

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

1 INTRODUCTION:

❖ OVERVIEW:

- ***Chronic kidney disease (CKD) is a long-term condition where the kidneys gradually lose their function over time. It is a significant public health issue, affecting millions of people worldwide, and if left untreated, can progress to end-stage renal disease (ESRD), requiring dialysis or kidney transplantation.***
- ***Early detection and management of CKD can significantly improve outcomes and prevent or delay the progression of the disease. Predicting the development of CKD in individuals at high risk can also help identify those who would benefit most from early intervention.***
- ***Several risk factors are associated with the development of CKD, such as hypertension, diabetes, obesity, family history, and age. Predictive models that integrate these risk factors can identify individuals at high risk of developing CKD and guide targeted screening and prevention strategies.***
- ***In this context, machine learning algorithms have shown promise in developing accurate and reliable predictive models for early CKD detection. By leveraging large datasets and advanced analytical techniques, these models can identify patterns and relationships between risk factors and disease outcomes, leading to more personalized and effective healthcare interventions.***

❖ **PURPOSE:**

- ***In the context of machine learning projects, the purpose of early kidney disease prediction is to develop accurate predictive models that can identify individuals at high risk of developing CKD. The use of machine learning algorithms can help identify patterns and relationships between risk factors and disease outcomes, leading to more personalized and effective healthcare interventions.***
- ***The development of predictive models for early CKD detection using machine learning can also lead to improved healthcare resource allocation, as targeted screening and prevention strategies can be employed for high-risk individuals. This can lead to more efficient use of healthcare resources and improved patient outcomes.***
- ***In addition, machine learning algorithms can help identify modifiable risk factors that contribute to the development of CKD, such as hypertension and diabetes. By addressing these risk factors early, healthcare providers can help prevent the onset of CKD in high-risk individuals, leading to better health outcomes and improved quality of life.***
- ***Overall, the development of machine learning models for early kidney disease prediction has the potential to make a significant impact on public health by reducing the burden of this chronic condition.***

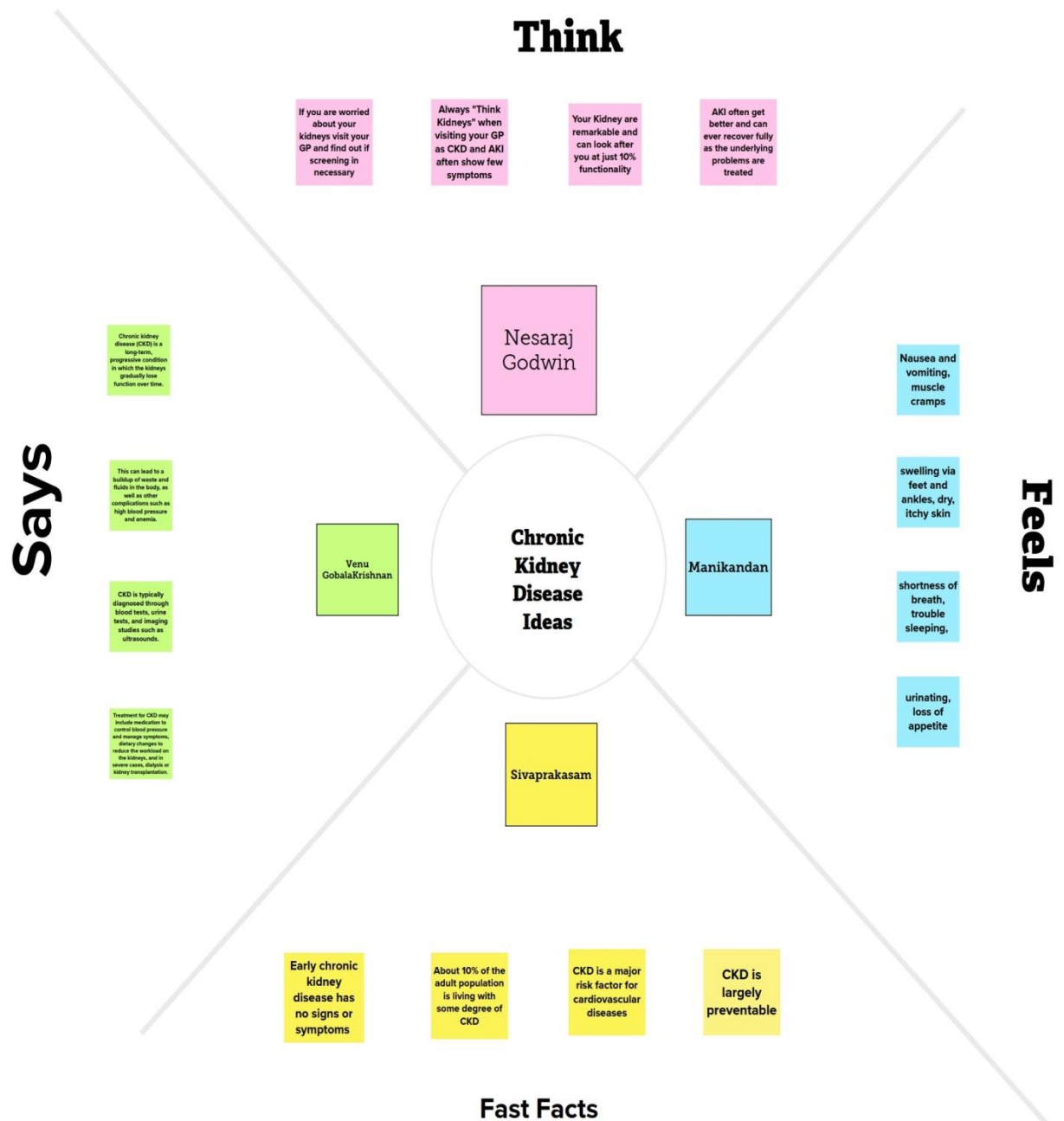
2 Problem Definition & Design Thinking:

❖ EMPATHY MAP



Build empathy of Chronic Kidney Disease Ideas

The information you add here should be representative of the observations and research you've done about your users.

