



# GOVERNMENT OF TAMILNADU

## Naan Muthalvan - Project-Based Experiential Learning

Early  
Prediction For Chronic Kidney Disease Detection: A Progressive Approach  
to Health Management

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# 1.INTRODUCTION

## 1.1 Overview:

Chronic Kidney Disease (CKD) is a major medical problem and can be cured if treated in the early stages. Usually, people are not aware that medical tests we take for different purposes could contain valuable information concerning kidney diseases. Consequently, attributes of various medical tests are investigated to distinguish which attributes may contain helpful information about the disease. The information says that it helps us to measure the severity of the problem, the predicted survival of the patient after the illness, the pattern of the disease and work for curing the disease. In today's world as we know most of the people are facing so many diseases and as this can be cured if we treat people in early stages this project can use a pretrained model to predict the Chronic Kidney Disease which can help in treatments of people who suffer from this disease.

## 1.2 Purpose

Chronic kidney disease (CKD) occurs when the kidneys gradually lose their function over time. There are several potential causes of CKD, including

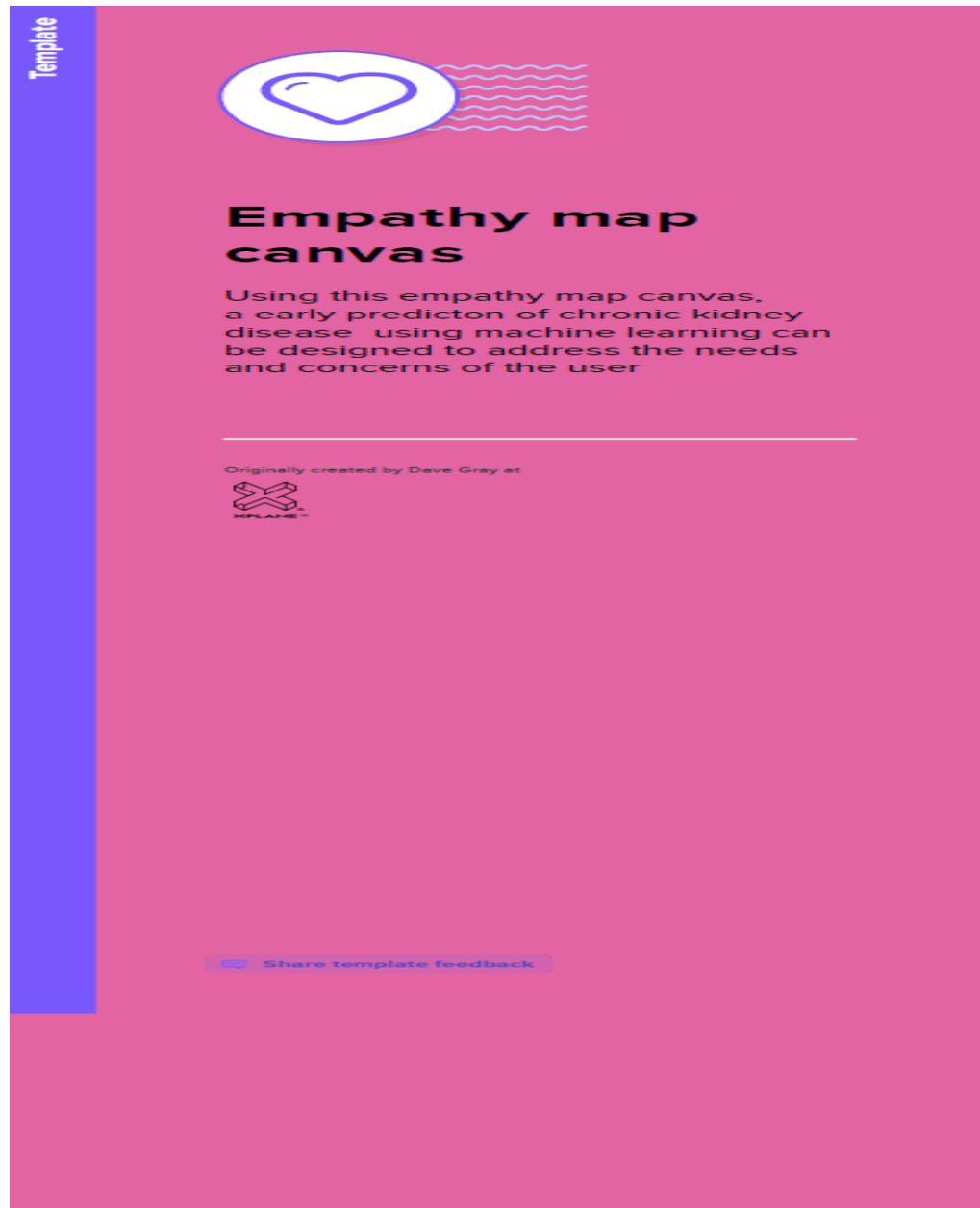
1. Diabetes: High blood sugar levels can damage the small blood vessels in the kidneys, making them less effective at filtering waste and excess fluid from the body.
2. High blood pressure: High blood pressure can damage the small blood vessels in the kidneys, as well as the delicate filtering units called nephrons.
3. Glomerulonephritis: This is an inflammation of the tiny filters in the kidneys, which can cause scarring and permanent damage.
4. Polycystic kidney disease: This is a genetic condition where cysts develop in the kidneys, impairing their function.

