EARLY DETECTION OF CHRONIC KIDNEY DISEASE USING MACHINE LEARNING

TEAM ID : PNT2022TMID48329

A PROJECT REPORT

Submitted by

M. PRIYADHARSHINI - 912019106701

V. KRISHNAKANTH - 912019104701

P. SWETHA - 912019106702

M. LOGANAYAKI - 912019104702

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1. INTRODUCTION

Project Overview

This article objects to predict Chronic Kidney Disease based on full features and important features of CKD dataset. Machine learning technique has become reliable for medical treatment. With the help of a machine learning classifier algorithms, the doctor can detect the disease on time. For this perspective, Chronic Kidney Disease prediction has been discussed in this article.. The important feature selection technique was also applied to the dataset. For each classifier, the results have been computed based on (i) full features, (ii) correlation-based feature selection, (iii) Wrapper method feature selection, (iv) Least absolute shrinkage and selection operator regression, (v) synthetic minority over-sampling technique with least absolute shrinkage and selection operator regression selected features, (vi) synthetic minority oversampling technique with full features. Along with machine learning models one deep neural network has been applied on the same dataset and it has been noted that

deep neural network achieved the highest accuracy of 99.6%.

Purpose

This research article primarily aims to predict whether a person has Chronic Kidney Disease or not. In this perception, seven different machine learning classifiers were applied on the dataset. All the algorithms were running with both full features and selected features. SMOTE was used for over sampling and all the results were recorded. All the machine learning model results were also compared with one deep neural network algorithm. Deep learning neural network was used with two hidden layers.

2. LITERATURE SURVEY

Existing Problem

SURVEY 1 : AUTHORS: Himanshu Kriplani, Bhumi Patel and Sudipta Roy

TITLE : Prediction of Chronic Kidney Diseases Using Deep Artificial Neural Network Technique.

METHODS : This project presents a method to detect the chronic kidney disease and methodologies to diagnose chronic kidney disease is a challenging problem which can reduce the cost of treatment. We studied 224 records of chronic kidney disease available on the UCI machine learning repository named chronic kidney diseases dating back to 2015. Our proposed method is based on deep neural network which predicts the presence or absence of chronic kidney disease with an accuracy of 97%. Compared to other available algorithms, the model we built shows better results which is implemented using the cross-validation technique to keep the model safe from over fitting. This automatic chronic kidney disease treatment helps reduce the kidney damage progression, but for this chronic kidney disease detection at initial stage is necessary.

SURVEY 2 :

AUTHORS : Hongquan Peng , Haibin Zhu , Chi WaAo Ieong , Tao Tao , Tsung Yang Tsai , Zhi Liu

TITLE : A two-stage neural network prediction of chronic kidney disease

METHODS : This paper presents a method to detect chronic kidney disease (CKD) plays a pivotal role in early diagnosis and treatment. Measured glomerular

filtration rate (mGFR) is considered the benchmark indicator in measuring the kidney function. However, due to the high resource cost of measuring mGFR, it is usually approximated by the estimated glomerular filtration rate, underscoring an urgent need for more precise and stable approaches. With the introduction of novel machine learning methodologies, prediction performance is shown to be significantly improved across all available data, but the performance is still limited because of the lack of models in dealing with ultra-high dimensional datasets. This study aims to provide a two-stage neural network approach for prediction of GFR and to suggest some other useful biomarkers obtained from the blood metabolites in measuring GFR. It is a composite of feature shrinkage and neural network when the number of features is much larger than the number of training samples. The results show that the proposed method outperforms the existing ones, such as convolution neural network and direct deep neural network.

SURVEY 3 :

AUTHORS : Deepak K N , Adhwaidh P S , Akshay P D , Athira K S , Jisna Jayan

TITLE : Chronic Kidney Disease Prediction system using Machine Learning

METHODS : This paper reviews and analyzes the Chronic kidney disease (CKD)

is a global health issue that causes a high rate of morbidity and mortality, as well as the onset of additional diseases. Because there are no clear symptoms in the early stages of CKD, people frequently miss it. Early identification of CKD allows patients to obtain timely treatment to slow the disease's progression. Due to their rapid and precise recognition capabilities, machine learning models can successfully assist doctors in achieving this goal. We propose a machine learning framework for diagnosing CKD in this paper. The CKD data set was taken from kaggle, which has a substantial number of missing values. We employ multiple machine learning methods such as DT, SVM, and DNN to analyze data from CKD patients with 21 characteristics and 400 records. The dataset is preprocessed by filling in missing data and normalizing it. To increase accuracy and save training time, the most relevant features from the dataset are chosen. Image processing and letter recognition are used to automatically input the attributes.

References

1.Q.-L. Zhang and D. Rothenbacher, ‘ Prevalence of chronic kidney disease in population-based studies: Systematic review,’ BMC Public Health, vol. 8, no. 1, p. 117, Dec. 2008.

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Cushman, L. A. McClure, B. B. Newsome, and G. Howard, ‘ Albuminuria and racial disparities in the risk for ESRD,’ J. Amer. Soc. Nephrol., vol. 22, no. 9, pp. 1721–1728, Aug. 2011.

3. M. K. Haroun, ‘ Risk factors for chronic kidney disease: A prospective study of 23,534 men and women in Washington County, Maryland,’ J. Amer. Soc. Nephrol., vol. 14, no. 11, pp. 2934–2941, Nov. 20.

4. W. D. Souza, L. C. D. Abreu, L. G. D. SilvaI, and I. M. P. Bezerra, ‘ Incidence of chronic kidney disease hospitalisations and mortality in Espírito Santo between 1996 to 2017,’ WisitCheungpasitporn, Univ. Mississippi Medical Center, Rochester, MN, USA, Tech. Rep., 2019, doi: 10.1371/journal.pone.0224889.

5. W. Mula-Abed, K. A. Rasadi, and D. Al-Riyami, ‘ Estimated glomerular filtration rate (eGFR): A serum creatinine-based test for the detection of chronic kidney disease and its impact on clinical practice,’ Oman Med. J., vol. 27, no. 4, pp. 339–340, 2012.