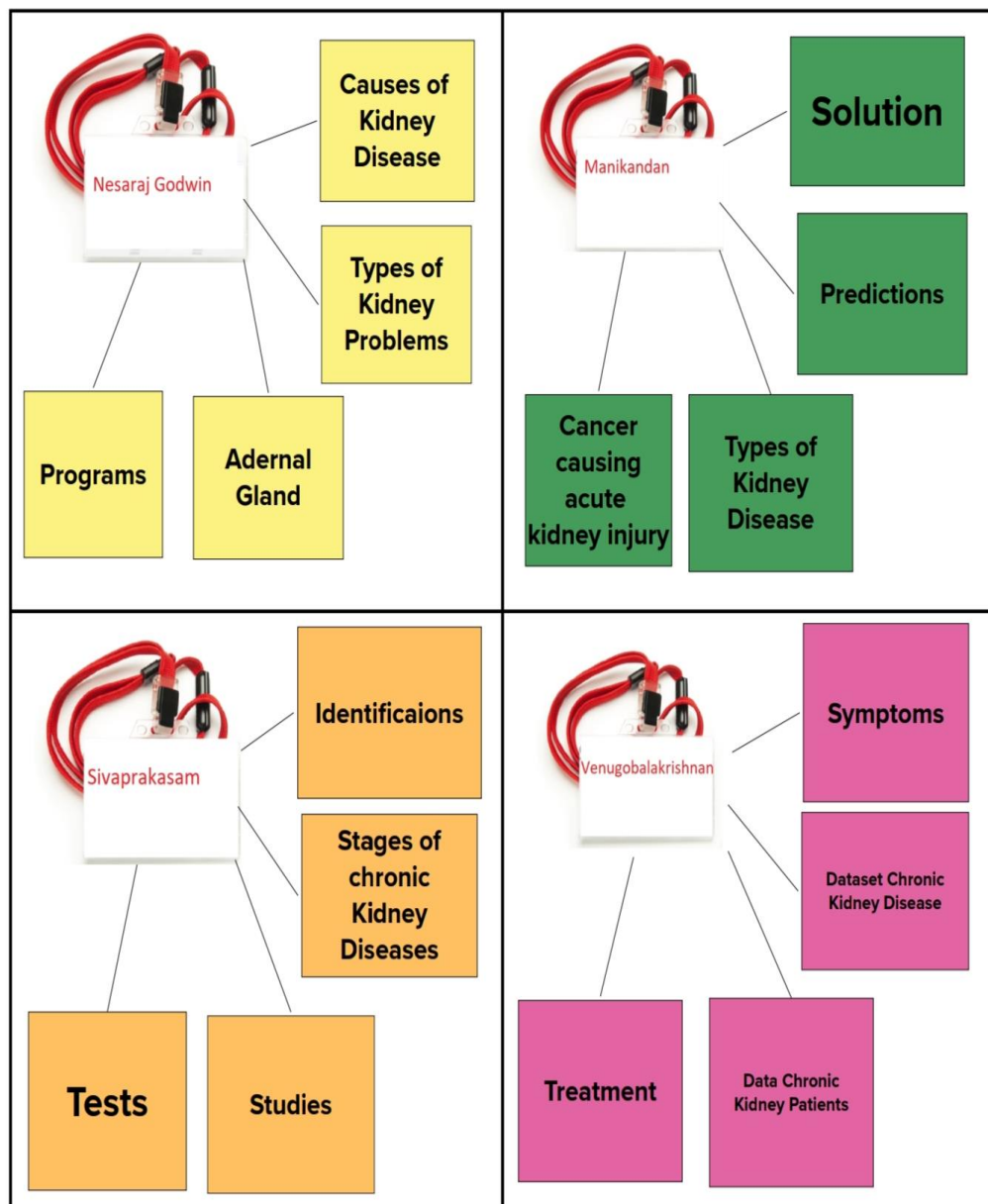


❖ BRAINSTORM:

2 Predictions of Chronic Kidney Disease

Write down any ideas that come to mind that address your problem statement.

2x2 Grid



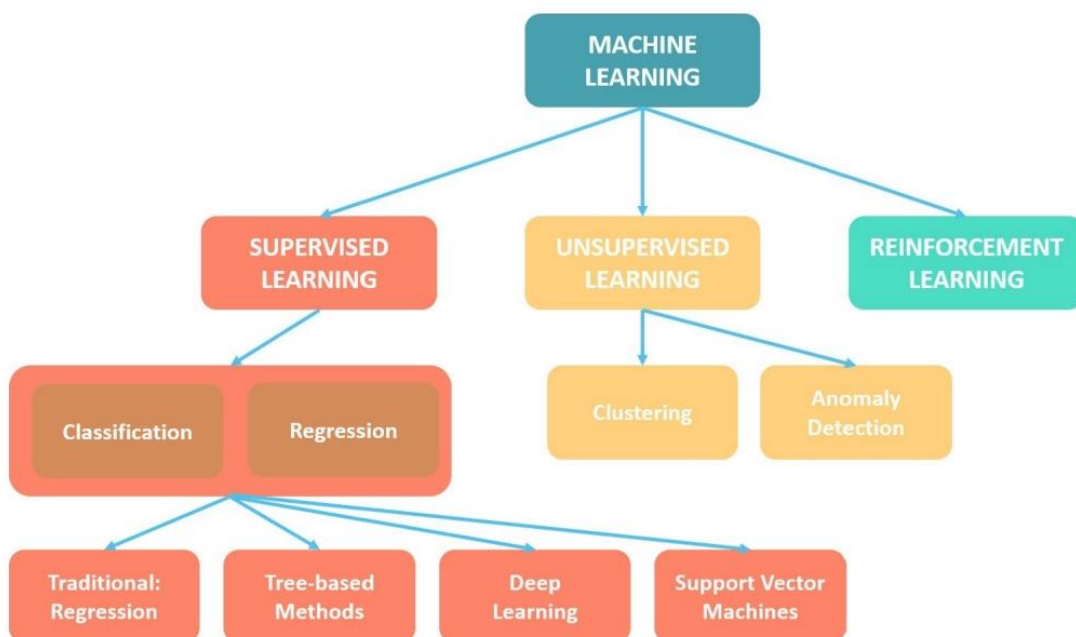
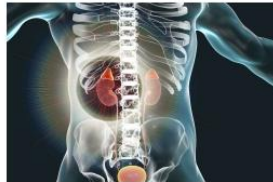
Chronic Kidney Disease



This article is about Chronic Kidney Disease, We discuss to identify the disease by Machine Learning Program

Kidney Gland

A small gland that makes steroid hormones, adrenaline, and noradrenaline. These hormones help control heart rate, blood pressure and other important body functions. There are two adrenal glands, one on top of each kidney. Also called suprarenal gland.

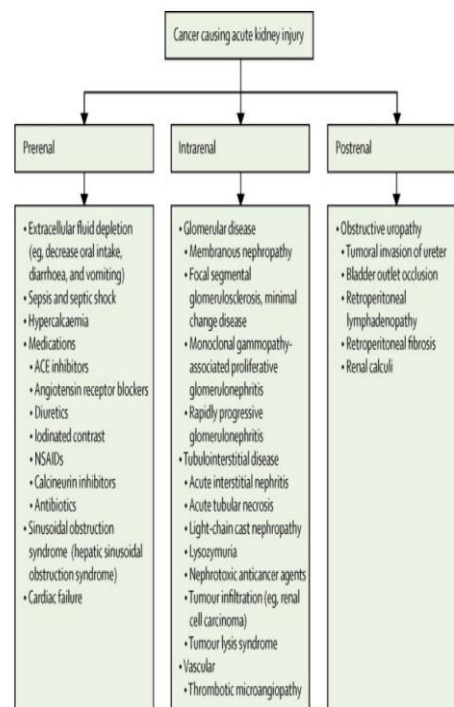
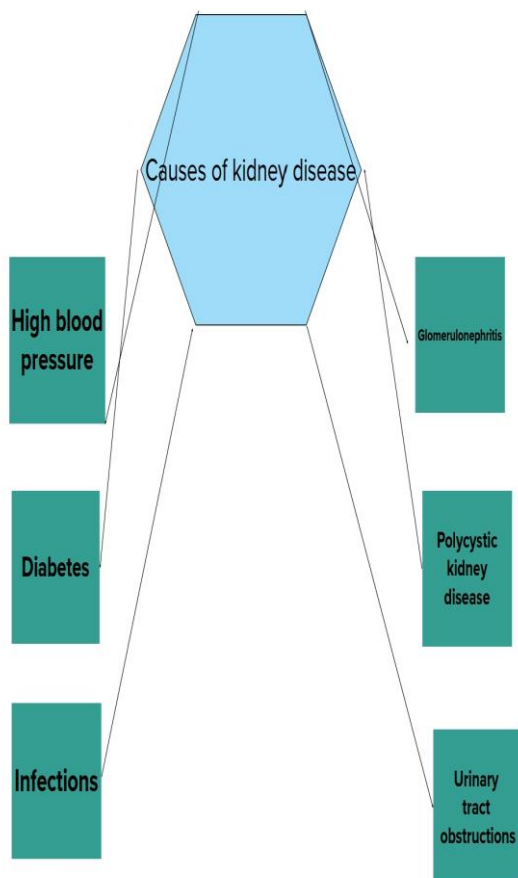
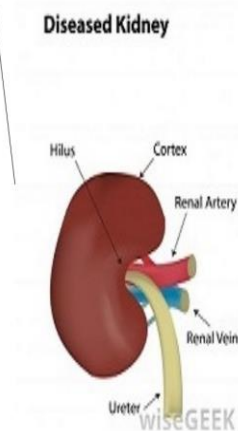
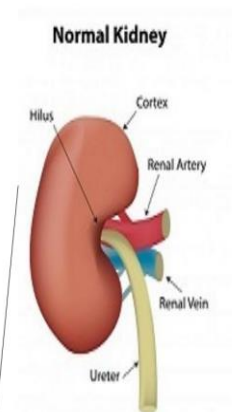
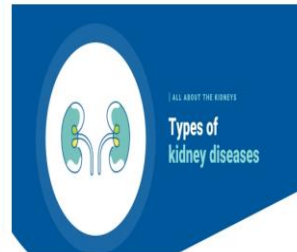
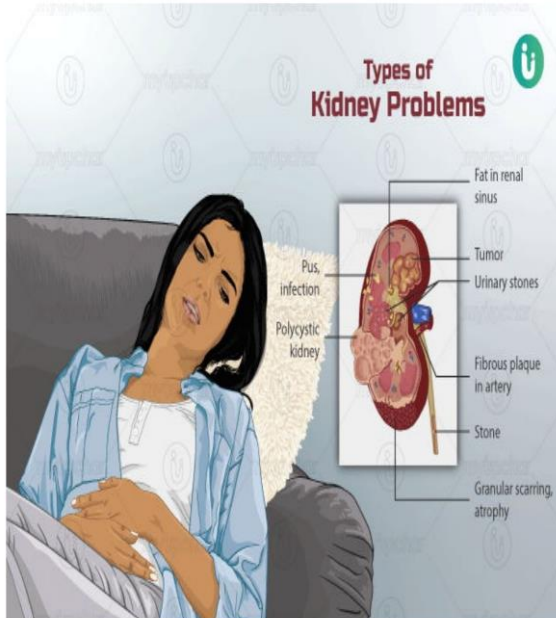