Other conditions: Certain autoimmune diseases, infections, and genetic disorders can also lead to D

The purpose of the kidneys is to filter waste and excess fluid from the blood, maintaining a healthy balance of electrolytes and other substances in the body.

## 2. Problemdefinition&DesignThinking

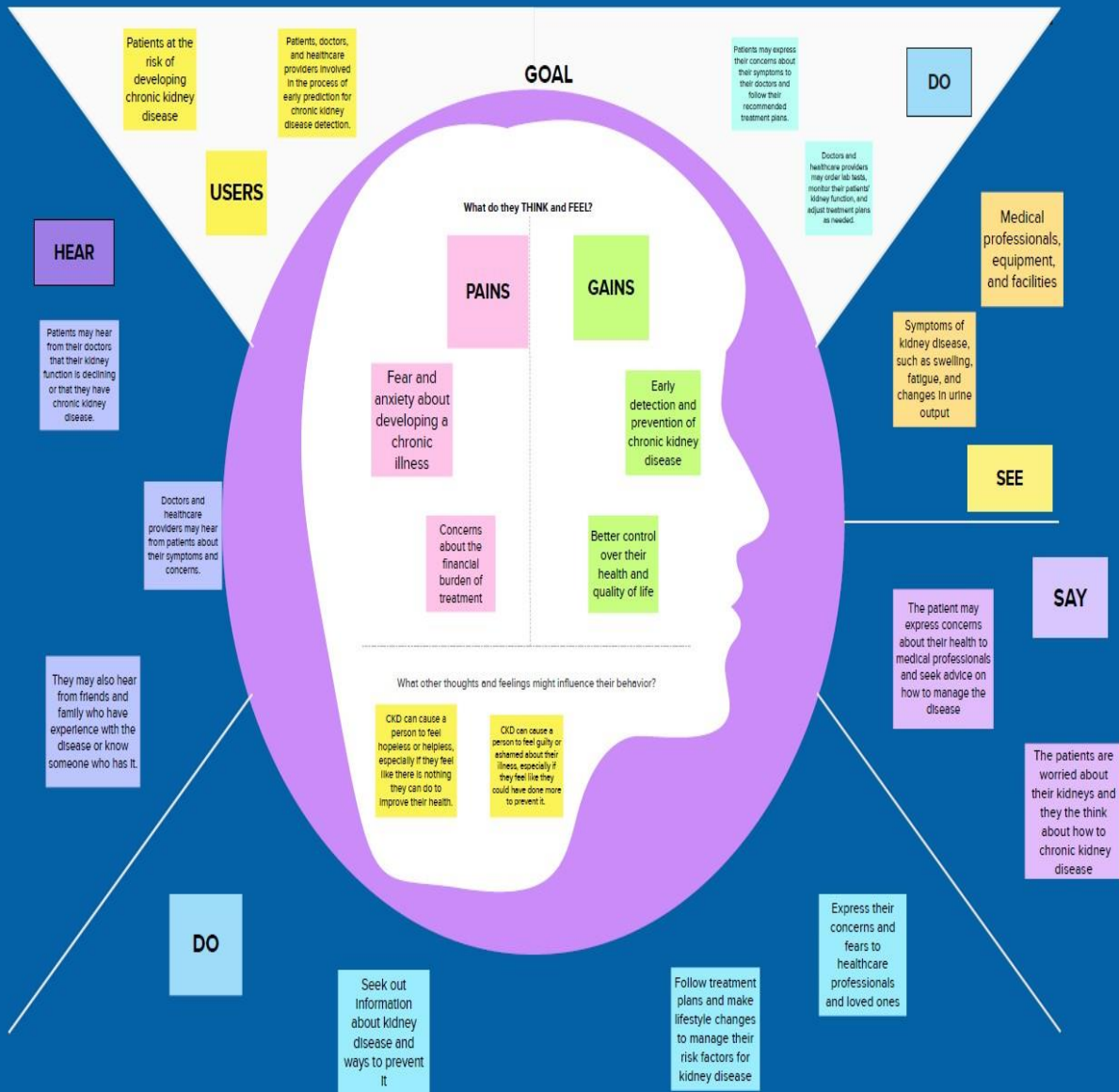
### 2.1 EmpathyMap





## Develop shared understanding and empathy

By using the empathy map, we are learn to develop ideas.



