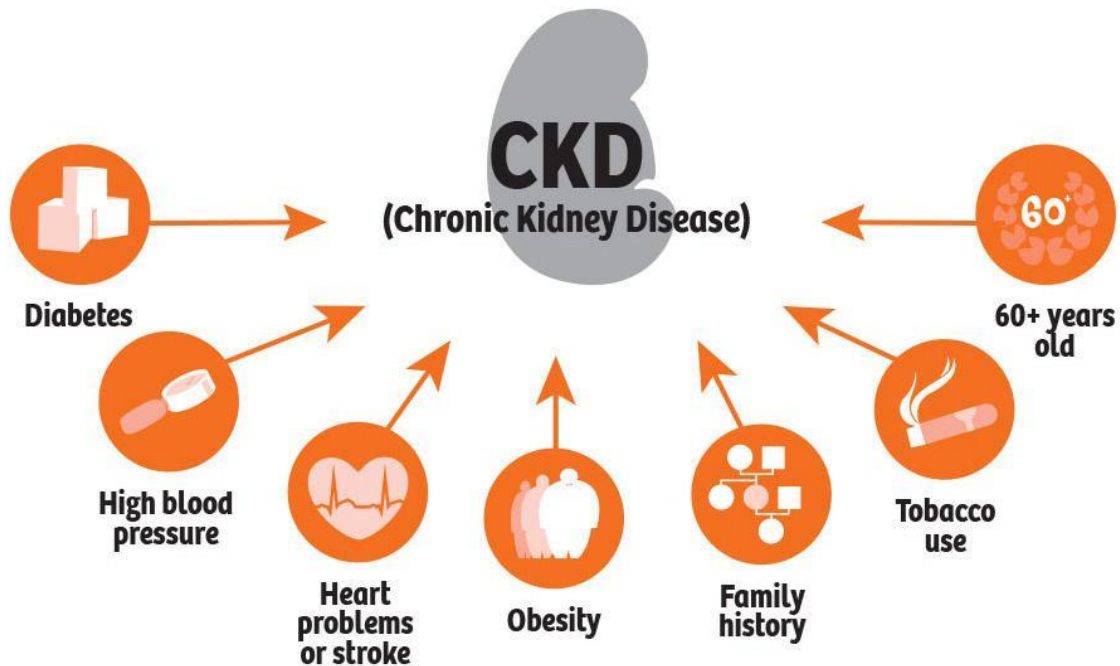


Chronic Kidney Decease



1. Problem statement: Prediction of chronic kidney decease from given dataset.
(Machine learning-supervised learning-classification)
2. Dataset details: Number of Columns-25 & Number of Rows-399.
3. Data preprocessing performed for column name – rbc, pc, pcc, ba.
4. Machine learning algorithms for classification using random forest produce good results.

```
[[45  0]
 [ 2 73]]
```

	precision	recall	f1-score	support
0	0.96	1.00	0.98	45
1	1.00	0.97	0.99	75
accuracy			0.98	120
macro avg	0.98	0.99	0.98	120
weighted avg	0.98	0.98	0.98	120

```
0.9834018801410106
0.9997037037037038
```

5. Other classification algorithm research values:

a. Decision Tree:

```
Fitting 3 folds for each of 12 candidates, totalling 36 fits
[[44  1]
 [ 3 72]]
      precision    recall  f1-score   support

     0       0.94       0.98       0.96         45
     1       0.99       0.96       0.97         75

 accuracy          0.97         120
 macro avg       0.96       0.97       0.96         120
 weighted avg    0.97       0.97       0.97         120

0.9668037602820211
0.9688888888888889
```

b. Random Forest:

```
[[45  0]
 [ 2 73]]
      precision    recall  f1-score   support

     0       0.96       1.00       0.98         45
     1       1.00       0.97       0.99         75

 accuracy          0.98         120
 macro avg       0.98       0.99       0.98         120
 weighted avg    0.98       0.98       0.98         120

0.9834018801410106
0.9997037037037038
```

c. Logistic Regression:

```
Fitting 3 folds for each of 4 candidates, totalling 12 fits
[[43  2]
 [ 0 75]]
      precision    recall  f1-score   support

     0       1.00       0.96       0.98         45
     1       0.97       1.00       0.99         75

 accuracy          0.98         120
 macro avg       0.99       0.98       0.98         120
 weighted avg    0.98       0.98       0.98         120

0.9832535885167464
0.9973333333333334
```

d. Naïve Bayes:

```
Fitting 3 folds for each of 1 candidates, totalling 3 fits
[[45  0]
 [ 2 73]]
      precision    recall  f1-score   support

     0       0.96       1.00       0.98         45
     1       1.00       0.97       0.99         75

 accuracy          0.98         120
 macro avg       0.98       0.99       0.98         120
weighted avg       0.98       0.98       0.98         120

0.9834018801410106
1.0
```

e. K Nearest neighbors:

```
Fitting 3 folds for each of 4 candidates, totalling 12 fits
[[41  4]
 [22 53]]
      precision    recall  f1-score   support

     0       0.65       0.91       0.76         45
     1       0.93       0.71       0.80         75

 accuracy          0.78         120
 macro avg       0.79       0.81       0.78         120
weighted avg       0.83       0.78       0.79         120

0.7866161616161614
0.8558518518518519
```

f. Support vector machine:

```
[[44  1]
 [ 1 74]]
      precision    recall  f1-score   support

     0       0.98       0.98       0.98         45
     1       0.99       0.99       0.99         75

 accuracy          0.98         120
 macro avg       0.98       0.98       0.98         120
weighted avg       0.98       0.98       0.98         120

0.9833333333333335
0.9985185185185185
```

6. Final Model analysis:

The trained model's development indicated that the Random Forest algorithm delivered the best performance across all parameters.

- It predicts a 98% total model performance.
- Type I error in the confusion matrix is 0, and Type II error is 2. Here, type I error has a lower value compared to type II error, which suggests that predicting chronic renal disease is accurate.
- A slight increase or decrease in recall and precision values. Only F1 score is taken into consideration for the best outcome.
- The ROC value for this problem is 0.9997, or almost 1, indicating that the random forest algorithm is effective.