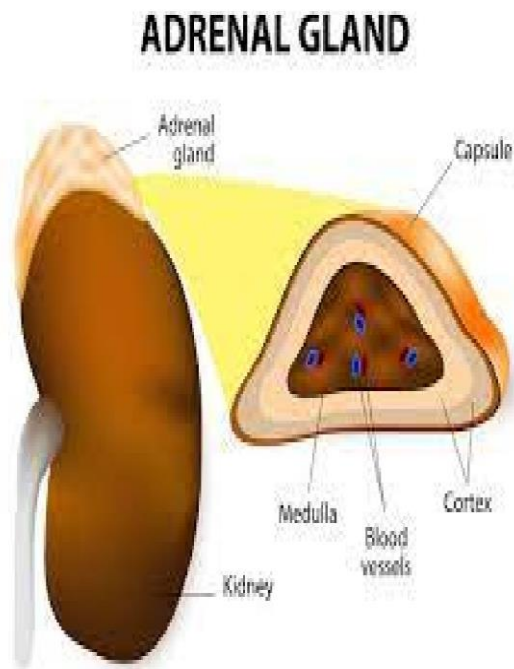
3

Group ideas

Take turns sharing your ideas while clustering similar or related notes as you go. Once all sticky notes have been grouped, give each cluster a sentence-like label. If a cluster is bigger than six sticky notes, try and see if you can break it up into smaller sub-groups.

Type your paragraph...

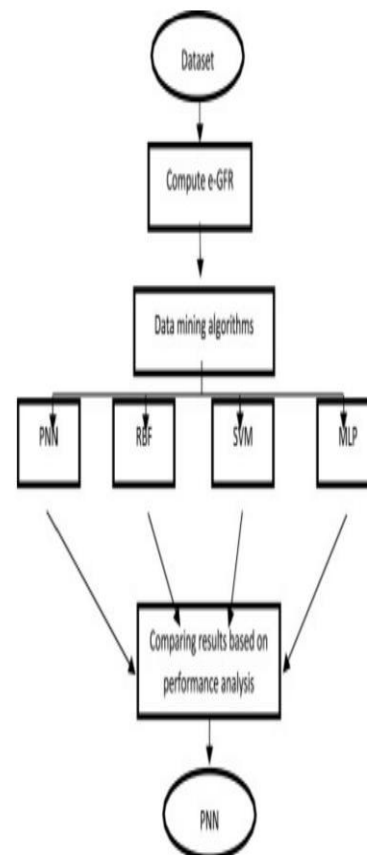
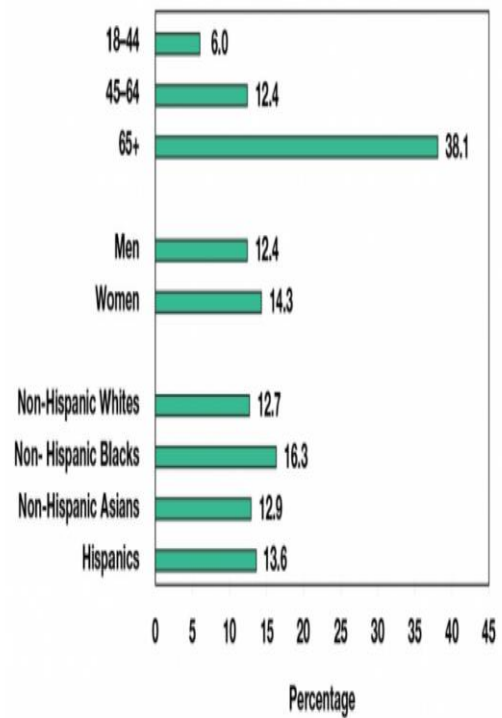




Stages of Chronic Kidney Disease

| Stage of CKD | eGFR result | What it means |
|--------------|--------------|---|
| Stage 1 | 90 or higher | <ul style="list-style-type: none"> - Mild kidney damage - Kidneys work as well as normal |
| Stage 2 | 60-89 | <ul style="list-style-type: none"> - Mild kidney damage - Kidneys still work well |
| Stage 3a | 45-59 | <ul style="list-style-type: none"> - Mild to moderate kidney damage - Kidneys don't work as well as they should |
| Stage 3b | 30-44 | <ul style="list-style-type: none"> - Moderate to severe damage - Kidneys don't work as well as they should |
| Stage 4 | 15-29 | <ul style="list-style-type: none"> - Severe kidney damage - Kidneys are close to not working at all |
| Stage 5 | less than 15 | <ul style="list-style-type: none"> - Most severe kidney damage - Kidneys are very close to not working or have stopped working (failed) |

Data of Kidney Disease in 2021



Chronic Kidney Disease (CKD)

Early Identification and Intervention in Primary Care

Step 1

Identify individuals at risk

Main clinical risk factors for CKD:

- Hypertension
- Diabetes
- CVD
- Family history of CKD

Consider other factors:

- Systemic disease affecting the kidneys (e.g. SLE)
- Obesity
- Genetic risk factors (e.g. ADPKD)
- Environmental exposures to nephrotoxins
- Demographics – older age, race/ethnicity
- History of AKI

Step 2

Test high-risk adults to detect CKD (not population-wide)

Evaluate kidney function – eGFR

- eGFR calculated based on serum creatinine and/or cystatin C

AND Evaluate kidney damage – albuminuria

- UACR or dipstick* (if UACR is unavailable)

If UACR ≥ 30 mg/g (>3 mg/mmol)

OR

eGFR <60 mL/min/1.73 m²

Re-test in 3 months

If low eGFR or high UACR are present for ≥ 3 months, diagnose CKD

If UACR <30 mg/g (<3 mg/mmol)

AND

eGFR >60 mL/min/1.73 m²

Re-test at least once a year*

Step 3

Diagnose CKD

Step 4

Stratify and treat (also see Table 2)

Risk categories for CKD progression, morbidity, and mortality; monitoring frequency (number of check-ups per year in parentheses); and nephrology consultation¹

| eGFR categories (mL/min/1.73 m ²) Description and range | Albuminuria categories | | | Risk category | Monitoring frequency (per year) | Nephrology consultation |
|--|---|--|---|---------------------------|--|-------------------------|
| | A1 Range <30 mg/g <3 mg/mmol | A2 Range 30–299 mg/g 3–29 mg/mmol | A3 Range ≥ 300 mg/g ≥ 30 mg/mmol | | | |
| ≥ 90 G1 | Monitor (1) | Treat (1) | Treat & consult (3) | Low risk | Stable disease OR NO CKD in absence of other markers of kidney damage. ¹ Requires measurements once a year or earlier in case of new symptoms / risk factors. | |
| 60–89 G2 | Monitor (1) | Treat (1) | Treat & consult (3) | Moderately increased risk | Requires measurements at least once a year | |
| 45–59 G3a | Treat (1) | Treat (2) | Treat & consult (3) | High risk | Requires measurements at least twice a year | |
| 30–44 G3b | Treat (2) | Treat & consult (3) | Treat & consult (3) | Very high risk | Treat in agreement with a nephrologist | |
| 15–29 G4 | Treat & consult (3) | Treat & consult (3) | Treat & consult (4+) | | Requires the closest monitoring at least four times a year (every 1–3 months) | |
| <15 G5 | Treat & consult (4+) | Treat & consult (4+) | Treat & consult (4+) | | | |

Adapted from de Zeeuw et al. 2022¹

Step 5

Nephrology consultation

Take action based on the risk categories for CKD progression, morbidity, and mortality, and monitoring frequency (see above).