## 2.2 Ideation&BrainstormingMap

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### Project Ideas

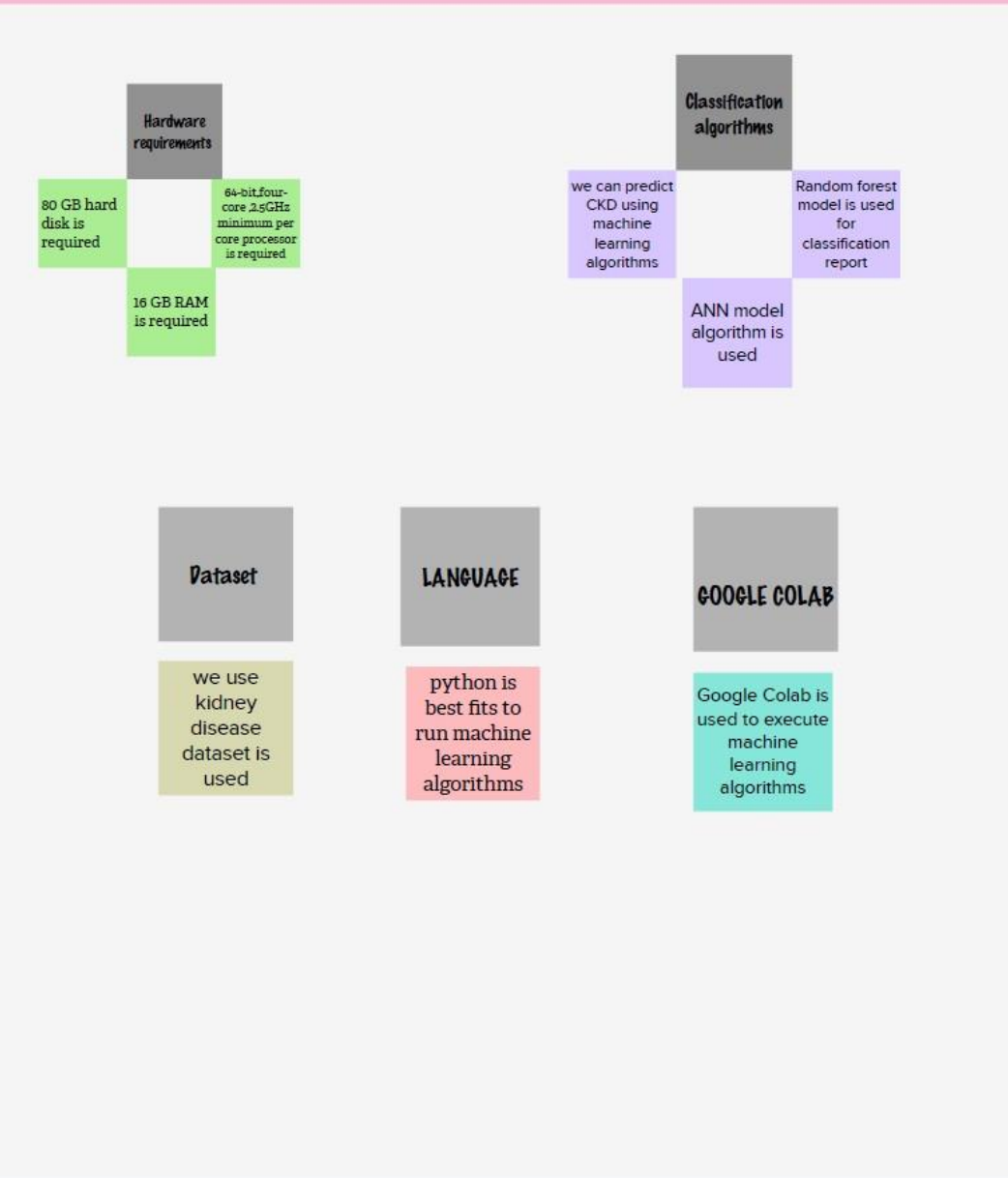
Quickly process  
large amount of  
university data  
Save cost  
associated with manual  
process



3

### Group ideas

1. 64-bit processor, 16 GB RAM is required
2. Machine learning algorithm such as ANN model, Logistic Regression, Random forest model is used.
3. Kidney disease dataset is used and it is imported from Kaggle.
4. Python language is used to execute Machine learning algorithms.
5. Google Colab is used for executing Machine learning algorithms.