6. A. S. Levey, D. Cattran, A. Friedman, W. G. Miller, J. Sedor, K. Tuttle, B. Kasiske, and T. Hostetter, ‘ Proteinuria as a surrogate outcome in CKD: Report of a scientific workshop sponsored by the national kidney foundation and the US food

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Problem Statement Definition

To identify and manage patients who have early stages of chronic kidney disease may slow or prevent the progression to end stage kidney disease and reduce cardiovascular complications caused due to diabetes and high blood.

Who does the problem affect?

Chronic Kidney Disease is more common in people aged 65years or older (38%) than in people aged 45–64 years (12%) or 18–44 years (6%). CKD is slightly more common in women in (14%)than men (12%).

What are the boundaries of the problem?

A disease in GFR may also be a marker of kidney disease and precedes the onset of kidney failure . Below 60 ml/min/1.73 m , the prevalence of complications of CKD Increases , as does the risk of cardiovascular disease . Albuminuria (ACR ≥ 30 mg/g), Urine sediment abnormalities, Electrolyte and other abnormalities due to

tubular disorders, Abnormalities detected by histology, Structural abnormalities detected by imaging, History of kidney transplantation.

What is the issue?

Kidneys are damaged and can't filter blood the way they should. The disease is called “chronic” because the damage to your kidneys happens slowly over a long period of time. Some of the common health problems caused by kidney disease include gout, anemia, bone disease, heart disease and fluid buildup.

Where is the issue occurring?

The kidneys grow larger and gradually lose the ability to function as they should. Chronic kidney disease occurs when a disease or condition impairs kidney function, causing kidney damage and high blood pressure.

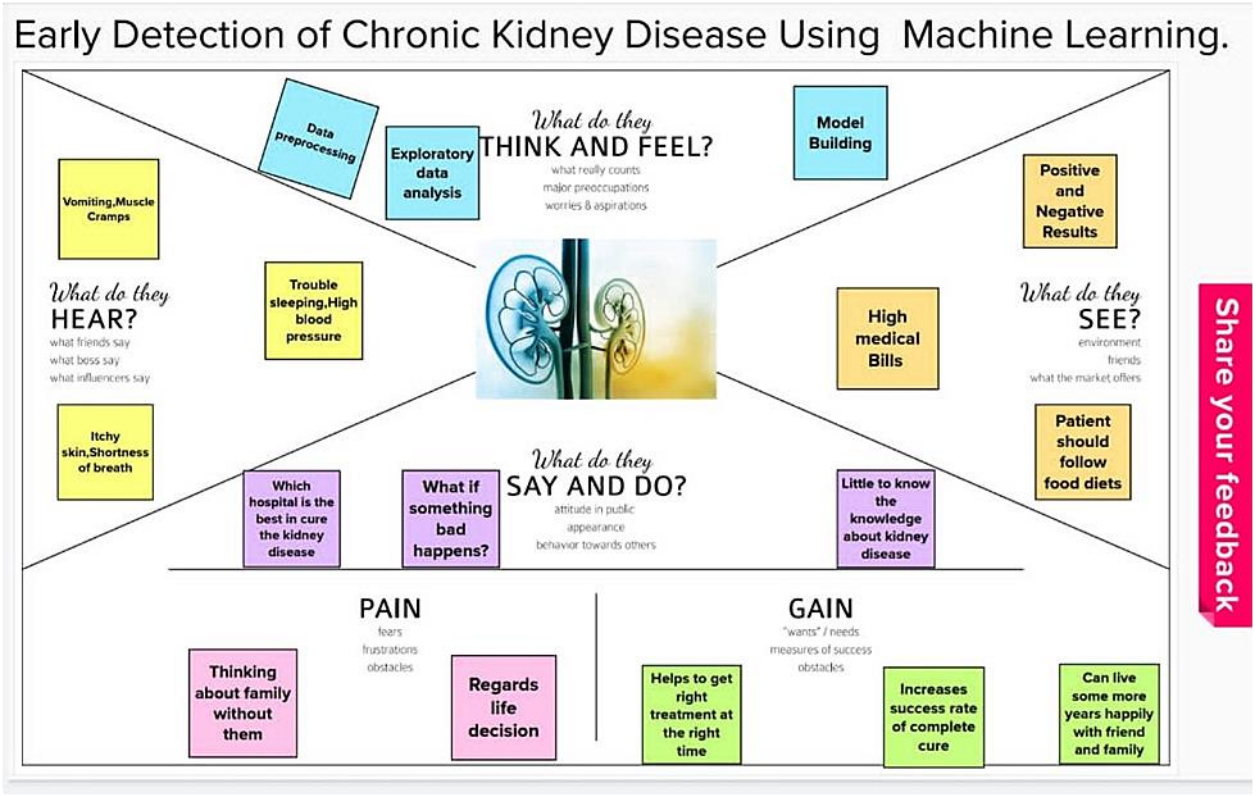
Why is it important that we fix the problem?

The older you get the more likely you are to have some degree of kidney disease.

This is important because CKD increases the risk of heart attack and stroke.

3. IDEATION & PROPOSED SOLUTION

Empathy Map Canvas



Ideation and Brainstorming:

1

Define your problem statement

What problem are you trying to solve? Frame your problem as a How Might We statement. This will be the focus of your brainstorm.

5 minutes

PROBLEM
How do we construct a model that predicts the chronic kidney disease?

PROBLEM
What causes kideney disease?

PROBLEM
How will the final output be like?

PROBLEM
Which algorithm is to be used to predict the disease?

2

Brainstorm

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

10 minutes

DHANA LAKSHMI R

By collecting details of an affected person and comparing it

Algorithm that may give higher accuracy can be used in the model

Based on certain diagnostic measures like Blood Pressure, albumin

We have to collect relevant dataset and decide an approach

ARTHI.S

Motivate the people to take test even for small symptoms

An Algorithm to be framed once the approach gets decided

Give awareness to people to maintain a healthy diet

Guiding the people to have a hygienic food

DILIP.G

Too much protein, glucose, or blood in your urine will be shown by urine test

Blood pressure readings show whether your blood pressure is abnorm

Discover patterns with the help of statistical/ historical data and train the model to predict CKD

The result of the prediction will be whether the person has CKD or not

SIVA SURIYAN M.S

Check the family history of a person if any person got affected

Different algorithms can be tested to find the correlation between various attributes

A panel of qualified doctors can be consulted to identify attributes of greatest significance

