# Classification Assignment

### Problem Statement or Requirement:

A requirement from the Hospital, Management asked us to create a predictive model which will predict the Chronic Kidney Disease (CKD) based on the several parameters. The Client has provided the dataset of the same.

- 1.) Identify your problem statement
- 2.) Tell basic info about the dataset (Total number of rows, columns)
- 3.) Mention the pre-processing method if you're doing any (like converting string to number nominal data)
- 4.) Develop a good model with good evaluation metric. You can use any machine learning algorithm; you can create many models. Finally, you have to come up with final model.
- 5.) All the research values of each algorithm should be documented. (You can make tabulation or screenshot of the results.)
- 6.) Mention your final model, justify why u have chosen the same.
- 1. The hospital wants a reliable predictive model to classify patients into CKD (Chronic Kidney Disease) or Not CKD, using medical attributes. You are tasked with building the best-performing classification model, evaluated using precision, recall, f1-score, and accuracy.

Domain: ML

Type: Supervised Learning

Objective: Binary Classification

```
In [84]: #importing the Libraies
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd

In [85]: # Reading the Dataset
dataset = pd.read_csv('CKD.csv')
```

2. Tell basic info about the dataset (Total number of rows, columns)

```
In [86]: print(f"\nRows: {dataset.shape[0]}")
print(f"\nColumns: {dataset.shape[1]}\n\n")

display(dataset)

# Displaying the dataset information
display(dataset.info())
```

Rows: 399 Columns: 25

	age	bp	sg	al	su	rbc	рс	рсс	ba	bgr	 pcv	wc	rc	htn	dm	cad	appet	pe	ane	classification
0	2.000000	76.459948	C	3.0	0.0	normal	abnormal	notpresent	notpresent	148.112676	 38.868902	8408.191126	4.705597	no	no	no	yes	yes	no	yes
1	3.000000	76.459948	С	2.0	0.0	normal	normal	notpresent	notpresent	148.112676	 34.000000	12300.000000	4.705597	no	no	no	yes	роог	no	yes
2	4.000000	76.459948	а	1.0	0.0	normal	normal	notpresent	notpresent	99.000000	 34.000000	8408.191126	4.705597	no	no	no	yes	роог	no	yes
3	5.000000	76.459948	d	1.0	0.0	normal	normal	notpresent	notpresent	148.112676	 38.868902	8408.191126	4.705597	no	no	no	yes	роог	yes	yes
4	5.000000	50.000000	С	0.0	0.0	normal	normal	notpresent	notpresent	148.112676	 36.000000	12400.000000	4.705597	no	no	no	yes	роог	no	yes
394	51.492308	70.000000	a	0.0	0.0	normal	normal	notpresent	notpresent	219.000000	 37.000000	9800.000000	4.400000	no	no	no	yes	роог	no	yes
395	51.492308	70.000000	C	0.0	2.0	normal	normal	notpresent	notpresent	220.000000	 27.000000	8408.191126	4.705597	yes	yes	no	yes	роог	yes	yes
396	51.492308	70.000000	C	3.0	0.0	normal	normal	notpresent	notpresent	110.000000	 26.000000	9200.000000	3.400000	yes	yes	no	роог	роог	no	yes
397	51.492308	90.000000	а	0.0	0.0	normal	normal	notpresent	notpresent	207.000000	 38.868902	8408.191126	4.705597	yes	yes	no	yes	роог	yes	yes
398	51.492308	80.000000	а	0.0	0.0	normal	normal	notpresent	notpresent	100.000000	 53.000000	8500.000000	4.900000	no	no	no	yes	роог	no	no

399 rows × 25 columns

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 399 entries, 0 to 398
Data columns (total 25 columns):
# Column
                   Non-Null Count Dtype
    -----
                    -----
0
    age
                    399 non-null
                                   float64
                    399 non-null
1
    bp
                                   float64
                    399 non-null
                                   object
2
    sg
                    399 non-null
3
    al
                                   float64
 4
    su
                    399 non-null
                                   float64
    rbc
                    399 non-null
                                   object
 6
    рс
                    399 non-null
                                   object
7
                    399 non-null
    рсс
                                   object
                    399 non-null
8
    ba
                                   object
 9
    bgr
                    399 non-null
                                   float64
 10
    bu
                    399 non-null
                                   float64
 11 sc
                    399 non-null
                                   float64
 12 sod
                    399 non-null
                                   float64
                    399 non-null
 13 pot
                                   float64
 14 hrmo
                    399 non-null
                                   float64
                    399 non-null
 15 pcv
                                   float64
                    399 non-null
 16 wc
                                   float64
 17 rc
                    399 non-null
                                   float64
 18 htn
                    399 non-null
                                   object
 19 dm
                    399 non-null
                                   object
 20 cad
                    399 non-null
                                   object
 21 appet
                    399 non-null
                                   object
 22 pe
                    399 non-null
                                   object
 23 ane
                    399 non-null
                                   object
24 classification 399 non-null
                                   object
dtypes: float64(13), object(12)
memory usage: 78.1+ KB
None
```

3.) Mention the pre-processing method if you're doing any (like converting string to number – nominal data)

```
In [87]: # Converting categorical variables to numerical
# as the classification column is categorical - ordinal data, will be converted to numerical values using LabelEncoder
from sklearn.preprocessing import LabelEncoder

# Handle categorical and numerical separately
categorical_cols = dataset.select_dtypes(include='object').columns
numerical_cols = dataset.select_dtypes(exclude='object').columns

# Fill missing values
dataset[categorical_cols] = dataset[categorical_cols].fillna(dataset[categorical_cols].mode().iloc[0])
```

```
dataset[numerical_cols] = dataset[numerical_cols].fillna(dataset[numerical_cols].mean())

# Encode categorical columns
label_encoders = {}
for col in categorical_cols:
    le = LabelEncoder()
    dataset[col] = le.fit_transform(dataset[col])
    label_encoders[col] = le
```

4. Develop a good model with good evaluation metric. You can use any machine learning algorithm; you can create many models. Finally, you have to come up with final model.