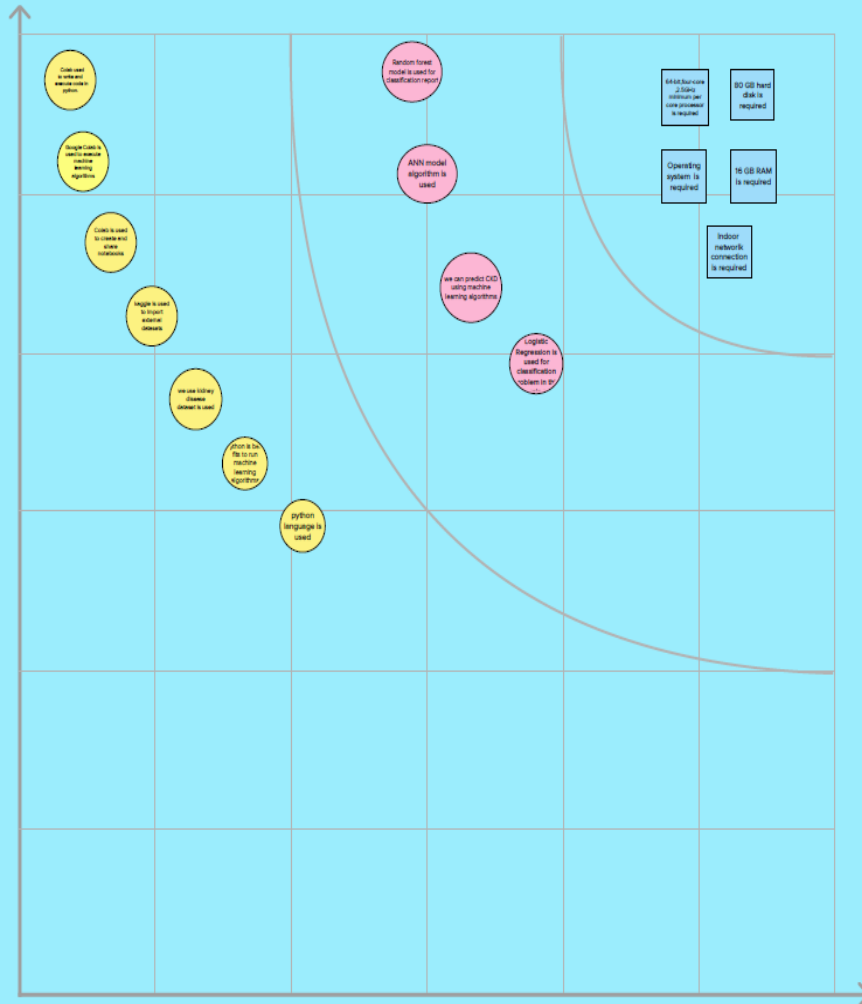


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## Prioritize

we prioritize the ideas based on the project requirements

🕒 20 minutes

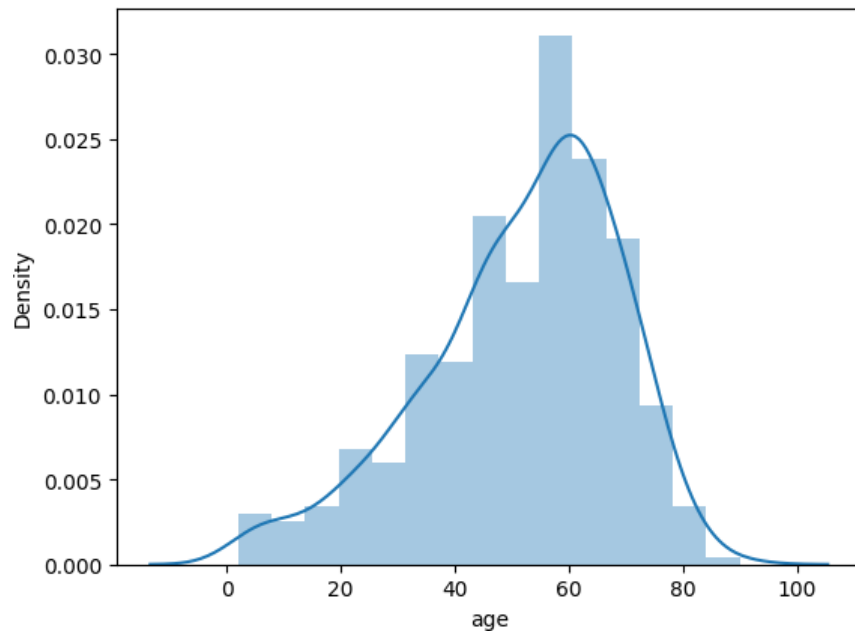




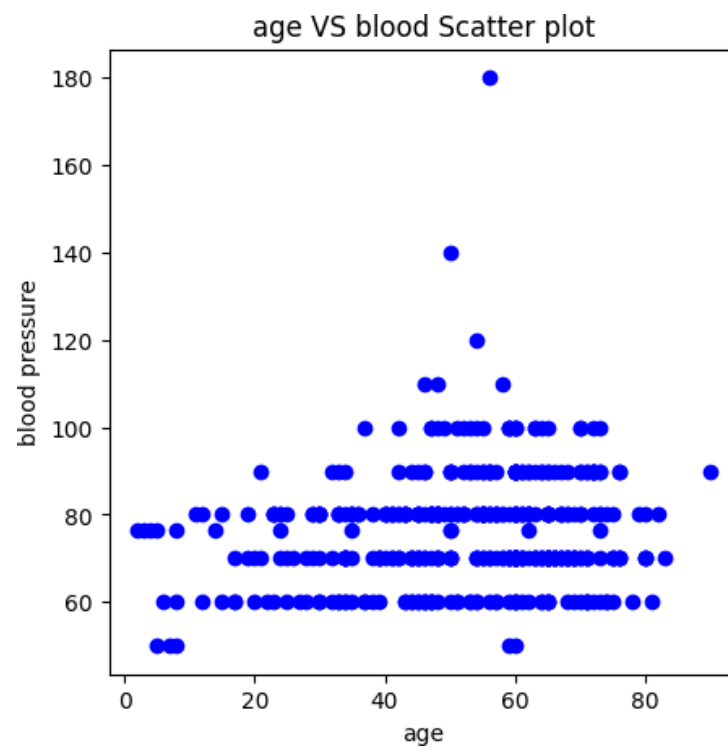
3.