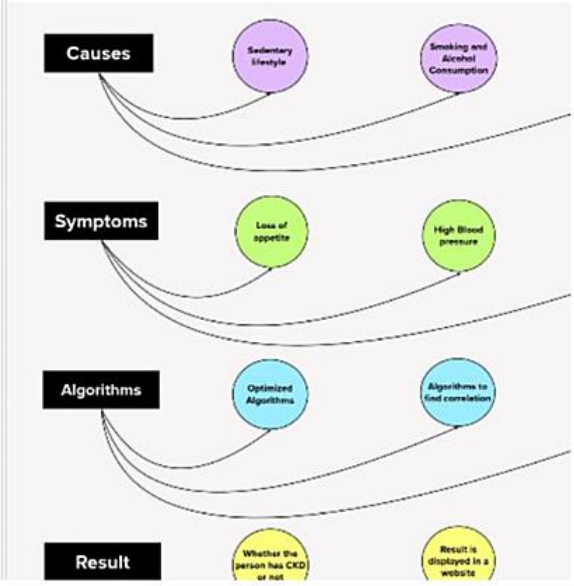
Smoking and excessive consumption alcohol increases blood pressure and cholesterol

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 and break it up into smaller sub-groups.

20 minutes

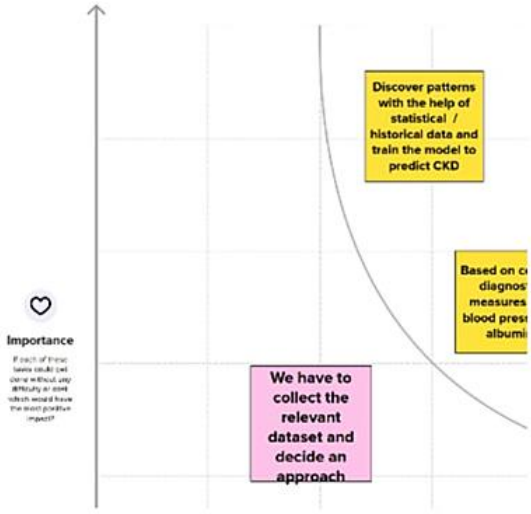


4

Prioritize

Your team should all be on the same page about what's important moving forward. Place your ideas on this grid to determine which ideas are important and which are feasible.

20 minutes



Proposed Solution

Problem Statement

Chronic Kidney Disease is a major concern for the global health care system. Chronic Kidney Disease is now wreaking havoc on society and is spreading at an alarming rate. Various efforts have been undertaken to advance early therapy to prevent the condition from progressing to CKD. Recent research suggest that some of the negative outcomes can be avoided with early identification and treatment.

Feasibility of idea

To predict the early onset of CKD, three Machine Learning techniques are used: ->Random Forest, Decision Tree, Support Vector Machines. Using these techniques, each algorithm's effectiveness is evaluated and the prediction of how many people have been affected by CKD is identified.

Novelty

The renal patient is recognized by undertaking two primary tests.

->A Blood Test to determine Glomerular FilterationRate(GFR).

->A Urine Test to determine Albumin.

Social Impact

As people don't undergo the general test of their health, early detection of CKD is not identified. This creates a great social impact of not being aware of CKD. As a result of this many people are getting affected by CKD.

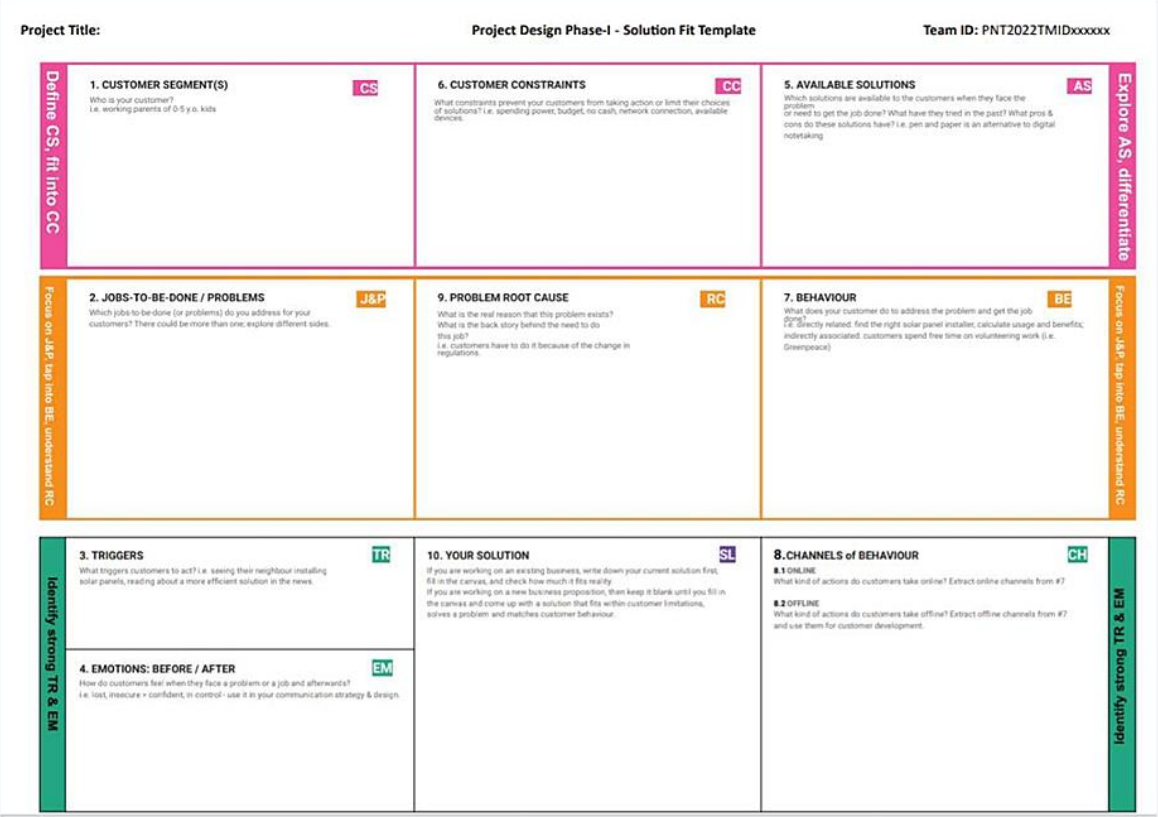
Business Model

The widespread use of Machine Learning of predicting the CKD in the Medical Industries promotes medical innovation, lowers medical expenses, and improves medical quality. In order to cure the CKD patients, the hospitals have been gaining business profit in recovering the patients.