Primary care practitioners should consult with a nephrologist while initiating treatment; some patients may be under the direct care of a nephrologist if indicated (see Table 3).

Symptoms of Chronic Kidney Disease

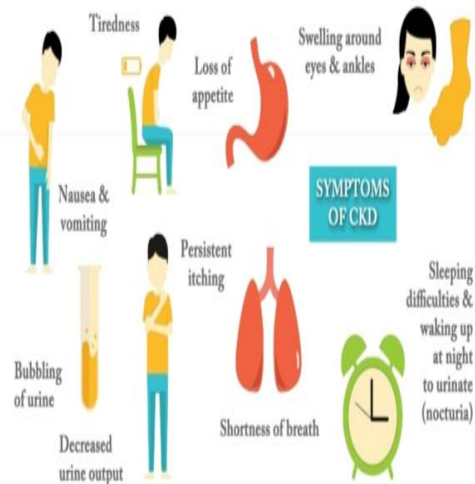


Table 1. Treat to slow CKD progression, reduce mortality risk, and manage comorbidities

| Lifestyle modification | |
|---|---|
| Smoking cessation; regular exercise; well-balanced diet (avoid excessive protein intake and processed food, limit sodium intake <2 g/day) | |
| Medical treatment | |
| Treat diabetes, hypertension, and CVD: Optimise blood pressure and glycemic control | Ensure guideline-directed medical treatment to slow down CKD progression and reduce CVD risk: maximally tolerated doses of ACEi/ARBs, SGLT2 inhibitors, nonsteroidal MRAs with proven benefits in renal and cardiovascular outcome trials for T2D; also consider lipid-lowering therapy (statins) and/or antiplatelet therapy (for patients with CKD at risk of atherosclerotic events) |
| Considerations | |
| Adjust dosing of medications based on eGFR; exercise caution when prescribing analgesics, antimicrobials, hypoglycemics, chemotherapeutics, or anticoagulants; avoid nephrotoxins (e.g. NSAIDs) and some contrast media | |

Table 2. Monitor for CKD progression and comorbidities

| CKD progression and comorbidities | What to monitor |
|-----------------------------------|--|
| CKD monitoring | eGFR, UACR, urinalysis (urine sediment) |
| CVD and dyslipidemia | Blood pressure, cardiovascular risk stratification, lipid status |
| Diabetes | Blood glucose, HbA1c |

Identify CKD complications: anemia, mineral and bone disorders, metabolic acidosis, etc.

Table 3. Additional considerations for nephrology consultation

- Unexplained, progressive decline in eGFR ≥ 5 mL/min/1.73 m² over 12 months or sudden decline in eGFR over days to weeks
- Unexplained significant albuminuria/proteinuria or hematuria
- Persistent hyperkalemia, resistant hypertension (defined as uncontrolled hypertension on three antihypertensive agents, including a diuretic), recurring kidney stones, or hereditary kidney diseases (e.g. ADPKD)
- Other complications identified (anemia, mineral and bone disorders, metabolic acidosis, etc.)

Consultation with a nephrologist can be for identifying other treatable causes or for developing a treatment plan. Although some patients may be maintained further in nephrology care, most will return to primary care.


3 RESULT:

❖ OVERVIEW:

| OBJECT NAME | FIELDS IN THE OBJECT | | | | | | | | | | | | | | | | | | | | | | |
|----------------------|--|-------------|------------|-----------------|------------------------|-----------------|--|----------------------|---------|----------------|---------|------------------|---------|---------------|---------|---------|---------|------------|---------|----------------|----------------------------|----------------|----------------------------|
| PREDICTION | <table><tr><th>FIELD LABEL</th><th>DATA TYPES</th></tr><tr><td>Age</td><td>Numeric</td></tr><tr><td>Gender</td><td>Categorical(Binary:Male/Female)</td></tr><tr><td>Body Mass Index(BMI)</td><td>Numeric</td></tr><tr><td>Blood Pressure</td><td>Numeric</td></tr><tr><td>Serum Creatinine</td><td>Numeric</td></tr><tr><td>Blood Glucose</td><td>Numeric</td></tr><tr><td>Albumin</td><td>Numeric</td></tr><tr><td>Hemoglobin</td><td>Numeric</td></tr><tr><td>Smoking Status</td><td>Categorical(Binary:Yes/No)</td></tr><tr><td>Family History</td><td>Categorical(Binary:Yes/No)</td></tr></table> | FIELD LABEL | DATA TYPES | Age | Numeric | Gender | Categorical(Binary:Male/Female) | Body Mass Index(BMI) | Numeric | Blood Pressure | Numeric | Serum Creatinine | Numeric | Blood Glucose | Numeric | Albumin | Numeric | Hemoglobin | Numeric | Smoking Status | Categorical(Binary:Yes/No) | Family History | Categorical(Binary:Yes/No) |
| FIELD LABEL | DATA TYPES | | | | | | | | | | | | | | | | | | | | | | |
| Age | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Gender | Categorical(Binary:Male/Female) | | | | | | | | | | | | | | | | | | | | | | |
| Body Mass Index(BMI) | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Blood Pressure | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Serum Creatinine | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Blood Glucose | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Albumin | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Hemoglobin | Numeric | | | | | | | | | | | | | | | | | | | | | | |
| Smoking Status | Categorical(Binary:Yes/No) | | | | | | | | | | | | | | | | | | | | | | |
| Family History | Categorical(Binary:Yes/No) | | | | | | | | | | | | | | | | | | | | | | |
| SVM | <table><tr><th>FIELD LABEL</th><th>DATA TYPES</th></tr><tr><td>Support Vectors</td><td>List of Numeric values</td></tr><tr><td>Kernal Function</td><td>Categorical(e.g linear, Polynomial, radial basis function)</td></tr></table> | FIELD LABEL | DATA TYPES | Support Vectors | List of Numeric values | Kernal Function | Categorical(e.g linear, Polynomial, radial basis function) | | | | | | | | | | | | | | | | |
| FIELD LABEL | DATA TYPES | | | | | | | | | | | | | | | | | | | | | | |
| Support Vectors | List of Numeric values | | | | | | | | | | | | | | | | | | | | | | |
| Kernal Function | Categorical(e.g linear, Polynomial, radial basis function) | | | | | | | | | | | | | | | | | | | | | | |

❖ ACTIVITY & SCREENSHOT:

Read the Dataset

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Run Code

```
In [1]: import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt


In [2]: pd.pandas.set_option('display.max_columns', None)

In [3]: dataset = pd.read_csv("kidney_disease.csv")
dataset.head()
```