## Result

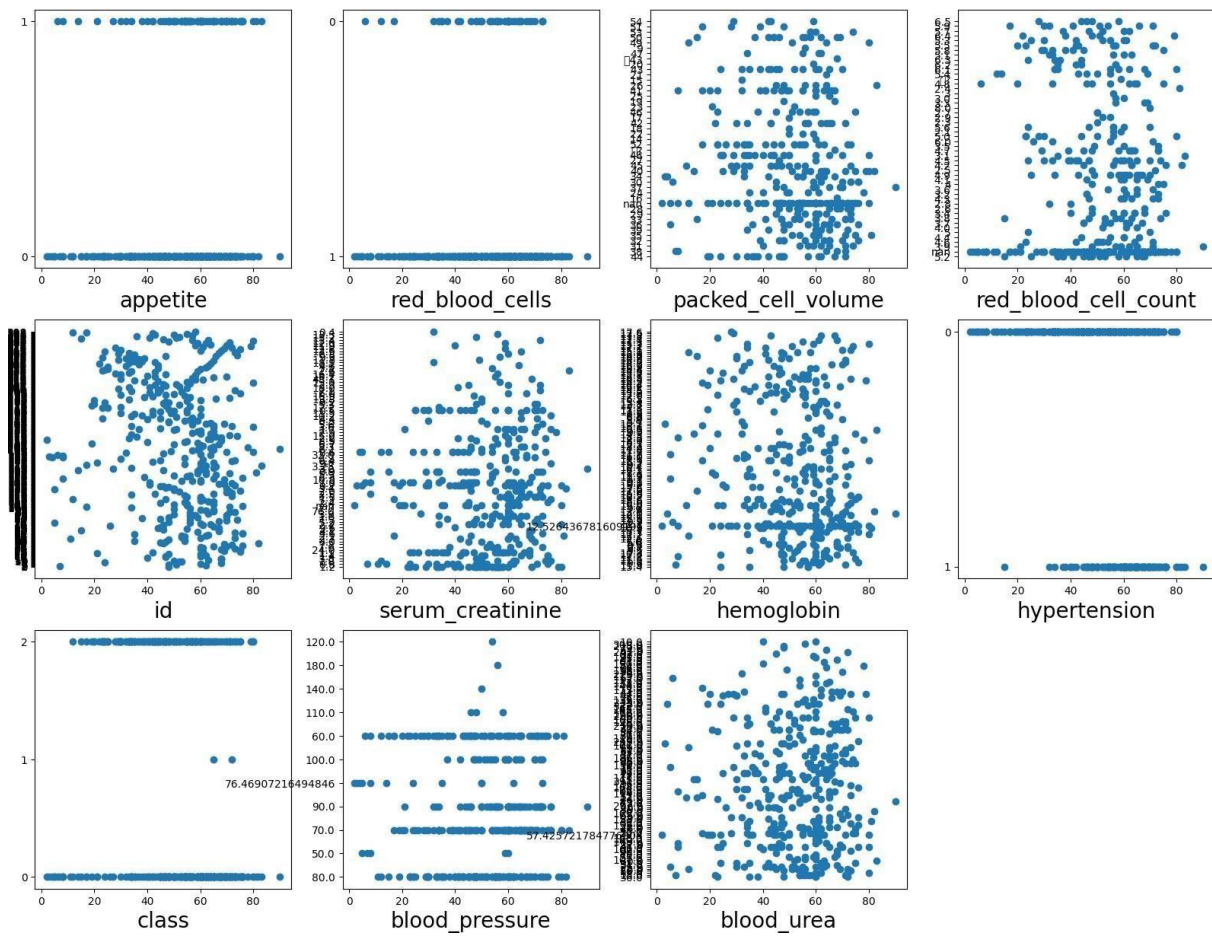
### 3.1 Univariateanalysis

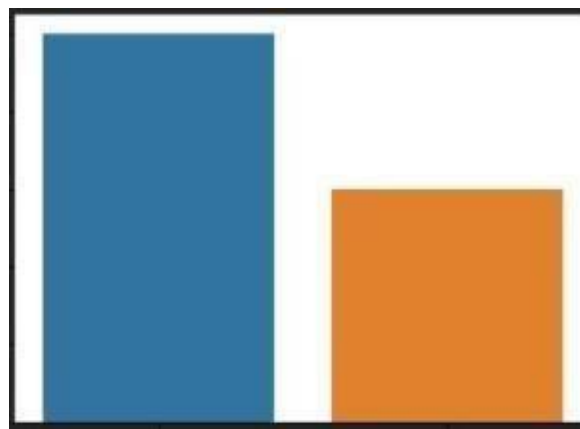
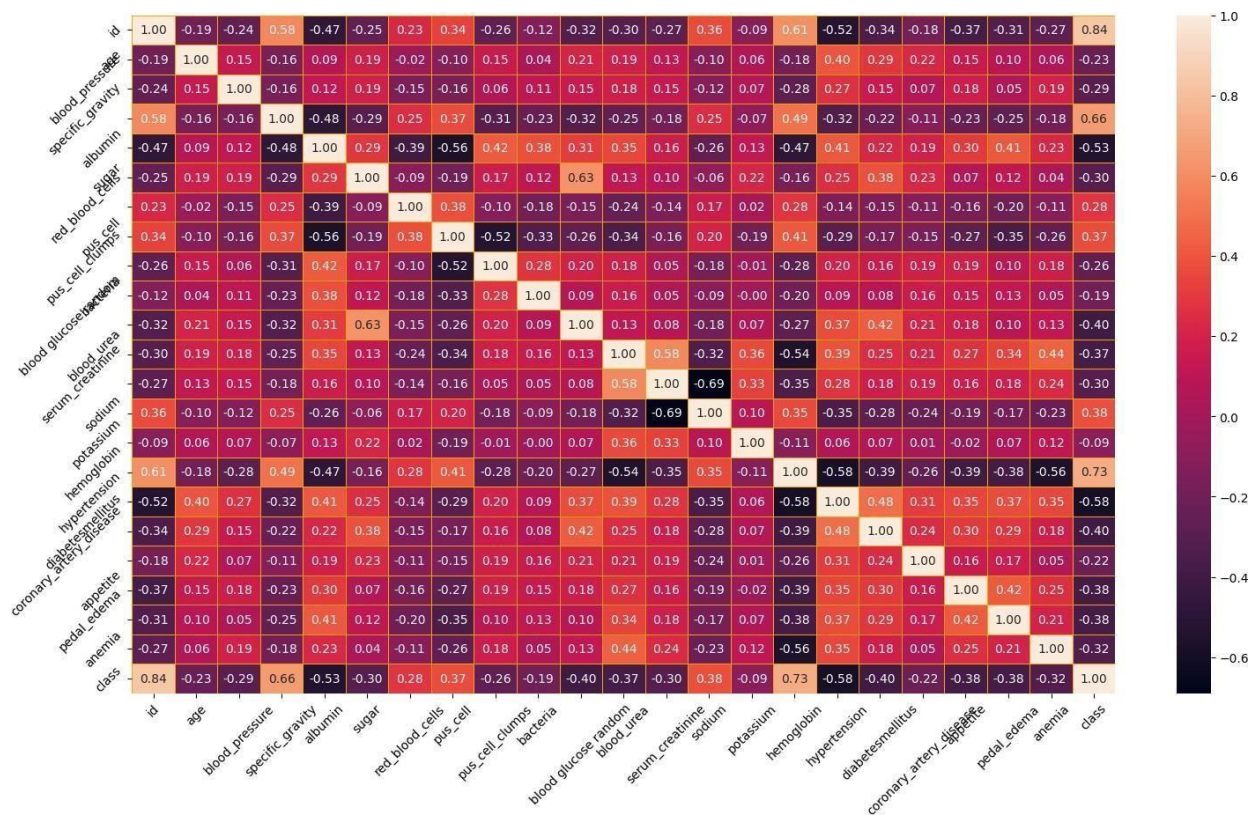