Scalability of solution

This Chronic Kidney Disease have been spreading widely now a days. Early prediction of CKD using Machine Learning that is more efficient to analyze the disease so that it can be cured on time.

Problem Solution Fit



4. REQUIREMENTS ANALYSIS

Functional Requirements

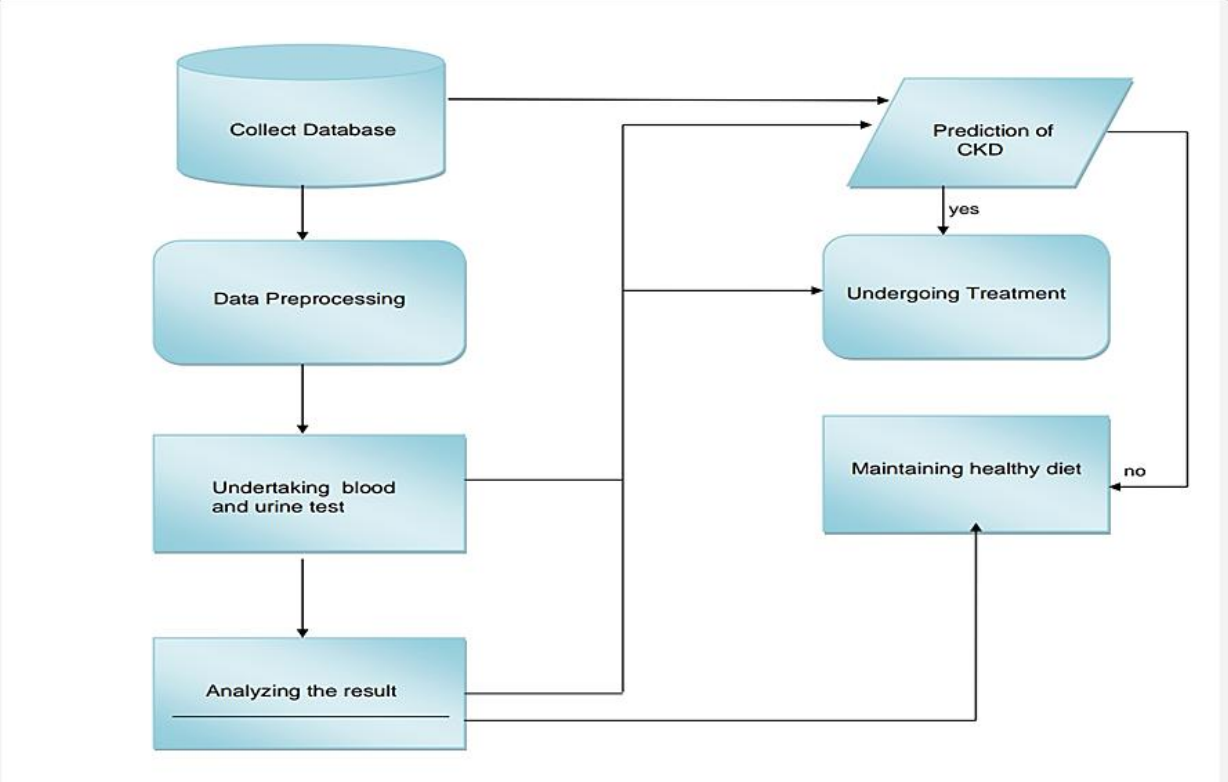
FR No.	Functional Requirement (Epic)	Sub Requirement (Story / Sub-Task)
FR 1	User Page	Users can go to home pag
FR 2	Details	Users can know more details about the CKD.
FR 3	User Requirements	Store past records Generate report for presence of CKD Diagnostic remedies for symptoms
FR 4	User Entry	Input form for pre-diagnostic test results
FR 5	Business Requirements	Quick diagnosis for CKD
FR 6	User Feedback	Allows users to submit feedback

Non- Functional Requirements

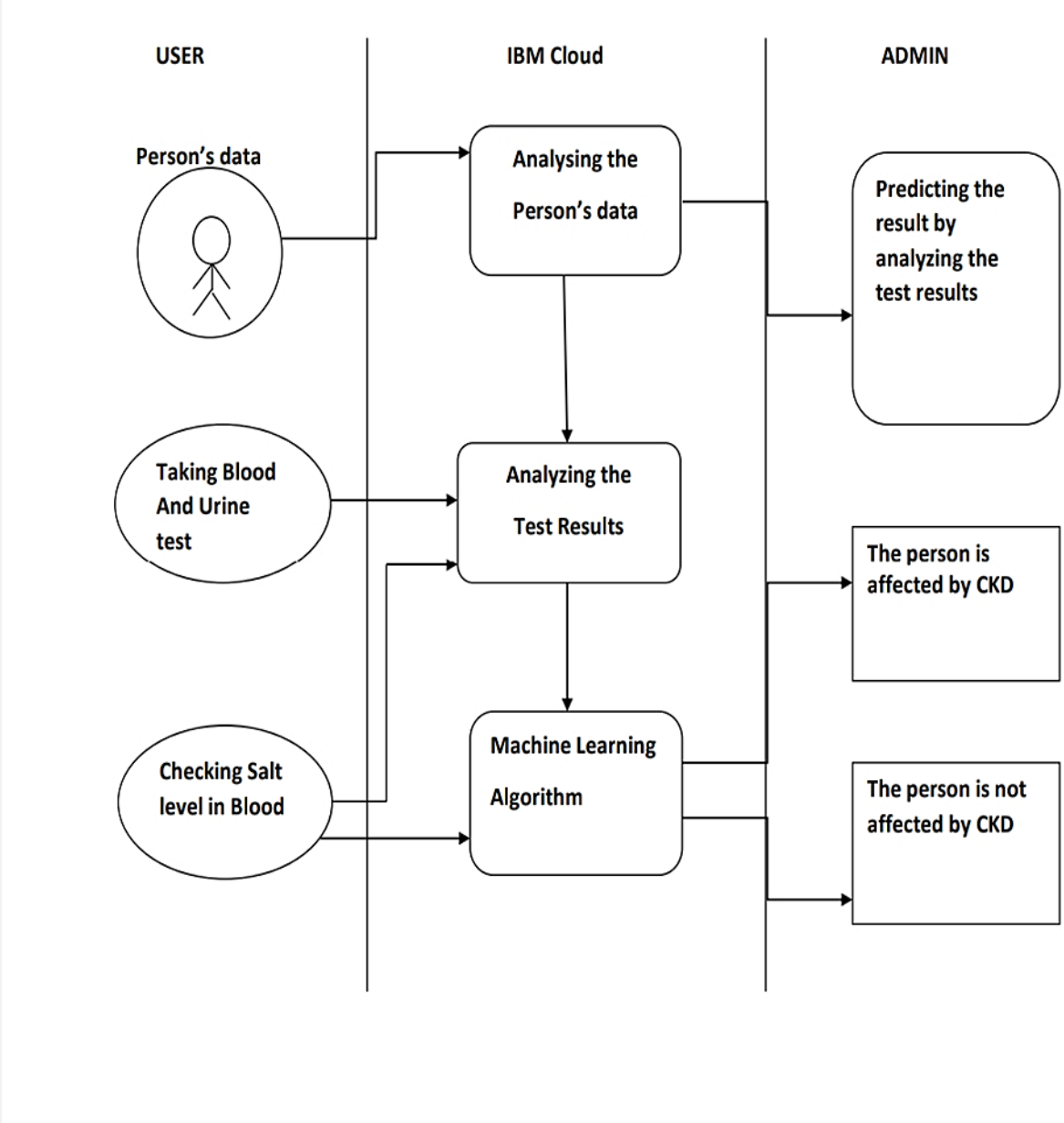
FR NO.	**Non-Functional Requirement**	**Description**
NFR - 1	Usability	Simple user-friendly interface for communication
NFR - 2	Security	Safeguard the details shared by users and maintain confidentiality
NFR - 3	Reliability	Diagnosis based on probability predicted by ML model must be reliable
NFR - 4	Performance	Reduction in overall time taken for diagnosis
NFR - 5	Availability	Available at any time to users from various places
NFR - 6	Scalability	Needs to support numerous users at once