```
In [88]: # Splitting the dataset into independent and dependent variables
    independent = dataset.drop('classification', axis=1)
    dependent = dataset['classification']

In [89]: #split into training set and test
    from sklearn.model_selection import train_test_split
    X_train, X_test, y_train, y_test = train_test_split(independent, dependent, test_size = 1/3, random_state = 0)

In [90]: from sklearn.preprocessing import StandardScaler
    StandardScaler = StandardScaler()
    X_train = StandardScaler.fit_transform(X_train)
    X_test = StandardScaler.transform(X_test)
```

5. All the research values of each algorithm should be documented. (You can make tabulation or screenshot of the results.)

### The report using LogisticRegression:

	precision	recall	f1-score	support
0	0.96	1.00	0.98	51
1	1.00	0.98	0.99	82
accuracy			0.98	133

macro avg 0.98 0.99 0.98 133 weighted avg 0.99 0.98 0.99 133

### The report using K-Nearest Neibour KNN:

	precision	recall	fl-score	support
0	0.93	1.00	0.96	51
1	1.00	0.95	0.97	82
accuracy			0.97	133

macro avg 0.96 0.98 0.97 133 weighted avg 0.97 0.97 0.97 133

### The report using GaussianNB:

	precision	recall	f1-score	support
0 1	0.91 1.00	1.00 0.94	0.95 0.97	51 82
ıracy			0.96	133

macro avg 0.96 0.97 0.96 133 weighted avg 0.97 0.96 0.96 133

The report using MultinomialNB:

accu

	precision	recall	f1-score	suppor
0	0.39 1.00	1.00	0.56 0.07	51 82
1		0.04		
accuracy			0.41	133

macro avg 0.70 0.52 0.32 133 weighted avg 0.77 0.41 0.26 133

### The report using BernoulliNB:

	precision	recall	f1-score	support
0 1	0.96 1.00	1.00 0.98	0.98 0.99	51 82
accuracy			0.98	133

macro avg 0.98 0.99 0.98 133 weighted avg 0.99 0.98 0.99 133

# The report using ComplementNB:

	precision	recall	f1-score	support
0	0.40	1.00	0.57	51
1	1.00	0.05	0.09	82
accuracy			0.41	133

macro avg 0.70 0.52 0.33 133 weighted avg 0.77 0.41 0.27 133

### The report using CategoricalNB:

	precision	recall	f1-score	support
0.0 1.0	0.98 1.00	1.00 0.99	0.99 0.99	51 82
accuracy			0.99	133

macro avg 0.99 0.99 0.99 133 weighted avg 0.99 0.99 0.99 133

# The report using SVC:

	precision	recall	f1-score	support
0 1	0.94 0.98	0.96 0.96	0.95 0.97	51 82
accuracy			0.96	133

macro avg 0.96 0.96 0.96 133 weighted avg 0.96 0.96 0.96 133

# The report using RandomForestClassifier :

	precision	recall	f1-score	support
0	1.00	1.00	1.00	51
1	1.00	1.00	1.00	82

accuracy	1.00	133

macro avg 1.00 1.00 1.00 133 weighted avg 1.00 1.00 1.00 133

### The report using DecisionTreeClassifier:

	precision	recall	f1-score	support
0	0.86	0.98	0.92	51
1	0.99	0.90	0.94	82
accuracy			0.93	133

macro avg 0.92 0.94 0.93 133 weighted avg 0.94 0.93 0.93 133

6. Mention your final model, justify why u have chosen the same.

After evaluating multiple classification algorithms, RandomForestClassifier was chosen as the final model due to its perfect classification performance with 100% accuracy, precision, recall, and F1-score on the test dataset. In addition to superior performance, its ability to handle both numerical and categorical data, resistance to overfitting, and interpretability through feature importance make it the most appropriate choice for chronic kidney disease prediction.

### Justification:

- 1. Superior Performance RandomForestClassifier delivers perfect classification on the test data. While other models like CategoricalNB, BernoulliNB, and LogisticRegression come very close with ~98–99% accuracy, RandomForest achieves a clean 1.00 across all evaluation metrics.
- 2. Handles Both Numerical and Categorical Data Random Forests are versatile—they do not require strict feature scaling or encoding schemes, unlike SVM or Naive Bayes models. This is particularly useful in medical datasets like CKD, which often contain mixed data types.
- 3. Robust to Noise & Outliers Ensemble methods like RandomForest are less likely to overfit, especially compared to Decision Trees or models that rely on distribution assumptions (e.g., GaussianNB).
- 4. Feature Importance Random Forests allow you to interpret feature importance, which is useful in a clinical setting to understand which parameters (e.g., serum creatinine, blood pressure) contribute most to CKD risk.

#### Final Model: RandomForestClassifier

### Performance:

Accuracy: 100%

Precision, Recall, F1-Score: All metrics are 1.00 for both classes.

Support: Balanced performance on both CKD (label 1) and non-CKD (label 0) classes.