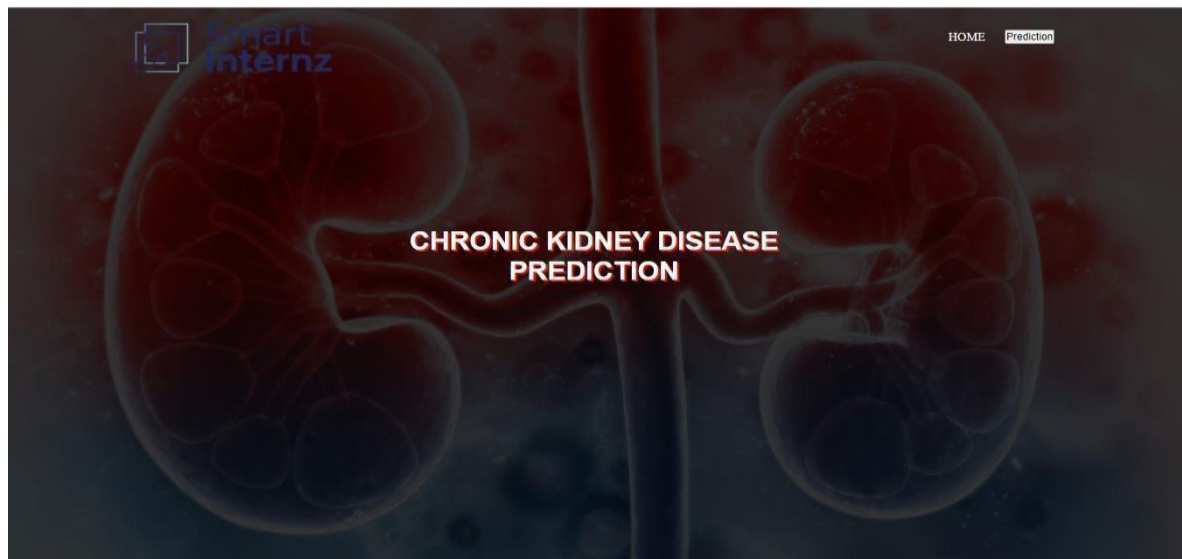
### 3.2 Bivariateanalysis



### 3.3 Multivariate analysis







*Chronic Kidney Disease*  
A Machine Learning Web App, Built with Flask

Enter your blood_urea
Enter your blood glucose random
Select anemia or not
Select coronary artery disease or not
Select pus_cell or not
Select red_blood_cell level
Select diabetesmellitus or not
Select pedal_edema or not

Predict



Input - Now, the user will give inputs to get the predicted result after clicking onto the submit button.

Chronic Kidney Disease

A Machine Learning Web App, Built with Flask

1

1

NO

NO

normal

normal

NO

NO

Predict

Chronic Kidney Disease

A Machine Learning Web App, Built with Flask

Prediction: Great! You DON'T have Chronic Kidney Disease

A meme featuring three Minions from the Despicable Me franchise. They are standing in a row, looking happy and excited. The text "IT'S TIME TO PARTYYYYYY!" is written in large, bold, white capital letters across the top of the image.