Out[3]:

| | id | age | bp | sg | al | su | rbc | pc | pcc | ba | bgr | bu | sc | sod | pot | hemo | pcv | wc | rc | htn | dm | cad | appet | pe | ane |
|---|----|------|------|-------|-----|-----|--------|----------|------------|------------|-------|------|-----|-------|-----|------|-----|------|-----|-----|-----|-----|-------|-----|-----|
| 0 | 0 | 48.0 | 80.0 | 1.020 | 1.0 | 0.0 | NaN | normal | notpresent | notpresent | 121.0 | 36.0 | 1.2 | NaN | NaN | 15.4 | 44 | 7800 | 5.2 | yes | yes | no | good | no | no |
| 1 | 1 | 7.0 | 50.0 | 1.020 | 4.0 | 0.0 | NaN | normal | notpresent | notpresent | NaN | 18.0 | 0.8 | NaN | NaN | 11.3 | 38 | 6000 | NaN | no | no | no | good | no | no |
| 2 | 2 | 62.0 | 80.0 | 1.010 | 2.0 | 3.0 | normal | normal | notpresent | notpresent | 423.0 | 53.0 | 1.8 | NaN | NaN | 9.6 | 31 | 7500 | NaN | no | yes | no | poor | no | yes |
| 3 | 3 | 48.0 | 70.0 | 1.005 | 4.0 | 0.0 | normal | abnormal | present | notpresent | 117.0 | 56.0 | 3.8 | 111.0 | 2.5 | 11.2 | 32 | 6700 | 3.9 | yes | no | no | poor | yes | yes |
| 4 | 4 | 51.0 | 80.0 | 1.010 | 2.0 | 0.0 | normal | normal | notpresent | notpresent | 106.0 | 26.0 | 1.4 | NaN | NaN | 11.6 | 35 | 7300 | 4.6 | no | no | no | good | no | no |

Handling Missing Values

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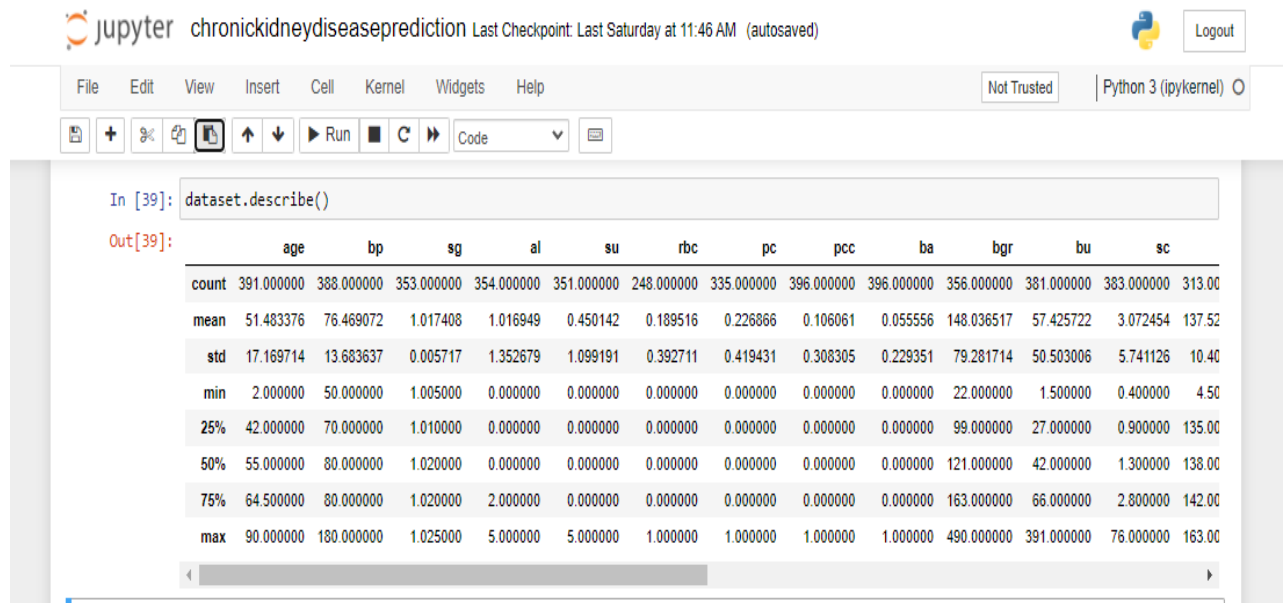
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Run Code

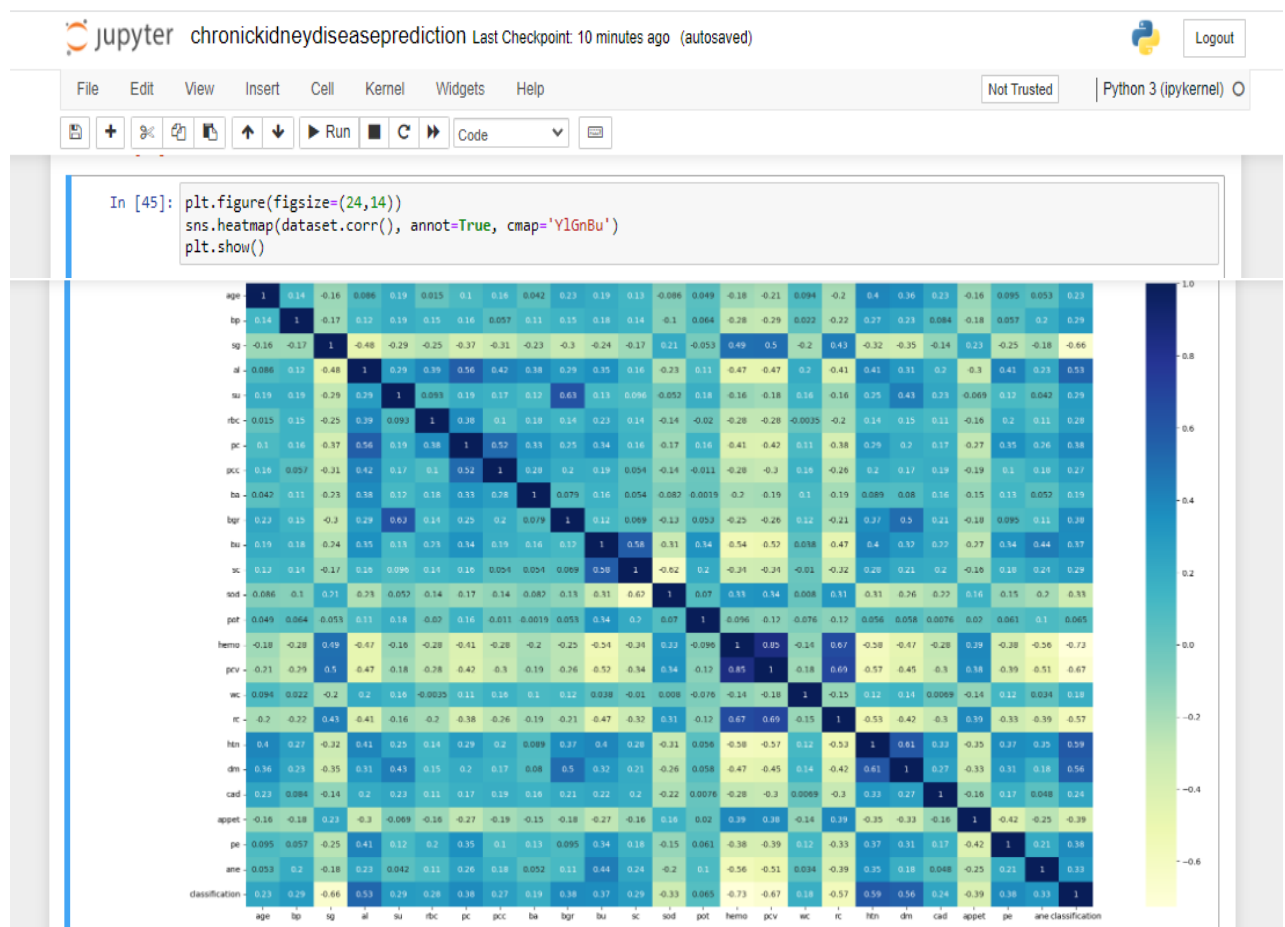
```
In [6]: dataset.isnull().sum()

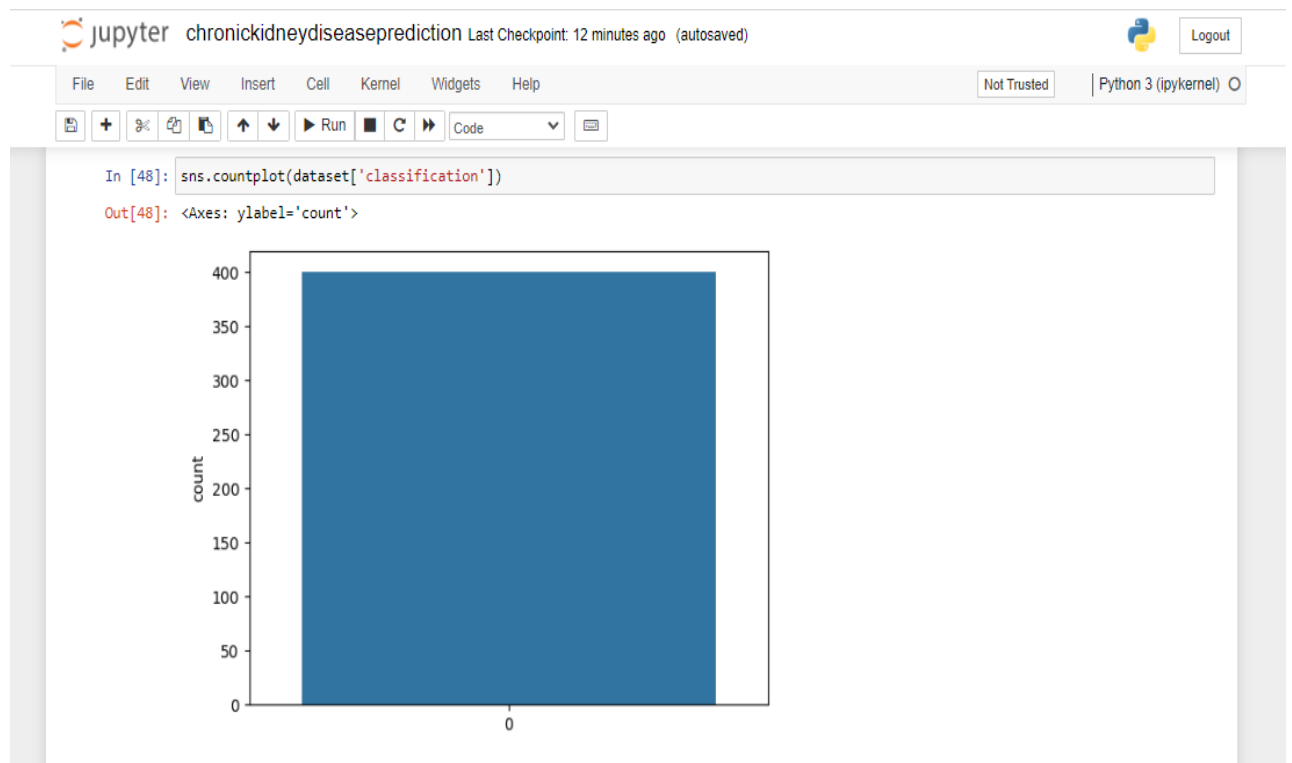
Out[6]:
age          9
bp          12
sg          47
al          46
su          49
rbc         152
pc          65
pcc          4
ba           4
bgr         44
bu          19
sc           17
sod          87
pot          88
hemo         52
pcv          70
wc         105
rc          130
htn           2
dm           2
cad           2
appet         1
pe            1
ane            1
classification 0
dtype: int64
```