5. PROJECT DESIGN

Data Flow Diagrams



Solution & Technical Architecture



User Stories

Create Home Page

As a user, I can collect the database of patients.

Preprocessing

As a user, I can preprocess the data from the database.

Testing

As a user, I can make the patients undergo two kinds of test (urine and blood test)

Analysis

As a user, I can analyse the test results

Prediction

As a user, I can predict CKD with the test results

Recovering Process

As a user, I can make the affected patients to take treatment

Awareness

As a user, I can advise the patients to maintain a healthy diet plan.

6. PROJECT PLANNING & SCHEDULING

Sprint Planning & Estimation

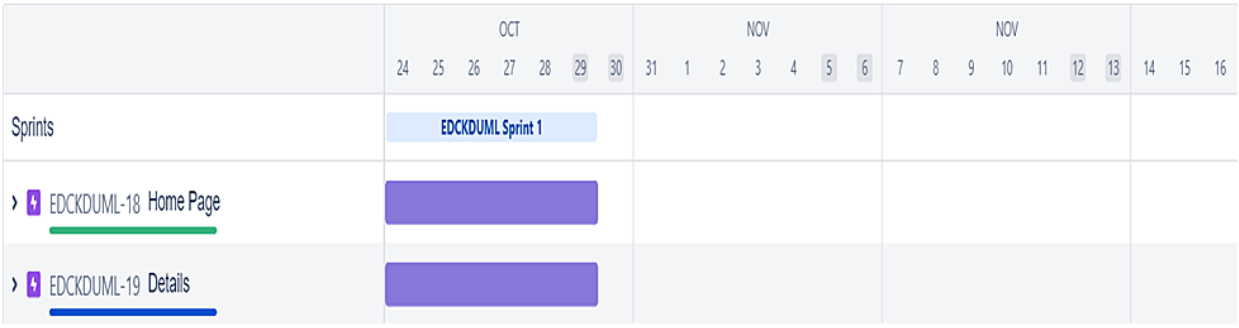
Sprint	Functional Requirement (Epic)	User Story Number	User Story /Task	Story Points	Priority
Sprint-1	Homepage	USN-1	As a user, I can create a website with homepage	5	High
Sprint-1	Details	USN-2	As a user, I can give the details about the disease	15	High
Sprint-2	Awareness	USN-3	As a user, I can create awareness about the disease	5	Low
Sprint-2		USN-4	As a user, I can give the details of the healthy diet	15	Medium
Sprint-3	Diagnosis	USN-5	As a user, I can enter the blood urea details	5	High
Sprint-3		USN-6	As a user, I can enter the glucose random.	10	High
Sprint-3		USN-7	As a user , I can select the test details of the patients	5	Medium
Sprint-4	Data analysis	USN-8	As a user, I can analyse the given data of the patient	5	High
Sprint-4	Result	USN-9	As a user, I can predict the results of the patient	10	High
Sprint-4		USN-10	As a user, I can view the results of the patient	5	High

Sprint Delivery Schedule

Sprint	Total Story Points	Duration	Sprint Start Date	Sprint End Date (Planned)	Story Points Completed (as on Planned End Date)	Sprint Release Date(Actual)
Sprint-1	20	6Days	24Oct2022	29Oct2022	20	29Oct2022
Sprint-2	20	6Days	31Oct2022	05Nov2022		
Sprint-3	20	6Days	07Nov2022	12Nov2022		
Sprint-4	20	6Days	14Nov2022	19Nov2022		

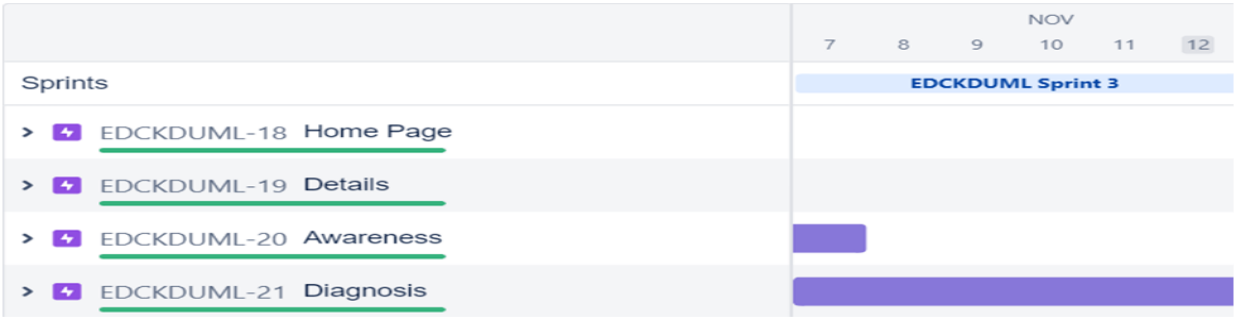
Reports from JIRA

SPRINT 1:



SPRINT 2:

SPRINT 3:



SPRINT 4:

7. CODING AND SOLUTIONING

Feature 1:

In our website, we have additionally created some more pages:

Details.html

```
<!DOCTYPE html>

<html>

<head>

<meta charset="UTF-8">

<title>About KIdney Disease</title>

<link rel="stylesheet" href="static/details.css">
```

</head>

<body bgcolor="#000">

<nav>

<label class="logo">Details about Kidney Disease</label>

<nav>

What is a Kidney Disease?

</nav>

<h1>DEFINITION:</h1>

<p>Chronic kidney disease, also called chronic kidney failure, involves a gradual loss
 of kidney function. Your kidneys filter wastes and excess fluids from your blood, which
 are then removed in your urine. Advanced chronic kidney disease can cause dangerous
 levels of fluid, electrolytes and wastes to build up in your body.

In the early stages of chronic kidney disease, you might have few signs or symptoms.
 You might not realize that you have kidney disease until the condition is advanced.

Treatment for chronic kidney disease focuses on slowing the progression of kidney damage,
 usually by controlling the cause. But, even controlling the cause might not keep
 kidney damage from progressing. Chronic kidney

disease can progress to end-stage kidney failure,
 which is fatal without artificial filtering (dialysis) or a kidney transplant.

<imgsrc="static/4.jpg" class="image">

<h1>SYMPTOMS:</h1>

Signs and symptoms of chronic kidney disease develop over time if
 kidney damage progresses slowly. Loss of kidney function can cause a
 buildup of fluid or body waste or electrolyte problems. Depending on how severe it is,
 loss of kidney function can cause:

<imgsrc="static/5.jpg">

<h1>The Symptoms are:</h1>

<p>Nausea

Vomiting

Loss of appetite

Fatigue and weakness

Sleep problems

Urinating more or less

Decreased mental sharpness

Muscle cramps

Swelling of feet and ankles

Dry, itchy skin

High blood pressure (hypertension) that's difficult to control

Shortness of breath, if fluid builds up in the lungs

Chest pain, if fluid builds up around the lining of the heart

</p

<imgsrc="static/6.jpg" class="causes">

<h1>CAUSES:</h1>

<h2>Chronic kidney disease occurs when a disease or condition impairs kidney function, causing kidney damage to worsen over several months or years.

Diseases and conditions that cause chronic kidney disease include:
</h2>

<p>

Type 1 or type 2 diabetes

High blood pressure

Glomerulonephritis (gloe-mer-u-low-nuh-FRY-tis), an inflammation of the kidney's filtering units (glomeruli)

Interstitial nephritis (in-tur-STISH-ulnuh-FRY-tis), an inflammation of the kidney's tubules and surrounding structures

Polycystic kidney disease or other inherited kidney diseases

Prolonged obstruction of the urinary tract, from conditions such as enlarged prostate, kidney stones and some cancers

Vesicoureteral (ves-ih-koe-yoo-REE-tur-ul) reflux, a condition that causes urine to back up into your kidneys

Recurrent kidney infection, also called pyelonephritis (pie-uh-low-nuh-FRY-

```
tis)<br>

</p>

</div>

</body>

</html>
```

Detail.css

```
*{
    margin: 0;
    padding: 0;
    font: 1em sans-serif;
    text-decoration: none;
    list-style: none;
    box-sizing: border-box;
}
.body{
    font-family: 'Courier New', Courier, monospace;
}
nav{
    background: grey;
```

```
    height: 30px;
    width: 100%;
}
label.logo{
    color: white;
    font-size: 35px;
    line-height: 20px;
    padding: 0 100px;
    font-weight: bolder;
    align-items: center;
    position: relative;
    left: 500px;
    top: 33px;
}
nav ul{
    font-size: 50px;
    font-weight: bold;
    font-style: italic;
    line-height: 300%;
    margin-top: 30px;
```

```
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/7.jpg);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.detail{
    margin-left: 20px;
    margin-top: 500px;
}
.details h1{
    font-size: 30px;
    color: white;
    margin-left: 20px;
```

```
    margin-top: 80px;
    font-weight: bolder;
    font-style: oblique;
}
.details p{
    font-size: 20px;
    font-weight: bold;
    line-height: 40px;
    color: white;
    margin-top: 30px;
    justify-content: space-between;
    margin-left: 30px;
}
.image{
    width: 35%;
    height: 80%;
    position: absolute;
    bottom: 0;
    right: 100px;
}
```

```
.content h1{
    font-size: 30px;
    font-weight: bolder;
    font-style: oblique;
    line-height: 80px;
    margin-left: 20px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.content p{
    font-size: 20px;
```

```
    font-weight: bold;
    line-height: 40px;
    margin-top: 0.50px;
    color: white;
    margin-left: 30px;
}
.pages{
    color: white;
}
.pages h1{
    font-size: 30px;
    font-weight: bold;
    line-height: 40px;
    margin-left: 20px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
```

```
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.pages p{
    margin-left: 30px;
    font-size: 20px;
    font-weight: bold;
    line-height: 40px;
    margin-left: 30px;
}
.causes{
    width: 50%;
    height: 500px;
    position: absolute;
    bottom: 0;
    right: 100px;
    left: 500px;
```

```
    top: 500px;
}
.c{
    font-size: 30px;
    color: white;
}
.c h1{
    font-size: 30px;
    font-weight: bolder;
    line-height: 80px;
    margin-left: 20px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
```

```
        background-position: 2000px 0;
    }
}
.c h2{
    font-size: 20px;
    font-weight: bolder;
    line-height: 50px;
    color: white;
    margin-left: 20px;
}
.c p{
    font-size: 20px;
    font-weight: bold;
    line-height: 40px;
    margin-left: 30px;
}
.d{
    width: 50%;
    height: 100px;
    margin-top: 30px;
```

```
    margin-left: 300px;
}
```