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### **3. ADVANTAGES & DISADVANTAGES**

#### **ADVANTAGES**

- Early detection and treatment of CKD can slow down or stop the progression of the disease, which can help prevent or delay the need for dialysis or a kidney transplant.
- CKD may be detected during routine blood or urine tests, which can lead to earlier detection of other health problems that may be contributing to the kidney disease.
- CKD may lead to changes in lifestyle that can improve overall health, such as following a healthy diet, quitting smoking, and increasing physical activity.

#### **DISADVANTAGES**

- CKD can increase the risk of serious health problems, such as heart disease, stroke, and bone disease.
- The symptoms of CKD may be vague and nonspecific, making it difficult to detect until the disease has progressed to a later stage.
- Treatment for CKD can be expensive and time-consuming, requiring regular doctor visits, medications, and potentially dialysis or a kidney transplant.
- CKD can have a significant impact on quality of life, including fatigue, sleep disturbances, and changes in appetite or body weight.
- CKD can also lead to feelings of anxiety, depression, or stress, especially if it requires major lifestyle changes or interferes with daily activities.

## 4. APPLICATION

Chronic kidney disease (CKD) has several applications in healthcare, including:

1. **Diagnosis:** CKD can be diagnosed using blood and urine tests that measure kidney function and detect any abnormalities. These tests can help healthcare providers identify the disease early, allowing for early intervention and better outcomes.
2. **Monitoring:** Once diagnosed, CKD can be monitored over time to track changes in kidney function and determine the effectiveness of treatment. Monitoring may include regular blood and urine tests, imaging studies, and other diagnostic tests.
3. **Treatment:** CKD can be treated using a variety of approaches, including medications, lifestyle changes, and dialysis or kidney transplant in advanced cases. Healthcare providers may use a combination of these treatments to manage the disease and prevent complications.
4. **Prevention:** Individuals at risk of CKD, such as those with diabetes, high blood pressure, or a family history of kidney disease, may benefit from early detection and intervention to prevent the disease from developing.
5. **Research:** CKD research has led to advancements in understanding the disease, identifying risk factors, and developing new treatments. Ongoing research in this area may lead to further improvements in CKD diagnosis, treatment, and prevention.

## 6.

## CONCLUSION

Individuals at risk of CKD should be vigilant about monitoring their kidney function and taking steps to prevent the disease from developing or progressing. Healthcare providers play a critical role

in diagnosing, monitoring, and treating CKD, and ongoing research in this area may lead to further improvements in CKD outcomes. Overall, managing CKD requires a comprehensive, multidisciplinary approach that involves healthcare providers, patients, and their families.

In conclusion, chronic kidney disease (CKD) is a serious medical condition in which the kidneys gradually lose their ability to function properly over time. While early detection and intervention can slow or stop the progression of the disease, CKD can increase the risk of serious health problems, such as heart disease and bone disease. Treatment for CKD can be expensive and time-consuming, and may include medications, lifestyle changes, and in some cases, dialysis or a kidney transplant.



## 7.

## FUTURESCOPE

There are several areas of future research and development in chronic kidney disease (CKD) that have the potential to improve diagnosis, treatment, and outcomes. Here are some examples:

1. **Early Detection:** Researchers are exploring new biomarkers that can detect CKD at an earlier stage, before significant kidney damage has occurred. These biomarkers may include urine and blood tests, imaging studies, and other diagnostic tools.
2. **Precision Medicine:** Advances in genetics and personalized medicine may lead to more targeted treatments for CKD based on an individual's unique genetic makeup and other factors.
3. **Regenerative Medicine:** Researchers are exploring new techniques for repairing or regenerating damaged kidney tissue, such as stem cell therapy and tissue engineering.
4. **Artificial Kidneys:** Development of artificial kidneys that can replace the function of damaged kidneys, potentially reducing the need for dialysis or transplantation.
5. **Telemedicine:** Telemedicine and remote monitoring technologies may improve access to care for patients with CKD, especially in underserved areas or regions with limited access to specialists.
6. **Patient Education:** Improved patient education and engagement can help individuals with CKD better understand their condition, manage their symptoms, and adhere to treatment plans.