DESCRIPTIVE STATISTICAL ANALYSIS:



FINDING CORRELATION BETWEEN THE INDEPENDENT COLUMNS:





SPLITTING THE DATA INTO TRAIN AND TEST:

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In [56]: `from sklearn.model_selection import train_test_split`
`X_train,X_test,y_train,y_test = train_test_split(X,y, test_size=0.3, random_state=33)`

In [57]: `print(X_train.shape)`
`print(X_test.shape)`

(280, 8)
(120, 8)

RANDOM FOREST MODEL :

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```
In [58]: from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
```

```
In [59]: from sklearn.ensemble import RandomForestClassifier
RandomForest = RandomForestClassifier()
RandomForest = RandomForest.fit(X_train,y_train)

y_pred = RandomForest.predict(X_test)

print('Accuracy:', accuracy_score(y_test,y_pred))
print(confusion_matrix(y_test,y_pred))
print(classification_report(y_test,y_pred))
```

Accuracy: 0.975
[[55 3]
 [0 62]]

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 1.00 | 0.95 | 0.97 | 58 |
| 1 | 0.95 | 1.00 | 0.98 | 62 |
| accuracy | | | 0.97 | 120 |
| macro avg | 0.98 | 0.97 | 0.97 | 120 |
| weighted avg | 0.98 | 0.97 | 0.97 | 120 |

DECISION TREE MODEL :

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```
In [51]: from sklearn.ensemble import ExtraTreesClassifier
import matplotlib.pyplot as plt
model=ExtraTreesClassifier()
model.fit(X,y)

plt.figure(figsize=(8,6))
ranked_features=pd.Series(model.feature_importances_,index=X.columns)
ranked_features.nlargest(24).plot(kind='barh')
```

| Feature | Importance |
|---------|------------|
| sg | 0.16 |
| dm | 0.14 |
| htn | 0.12 |
| hemo | 0.12 |
| al | 0.10 |
| rc | 0.09 |
| appet | 0.08 |
| pc | 0.07 |
| bgr | 0.06 |
| bu | 0.05 |
| rbc | 0.04 |
| ane | 0.04 |
| bp | 0.03 |
| sod | 0.03 |
| age | 0.02 |
| su | 0.02 |
| pot | 0.02 |
| wc | 0.02 |
| pcc | 0.02 |
| cad | 0.02 |
| ba | 0.01 |

CLASSIFICATION :

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Code


```
In [92]: from sklearn.ensemble import AdaBoostClassifier
AdaBoost = AdaBoostClassifier()
AdaBoost = AdaBoost.fit(X_train,y_train)

y_pred = AdaBoost.predict(X_test)

print('Accuracy:', accuracy_score(y_test,y_pred))
print(confusion_matrix(y_test,y_pred))
print(classification_report(y_test,y_pred))
```

Accuracy: 1.0
[[58 0]
 [0 62]]

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 1.00 | 1.00 | 1.00 | 58 |
| 1 | 1.00 | 1.00 | 1.00 | 62 |
| accuracy | | | 1.00 | 120 |
| macro avg | 1.00 | 1.00 | 1.00 | 120 |
| weighted avg | 1.00 | 1.00 | 1.00 | 120 |

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Code

```
In [93]: from sklearn.ensemble import GradientBoostingClassifier
GradientBoost = GradientBoostingClassifier()
GradientBoost = GradientBoost.fit(X_train,y_train)

y_pred = GradientBoost.predict(X_test)

print('Accuracy:', accuracy_score(y_test,y_pred))
print(confusion_matrix(y_test,y_pred))
print(classification_report(y_test,y_pred))
```

Accuracy: 0.975
[[55 3]
 [0 62]]

| | precision | recall | f1-score | support |
|--------------|-----------|--------|----------|---------|
| 0 | 1.00 | 0.95 | 0.97 | 58 |
| 1 | 0.95 | 1.00 | 0.98 | 62 |
| accuracy | | | 0.97 | 120 |
| macro avg | 0.98 | 0.97 | 0.97 | 120 |
| weighted avg | 0.98 | 0.97 | 0.97 | 120 |