Feature 2:

Diagnosis.html

```
<!DOCTYPE html>
<html lang="en">
<head>
<title>Diagnosis of test</title>
<link rel="stylesheet" href="static/diagnosis.css">
</head>
<body bgcolor="black">
<div class="tests">
<h1>What tests do doctors use to diagnose and monitor kidney disease?</h1>
<h2>To check for kidney disease, health care providers use:</h2>
<p>A blood test that checks how well your kidneys are filtering your blood,<br>
called GFR. GFR stands for glomerular filtration rate.<br>
    A urine test to check for albumin.<br> Albumin is a protein that can pass
into the urine when the kidneys are damaged.<br>
    If you have kidney disease, your health care provider will use the same
<br>two tests to help monitor your kidney disease and make sure your treatment
plan is working.<br></p>
```

<h1>Blood test for GFR</h1>

<h2>Your health care provider will use a blood test to check your kidney
function.
 The results of the test mean the following:
</h2>

A GFR of 60 or more is in the normal range.
 Ask your health care provider when your GFR should be checked again.

A GFR of less than 60 may mean you have kidney disease.
 Talk with your health care provider about how to keep your kidney health at
 this level.

A GFR of 15 or less is called kidney failure.
 Most people below this level need dialysis or a kidney transplant.
 Talk with your health care provider about your treatment options.

<imgsrc="static/11.jpg" class="gfr">

<imgsrc="static/12.png" class="image">

</body>

</html>

Diagnosis.css

```
*{
    margin: 0;
    padding: 0;
    text-decoration: none;
    list-style: none;
    box-sizing: border-box;
}
.body{
    font-family: 'Courier New', Courier, monospace;
}
.tests h1{
    font-size: 50px;
    font-weight: bolder;
    margin-left: 0px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/13.png);
    -webkit-background-clip:text;
    background-position: 0 0;
```

```
 animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.tests{
    color: #fff;
}
.tests h2{
    font-size: 40px;
    font-weight: bold;
    margin-left: 40px;
    margin-top: 40px;
    color: #fff;
}
.tests p{
    margin-top: 50px;
```

```
    margin-bottom: 100px;
   margin-left: 40px;
    line-height: 70px;
    font-size: 40px;
    color: #fff;
}
.gfrimg{
    width: 20%;
    height: 50%;
    position: absolute;
    bottom: 0;
    right: 0px;
    top: 100px;
}
.gfr h1{
    font-size: 50px;
    top: 0px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/13.png);
```

```
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.gfr h2{
    font-size: 40px;
    font-weight: bold;
    line-height: 70px;
    top: 50px;
    color: #fff;
}
.gfr p{
    font-size: 40px;
    margin-left: 40px;
    margin-top: 40px;
```

```
    line-height: 70px;
    color: #fff;
}
.image img{
    width: 25%;
    height: 50%;
    position: absolute;
    bottom: 0;
    right: 0px;
    top: 800px;
}
```

8. TESTING

Test Cases:

Test Case ID	Feature Type	Component	Test Scenario	Pre-Requisite	Steps to execute	Test Data	Expected Result	Actual Result	Status
Home_TC_001	UI	Home Page	When user clicks on my website, the user can see the home page I've created and it is verified successfully.	HTML , CSS and Flask for integration	1.Create HTML files 2. Run using python	https://127.0.0.1:5000/	Result was not as expected	Actual result was good	Pass
Prediction_TC_002	UI	Prediction Page	Verify the input values valid or not	Flask Server, Python, anaconda prompt	1.Create HTML files 2. Run using Flask	https://127.0.0.1:5000/prediction.html	Working as expected	Satisfied as expected	Pass
Result_TC_003	UI	Result Page	Verify whether the user is able to predict	HTML,CSS, Visual Code	1.Create HTML files 2. Run using Flask	https://127.0.0.1:5000/predict			Pass

User Acceptance Testing

Defect Analysis

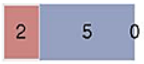
Resolution	Severity1	Severity2	Severity3	Severity4	Subtotal
By Design	10	4	2	3	20
Duplicate	1	0	3	0	4
External	2	3	0	1	6
Fixed	11	2	4	20	37
Not Reproduced	0	0	1	0	1
Skipped	0	0	1	1	2
Won't Fix	0	5	2	1	8
Totals	24	14	13	26	77

Test Case Analysis

Section	Total Cases	Not Tested	Fail	Pass
Print Engine	7	0	0	7
Client Application	51	0	0	51
Security	2	0	0	2
Outsource Shipping	3	0	0	3
Exception Reporting	9	0	0	9
Final Report Output	4	0	0	4
Version Control	2	0	0	2

9. RESULTS:

Performance metrics

Category (# of Measures)	Subcategories	Measure Validity Rating
CKD Prevention (7)	Hypertension, Diabetes	
Slowing CKD Progression (2)	Hypertension/CKD	
CKD Management (2)	Advance Care Planning, Lipid Testing	
Advanced CKD/Kidney Replacement (1)	Dialysis Access	
Dialysis Management (28)	Dialysis Access, Adequacy, Anemia, ESRD-related Complications, Transplant Referral, Advance Care Planning, Care Coordination	
Broad Measures (18)	Preventive Care, Medication Reconciliation and Safety, Advance Care Planning, Falls, Complications/Misc.	
PROMs (2)	PROMs	

10. ADVANTAGES AND DISADVANTANGES

Advantages:

- Resulted in good accuracy.
- Customers can easily predict CKD.
- This machine learning model helps us in analysing the CKD.

Disadvantages:

- At some point algorithm may fail.
- Correct prediction may not be done.

11. CONCLUSION

This work examines the ability to detect CKD using machine learning algorithms while considering the least number of tests or features. We approach this aim by applying machine learning classifier logistic regression on a big dataset of around 4000 records. We found that haemoglobin, albumin, and specific gravity have the most impact to predict the CKD.