## 8.APPENDIX

### A.SOURCECODE

```
import pandas as pd
import numpy as np
from collections import Counter
import matplotlib.pyplot as plt
import seaborn as sns
import missingno as msno
from sklearn.metrics import accuracy_score, confusion_matrix
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import LabelEncoder
from sklearn.linear_model import LogisticRegression
import pickle

data = pd.read_csv("/content/kidney_disease.csv")
data.head()

data.columns
data.columns = ['id', 'age', 'blood_pressure', 'specific_gravity',
                 'albumin', 'sugar', 'red_blood_cells', 'pus_cell', 'pus_cell_clumps',
                 'bacteria', 'blood_glucose_random', 'blood_urea', 'serum_creatinine',
                 'sodium', 'potassium', 'hemoglobin', 'packed_cell_volume',
                 'white_blood_cell_count', 'red_blood_cell_count', 'hypertension',
                 'diabetes_mellitus', 'coronary_artery_disease', 'appetite',
                 'pedal_edema', 'anemia', 'class']

data.columns

data.info()
data.isnull().any()

data['blood_glucose_random'].fillna(data['blood_glucose_random'].mean(), inplace=True)
data['blood_pressure'].fillna(data['blood_pressure'].mean(), inplace=True)
data['blood_urea'].fillna(data['blood_urea'].mean(), inplace=True)
data['hemoglobin'].fillna(data['hemoglobin'].mean(), inplace=True)
data['packed_cell_volume'].fillna(data['packed_cell_volume'].mean(), inplace=True)
data['potassium'].fillna(data['potassium'].mean(), inplace=True)
data['red_blood_cell_count'].fillna(data['red_blood_cell_count'].mean(), inplace=True)
```

```

data['serum_creatinine'].fillna(data['blood_urea'].mean(), inplace=True)
data['sodium'].fillna(data['sodium'].mean(), inplace=True) data['white_blood_cell_count'].fillna(data['white_blood_cell_count'].mean(), inplace=True)

data['age'].fillna(data['age'].mode()[0], inplace=True) data['hypertension'].fillna(data['hypertension'].mode()[0], inplace=True)
)
data['pus_cell_clumps'].fillna(data['pus_cell_clumps'].mode()[0], inplace=True) data['appetite'].fillna(data['appetite'].mode()[0], inplace=True)
) data['albumin'].fillna(data['albumin'].mode()[0], inplace=True) data['pus_cell'].fillna(data['pus_cell'].mode()[0], inplace=True) data['red_blood_cells'].fillna(data['red_blood_cells'].mode()[0], inplace=True)
data['coronary_artery_disease'].fillna(data['coronary_artery_disease'].mode()[0], inplace=True) data['bacteria'].fillna(data['bacteria'].mode()[0], inplace=True) data['anemia'].fillna(data['anemia'].mode()[0], inplace=True) data['sugar'].fillna(data['sugar'].mode()[0], inplace=True) data['diabetesmellitus'].fillna(data['diabetesmellitus'].mode()[0], inplace=True) data['pedal_edema'].fillna(data['pedal_edema'].mode()[0], inplace=True) data['specific_gravity'].fillna(data['specific_gravity'].mode()[0], inplace=True)

data.isnull().sum()

catcols=set(data.dtypes[data.dtypes=='O'].index.values)
print(catcols)

for i in catcols:
    print("Columns:", i)
    print(c(data[i]))
    print('*'*120+'\n')

catcols.remove('red_blood_cell_count')
catcols.remove('packed_cell_volume')
catcols.remove('white_blood_cell_count')
print(catcols)

catcols=['anemia', 'pedal_edema', 'appetite', 'bacteria', 'class', 'coronary_artery_disease', 'diabetesmellitus', 'hypertension', 'pus_cell', 'pus_cell_clumps', 'red_blood_cells']

from sklearn.preprocessing import LabelEncoder
for i in catcols:

```

```

print("LELENCODINGOF:", i)
LEi=LabelEncoder()print(c(data
[i]))
data[i]=LEi.fit_transform(data[i])p
rint(c(data[i]))
print(""*100)

contcols=set(data.dtypes[data.dtypes!='O'].index.values)
print(contcols)

foriincontcols:
    print("ContinousColumns:", i) pri
    nt(c(data[i]))
    print(''*120+'\n')

contcols.remove('specific_gravity')
contcols.remove('albumin')contcols
.remove('sugar')print(contcols)

contcols.add('red_blood_cell_count')c
ontcols.add('packed_cell_volume')cont
cols.add('white_blood_cell_count')prin
t(contcols)

contcols.add('specific_gravity')
contcols.add('albumin')contcols
.add('sugar')print(contcols)

data['coronary_artery_disease']=data.coronary_artery_disease.replace('\
tno', 'no')
c(data['coronary_artery_disease'])

data['diabetesmellitus']=data.diabetesmellitus.replace(to_replace={'\tno': 'no'
: '\tyes': 'yes': 'yes': ''}
c(data['diabetesmellitus'])

data.describe()sns.di
stplot(data.age)
import matplotlib.pyplot as
pltfig=plt.figure(figsize=(5,5))plt.scatter(data['age'], dat
a['blood_pressure'], color='blue')plt.xlabel('age')
plt.ylabel('blood
pressure')plt.title("agevsbloodscatte
rplot")
plt.figure(figsize=(20,15),
facecolor='white')plotnumber=1

```

```

for column in contcols:
    if
        plotnumber <= 11:
            ax =
                plt.subplot(3, 4, plotnumber)
                plt.scatter(data['age'], data[column])
                plt.xlabel(column, fontsize=20)

            plotnumber += 1
plt.show()
plt.figure(figsize=(20, 15),
facecolor='white')
plotnumber = 1
for column in contcols:
    if
        plotnumber <= 11:
            ax =
                plt.subplot(3, 4, plotnumber)
                plt.scatter(data['age'], data[column])
                plt.xlabel(column, fontsize=20)

            plotnumber += 1
plt.show()
f, ax = plt.subplots(figsize=(18, 10))
sns.heatmap(data.corr(), annot=True,
fmt=".2f", ax=ax, linewidths=0.5, linecolor="orange")
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()
sns.countplot(data['class'])
])
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
x_bal = sc.fit_transform(x)
selcols = ['red_blood_cells', 'pus_cell', 'bloodglucose', 'blood_urea', 'pedal_edema', 'anemia', 'diabetesmellitus', 'coronary_artery_disease']
x = pd.DataFrame(data, columns=selcols)
y = pd.DataFrame(data, columns=['class'])
print(x.shape)
print(y.shape)
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, random_state=2)

```