MODEL DEPLOYMENT

Integrate with Wen Framework

1. Building HTML Pages
2. Building server side script
3. Run the web application

BUILDING HTML PAGES

- app.py
- index.html
- result.html

```
app.py X
1 from flask import Flask, render_template, request
2 import numpy as np
3 import pickle
4
5
6 app = Flask(__name__)
7 model = pickle.load(open('Kidney.pkl', 'rb'))
8
9 @app.route('/', methods=['GET'])
10 def Home():
11     return render_template('index.html')
12
13 @app.route("/predict", methods=['POST'])
14 def predict():
15     if request.method == 'POST':
16         sg = float(request.form['sg'])
17         htn = float(request.form['htn'])
18         hemo = float(request.form['hemo'])
19         dm = float(request.form['dm'])
20         al = float(request.form['al'])
21         appet = float(request.form['appet'])
22         rc = float(request.form['rc'])
23         pc = float(request.form['pc'])
24
25         values = np.array([[sg, htn, hemo, dm, al, appet, rc, pc]])
26         prediction = model.predict(values)
27
28         return render_template('result.html', prediction=prediction)
29
30
31 if __name__ == "__main__":
32     app.run(debug=True)
33
34
```

...ease-Prediction-Project-main (1)\Chronic-Kidney-Disease-Prediction-Project-main\templates\index.html

app.py × index.html × result.html ×

```
1 <!DOCTYPE html>
2 <html lang="en">
3
4 <head>
5 <meta charset="UTF-8">
6 <title>Chronic Kidney Disease Model</title>
7
8 </head>
9 <body>
10 <div style="color:black;" class="container">
11 <h2 class='container-heading'><span class="heading_font">Chronic Kidney Disease Prediction
12 </div>
13
14 <div style="color:black;" class="ml-container">
15 <form action="{{ url_for('predict') }}" method="POST">
16 <br>
17 <br>
18 <h3>Specific Gravity</h3>
19 <input id="first" name="sg" placeholder="Ex: (1.005,1.010,1.015,1.020,1.025)" required
20 <br>
21 <h3>Hyper Tension</h3>
22 <input id="second" name="htn" placeholder="Yes = 1, No=0" required="required">
23 <br>
24 <h3>Hemoglobin</h3>
25 <input id="third" name="hemo" placeholder="in gms" required="required">
26 <br>
27 <h3>Diabetes Mellitus</h3>
28 <input id="fourth" name="dm" placeholder="Yes = 1, No=0" required="required">
29 <br>
30 <h3>Albumin</h3>
31 <input id="fifth" name="al" placeholder="(0,1,2,3,4,5)" required="required">
32 <br>
33 <h3>Appetite</h3>
34 <input id="sixth" name="appet" placeholder="Good = 1, Poor = 0" required="required">
35 <br>
```

```
36 <h3>Red Blood Cell Count</h3>
37 <input id="seventh" name="rc" placeholder="in Millions/cmm" required="required">
38 <h3>Pus Cell</h3>
39 <input id="eight" name="pc" placeholder="Normal = 0, Abnormal = 1" required="required"
40 <br>
41 <br>
42 <br>
43 <button id="sub" type="submit ">Submit</button>
44 <br>
45 <br>
46 <br>
47 <br>
48 </form>
49 </div>
50
51
52
53
54 <style>
55 /* Background Image */
56 body
57 {
58 background-image:url("https://raw.githubusercontent.com/SagarDhandare/Chronic-Kidney-Disease-Predi
59 height: 100%;
60
61
62 /* Center and scale the image nicely */
63 background-position: center;
64 background-repeat: no-repeat;
65 background-size: 100% 100%;
66
67
68
69 /* Color */
70 body{
```

```
71 font-family: Arial, Helvetica,sans-serif;
72 text-align: center;
73 margin: 0;
74 padding: 0;
75 width: 100%;
76 height: 100%;
77 display: flex;
78 flex-direction: column;
79 }
80
81
82 /* Heading Font */
83 .container-heading{
84 margin: 0;
85 }
86
87 .heading_font{
88 color: #black;
89 font-family: 'Pacifico', cursive;
90 font-size: 50px;
91 font-weight: normal;
92 }
93
94
95
96 /* Box */
97 #first {
98 border-radius: 14px;
99 height: 30px;
100 width: 300px;
101 font-size: 18px;
102 text-align: center;
103 }
104
105 #second {
```

```

106         border-radius: 14px;
107         height: 25px;
108         width: 160px;
109         font-size: 20px;
110         text-align: center;
111     }
112
113     #third {
114         border-radius: 14px;
115         height: 25px;
116         width: 120px;
117         font-size: 20px;
118         text-align: center;
119     }
120
121     #fourth {
122         border-radius: 14px;
123         height: 25px;
124         width: 160px;
125         font-size: 20px;
126         text-align: center;
127     }
128
129     #fifth {
130         border-radius: 14px;
131         height: 25px;
132         width: 130px;
133         font-size: 20px;
134         text-align: center;
135     }
136
137     #sixth {
138         border-radius: 14px;
139         height: 25px;
140         width: 200px;

```

```

141         font-size: 20px;
142         text-align: center;
143     }
144
145     #seventh {
146         border-radius: 14px;
147         height: 25px;
148         width: 180px;
149         font-size: 20px;
150         text-align: center;
151     }
152
153     #eight {
154         border-radius: 14px;
155         height: 25px;
156         width: 260px;
157         font-size: 20px;
158         text-align: center;
159     }
160
161     /* Submit Button */
162     #sub {
163         width: 120px;
164         height: 43px;
165         text-align: center;
166         border-radius: 14px;
167         font-size: 18px;
168     }
169
170
171
172
173
174     <link rel="stylesheet" href="https://cdnjs.cloudflare.com/ajax/libs/font-awesome/4.7.0/css/font-aw
175

```

```
</style>
```

```
</body>
```

```
</html>
```

Chronic Kidney Disease Prediction

Specific Gravity

1020

Hyper Tension

0

Hemoglobin

15.8

Diabetes Mellitus

0

Albumin

1

Appetite

1

Red Blood Cell Count

6.1

Pus Cell

0

Submit


```
...ase-Prediction-Project-main (1)\Chronic-Kidney-Disease-Prediction-Project-main\templates\result.html
index.html × result.html ×
1 <!DOCTYPE html>
2 <html lang="en">
3
4 <head>
5 <meta charset="UTF-8">
6 <meta name="viewport" content="width=device-width, initial-scale=1.0">
7 <title>Chronic Kidney Disease Result</title>
8 </head>
9
10 <body>
11
12 <div style="color:black;" class="container">
13 <form action="{% url_for('predict') %}" method="post">
14 <h2 class='container-heading'><span class="heading_font">Chronic Kidney Disease Predict
15
16 <br><br><br><br>
17
18 <!-- Result -->
19 <div style="color:black;" class="results">
20 {% if prediction==1 %}
21 <h1><span class='danger'>Oops! 😞<br><br>You have CHRONIC KIDNEY DISEASE.<br><br>Pl
22 <br><br><br><br><br><br>
23 <span class='safe'>🎉 Congratulation! 🎉<br><br>You DON'T have Chronic Kidney I
26 
29 </form>
30
31 </div>
32 <div>
33 <br><br> <br><br><br><br><br><br><br><br>
34 </div>
35
```

