	-1.005111	0.453509
3	-0.278051	0.115432	0.582910	1	0.896182	-0.185735
4	-1.146112	0.115432	0.582910	0	-1.480434	-0.580774

SPLITTING DATA

Split training features into X & y using pandas)

```
In [44]: X = training_features.loc[:, training_features.columns != 'Man of the Match']
y = training_features['Man of the Match']
```

Split X & y for training and testing

```
In [67]: from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2)
print('Train dataset shape: {} {}'.format(X_train.shape, y_train.shape))
print('Test dataset shape: {} {}'.format(X_test.shape, y_test.shape))

Train dataset shape: (102, 5) (102,)
Test dataset shape: (26, 5) (26,)
```

Split the data into

- 1. Training set (80%)
- 2. Test set (20%)



We used following learning models:

- 1. Logistic Regression
- 2. Decision Tree classifier
- 3. RandomForest classifier

```
In [68]: from sklearn.linear_model import LogisticRegression
    from sklearn.tree import DecisionTreeClassifier
    from sklearn.model_selection import GridSearchCV, RandomizedSearchCV
    from sklearn.ensemble import RandomForestClassifier
```

Logistic Regression

Logistic regression is used for solving the classification problems. It is a significant machine learning algorithm because it has the ability to provide probabilities and classify new data using continuous and discrete datasets. Since it predicts the output of a categorical dependent variable, the outcome must be a categorical or discrete value. It can be either Yes or No, 0 or 1, true or False, etc. but instead of giving the exact value as 0 and 1, it gives the probabilistic values which lie between 0 and 1.

```
In [69]: model_1 = LogisticRegression()
model_1.fit(X_train, y_train)
Out[69]: LogisticRegression()
```

DECISION TREE CLASSIFIER

It is a Supervised learning technique that can be used for both classification and Regression problems, but mostly it is preferred for solving Classification problems. It is a tree-structured classifier, where internal nodes represent the features of a dataset, branches represent the decision rules and each leaf node represents the outcome.

In this there are two nodes, which are the Decision Node and Leaf Node. Decision nodes are used to make any decision and have multiple branches, whereas Leaf nodes are the output of those decisions and do not contain any further branches.

```
In [70]: model_2 = DecisionTreeClassifier()
model_2.fit(X_train, y_train)
Out[70]: DecisionTreeClassifier()
```

RANDOM FOREST CLASSIFIER

- Random Forest is a classifier that contains a number of decision trees on various subsets of the given dataset and takes the average to improve the predictive accuracy of that dataset.
- Instead of relying on one decision tree, the random forest takes the prediction from each tree and based on the majority votes of predictions, and it predicts the final output.
- The greater number of trees in the forest leads to higher accuracy and prevents the problem of overfitting.

```
In [71]: model_3 = RandomForestClassifier()
model_3.fit(X_train, y_train)
```

Out[71]: RandomForestClassifier()

MODEL PREDICTION AND EVALUATION

We precdicted Man of the match using all three models on Test dataset(X_test).

We evaluated our models by calculating ACCURACY, PRECISION, RECALL and F1 SCORE.

ACCURACY - The simplest way of reporting the effectiveness of an algorithm is by calculating its accuracy. Accuracy is the ratio of number of correct predictions to the total number of predictions made.

RECALL - Recall measures the percentage of relevant items that your classifier found.

PRECISION - Precision is the ratio of correctly predicted positive observations to the total predicted positive observations

F1 SCORE -F1 Score is the Harmonic Mean between precision and recall. The range for F1 Score is [0, 1]. It tells you how precise your classifier is , as well as how robust it is (it does not miss a significant number of instances).

Model evaluation scores

```
In [498]: model_evaluation_data = {
    'Model' : models,
    'Accuracy' : accuracy_scores,
    'Precision' : precision_scores,
    'Recall' : recall_scores,
    'F1 score' : f1_scores
}
model_evaluation_df = pd.DataFrame(model_evaluation_data)
model_evaluation_df
```

Out[498]:

	Model	Accuracy	Precision	Recall	F1 score
0	Logistic Regression	0.846154	0.769231	0.909091	0.833333
1	Decision Tree	0.730769	0.666667	0.727273	0.695652
2	Random Forest	0.730769	0.642857	0.818182	0.720000