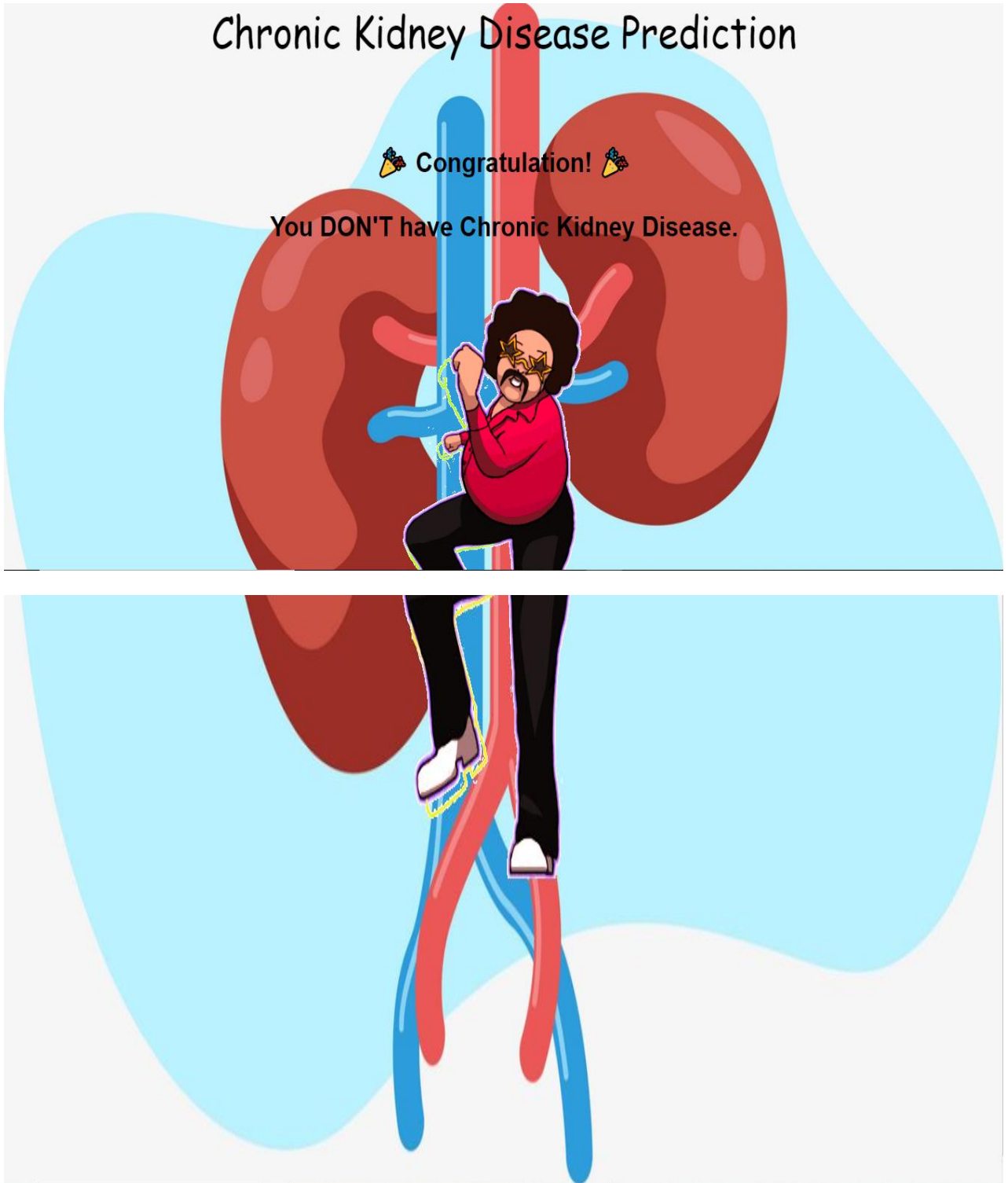
```
36
37 <style>
38
39 /* Background Image */
40 body
41 {
42 background-image:url("https://raw.githubusercontent.com/SagarDhandare/Chronic-Kidney-Disease-Predic
43 height: 100%;
44
45 /* Center and scale the image nicely */
46 background-position: center;
47 background-repeat: no-repeat;
48 background-size: 100% 100%;
49
50 }
51
52 /* Color */
53 body{
54 font-family: Arial, Helvetica,sans-serif;
55 text-align: center;
56 margin: 0;
57 padding: 0;
58 width: 100%;
59 height: 100%;
60 display: flex;
61 flex-direction: column;
62 }
63
64
65 /* Heading Font */
66 .container-heading{
67 margin: 0;
68 }
69
70 .heading_font{
```

```
71 color: #black;
72 font-family: 'Pacifico', cursive;
73 font-size: 50px;
74 font-weight: normal;
75 }
76
77
78
79 <link rel="stylesheet" href="https://cdn.jsdelivr.net/npm/font-awesome@4.7.0/css/font-awe
80 </style>
81
82
83
84 </body>
85
86 </html>
```

Chronic Kidney Disease Prediction

🎉 Congratulation! 🎉

You DON'T have Chronic Kidney Disease.



4 TRAILHEAD PROFILE PUBLIC URL:

Team Lead- <https://trailblazer.me/id/mastergodwin>
Team Member 1- <https://trailblazer.me/id/mkandanv1>
Team Member 2- <https://trailblazer.me/id/sprakasam6>
Team Member 3-. <https://trailblazer.me/id/vgobalakrishnan>

5 ADVANTAGES AND DISADVANTAGES:

❖ ADVANTAGES:

- ***Early detection: Machine learning models can predict the likelihood of developing CKD based on a set of patient characteristics, such as age, sex, medical history, and laboratory test results. Early detection can lead to early intervention and management, which can help slow or halt the progression of the disease.***
- ***Improved accuracy: Machine learning algorithms are designed to analyze vast amounts of data and identify patterns that may not be apparent to human observers. This can improve the accuracy of CKD diagnosis and reduce the risk of misdiagnosis.***
- ***Personalized treatment: Machine learning models can be used to identify the most effective treatment options for individual patients based on their specific characteristics.***

This can lead to more personalized and effective treatment plans.

- ***Cost-effective: Machine learning algorithms can process large amounts of data quickly and efficiently, reducing the time and cost of traditional diagnostic methods.***

❖ DISADVANTAGES:

- ***Data quality: Machine learning models require high-quality data to make accurate predictions. Poor quality data can result in inaccurate predictions and unreliable outcomes.***
- ***Bias: Machine learning algorithms may be biased towards certain groups of patients, leading to unfair or discriminatory outcomes. It is essential to ensure that the data used to train the algorithm is diverse and representative of the population being studied.***
- ***Interpretability: Machine learning models can be difficult to interpret, making it challenging to understand how they arrive at their predictions. This can be a concern for clinicians who may need to explain the reasoning behind the algorithm's predictions to patients.***
- ***Privacy concerns: Machine learning models require large amounts of patient data to train and test the algorithm. There are concerns about how this data is collected, stored, and used, and there is a need to ensure that patient privacy is protected.***

6 APPLICATIONS:

- ***Logistic Regression: Logistic regression is a statistical model used to analyze the relationship between a dependent***

variable and one or more independent variables. In CKD prediction, it can be used to identify the risk factors for the disease.

- ***Decision Trees: Decision trees are a type of machine learning algorithm that can be used for classification and regression analysis. They are useful in identifying the most important risk factors for CKD and predicting the likelihood of the disease.***
- ***Random Forests: Random forests are an ensemble learning method that uses multiple decision trees to improve the accuracy of predictions. They are used in CKD prediction to identify the most significant risk factors and their interactions with other variables.***
- ***Support Vector Machines (SVM): SVM is a machine learning algorithm that can be used for classification and regression analysis. It is useful in predicting the likelihood of CKD based on patient data and identifying the most important risk factors for the disease.***
- ***Neural Networks: Neural networks are a type of machine learning algorithm that uses artificial intelligence to learn from data. They are useful in CKD prediction as they can identify complex relationships between various patient factors and predict the likelihood of the disease.***

7 CONCLUSION:

Neural Networks: Neural networks are a type of machine learning algorithm that uses artificial intelligence to learn from data. They are useful in CKD prediction as they can identify complex relationships between various patient factors and predict the likelihood of the disease.

8 FEATURE SCOPE:

- **Demographic data:** Age, gender, race, and ethnicity can all be considered as potential risk factors for CKD
- **Medical history:** Previous diagnoses of hypertension, diabetes, cardiovascular disease, and other chronic conditions are commonly considered as risk factors for CKD.
- **Laboratory test results:** Measures of kidney function, such as serum creatinine and estimated glomerular filtration rate (eGFR), as well as electrolyte levels and other blood chemistry values, can provide insight into the patient's kidney health. **Clinical examination data:** Blood pressure, body mass index (BMI), and other physical measurements can provide additional information about a patient's overall health status.
- **Medication history:** Some medications can cause or exacerbate kidney damage, so medication history can also be considered as a risk factor for CKD.
- **Environmental factors:** Exposure to environmental toxins such as lead, cadmium, and arsenic can increase the risk of CKD, so environmental factors may also be considered.