We conclude that by using Machine Learning algorithm , we can predict whether a person is suffering from CKD or not. This effective supervised learning algorithm helps us to analyze the datasets and helps us conclude about the disease. Thus, this project helps customers to easily predict the CKD by giving their data

12. FUTURE SCOPE

This work will be considered as basement for the healthcare system for CKD

patients. Also extension to this work is that implementation of deep learning since deep learning provides high-quality performance than machine learning algorithm. In order to help in reducing the prevalence of CKD, we plan to predict if a person with CKD risk factors such as diabetes, hypertension, and family history of kidney failure will have CKD in the future or not by using appropriate dataset.

1. APPENDIX

Source code:

HTML CODING:

Index.html:

```
<!DOCTYPE html>
<html lang="en"
<head>
<title>Kidney Disease Prediction</title>
<link rel="stylesheet" type="text/css" href="static/style.css">
</head>
<body bgcolor="black">
<video src="static/video.mp4" type="video/mp4" autoplay muted
loop></video>
```

```
<section>
<header class="header">
<div class="nav-bar">
<label class="logo">KIDNEY DISEASE</label>
<div class="menu">
<ul>
<li><a href="index.html">HOME</a></li>
<li><a href="details.html">DETAILS</a></li>
<li><a href="awareness.html">AWARENESS</a></li>
<li><a href="diagnosis.html">DIAGNOSIS</a></li>
<li><a href="prediction.html">PREDICTION</a></li>
</ul>
</div>
</div>
</header>
</section>
<div class="content">
<h1>Chronic Kidney Disease Prediction</h1>
<p>Here you can know more details about Kidney Disease and Treatments</p>
```

```
</div>

</body>

</html>
```

Details.html:

```
<!DOCTYPE html>

<html>

<head>

<meta charset="UTF-8">

<title>About KIdney Disease</title>

<link rel="stylesheet" href="static/details.css">

</head>

<body bgcolor="#000">

<nav>

<label class="logo">Details about Kidney Disease</label>

<nav>

<ul>What is a Kidney Disease?</ul>

</nav>

<div class="details">

<h1>DEFINITION:</h1>
```

Chronic kidney disease, also called chronic kidney failure, involves a gradual loss
 of kidney function. Your kidneys filter wastes and excess fluids from your blood, which
 are then removed in your urine. Advanced chronic kidney disease can cause dangerous
 levels of fluid, electrolytes and wastes to build up in your body.

In the early stages of chronic kidney disease, you might have few signs or symptoms.
 You might not realize that you have kidney disease until the condition is advanced.

Treatment for chronic kidney disease focuses on slowing the progression of kidney damage,
 usually by controlling the cause. But, even controlling the cause might not keep
 kidney damage from progressing. Chronic kidney disease can progress to end-stage kidney failure,
 which is fatal without artificial filtering (dialysis) or a kidney transplant.

<imgsrc="static/4.jpg" class="image">

<h1>SYMPTOMS:</h1>

Signs and symptoms of chronic kidney disease develop over time if
kidney damage progresses slowly. Loss of kidney function can cause a
buildup of fluid or body waste or electrolyte problems. Depending on how
severe it is,
 loss of kidney function can cause:

<img src="static/5.jpg">

<h1>The Symptoms are:</h1>

Nausea

Vomiting

Loss of appetite

Fatigue and weakness

Sleep problems

Urinating more or less

Decreased mental sharpness

Muscle cramps

Swelling of feet and ankles

Dry, itchy skin

High blood pressure (hypertension) that's difficult to control

Shortness of breath, if fluid builds up in the lungs

Chest pain, if fluid builds up around the lining of the heart

<imgsrc="static/6.jpg" class="causes">

<h1>CAUSES:</h1>

<h2>Chronic kidney disease occurs when a disease or condition impairs kidney function, causing kidney damage to worsen over several months or years.
 Diseases and conditions that cause chronic kidney disease include:
</h2>

Type 1 or type 2 diabetes

High blood pressure

Glomerulonephritis (gloe-mer-u-low-nuh-FRY-tis), an inflammation of the kidney's filtering units (glomeruli)

Interstitial nephritis (in-tur-STISH-ulnuh-FRY-tis), an inflammation of the kidney's tubules and surrounding structures

Polycystic kidney disease or other inherited kidney diseases

Prolonged obstruction of the urinary tract, from conditions such as enlarged prostate, kidney stones and some cancers

Vesicoureteral (ves-ih-koe-yoo-REE-tur-ul) reflux, a condition that causes urine to back up into your kidneys

Recurrent kidney infection, also called pyelonephritis (pie-uh-low-nuh-FRY-tis)

</p>

</body>

</html>

Awareness.html:

<!DOCTYPE html>

<html lang="en">

<head>

<title>Awareness about Kidney Disease</title>

<link rel="stylesheet" href="static/awareness.css">

</head>

<body bgcolor="#000">

<imgsrc="static/10.jpg">

<h1>AWARENESS</h1>

<h3>This page gives you more awareness about the kidney disease</h3>

<p>
Public awareness and media coverage of kidney disease is relatively low
compared to other conditions, such as heart disease or cancer.

 Many people we spoke to knew that the kidneys are a vital organ,
 that they act as a filter to remove waste products from the blood
 stream and that they are somehow connected to ‘the waterworks’.

 However, the term ‘kidney disease’ may make people think of kidney
 failure, transplants and dialysis.

 As Jim put it, “I knew you could get kidney failure and things like that,
 but I didn’t realise you could just get something small wrong with
them”.

 Some people thought there should be more advertising of
 kidney disease,

especially what causes it, why kidneys are important
 for the body to stay healthy and what people can do to keep their kidneys healthy.

The urgency of making patients aware of CKD could be questioned on the grounds
 that many patients die before they progress to a more severe stage
 of CKD and many are already being treated for diabetes and/or hypertension.28,29

 However, there are still compelling reasons why patients would benefit from awareness of
 their CKD.

 First, they could be made aware of medication exposures that could influence progression, including over-the-counter
 nonsteroidal anti-inflammatory agents and contrast agents used in imaging tests.

 In addition, there is evidence that appropriate early treatment (with medications and
hypertension control) could slow progression of CKD.29

Also, patients should be aware that current and future medications could require dose adjustment
 in the setting of CKD.

 Many conditions, such as heart disease and cognitive decline, may increase
 with severity of CKD, and awareness of disease may motivate patients to adopt preventive
 strategies for these conditions.

 Awareness might also make patients more vigilant with adherence to dietary recommendations for comorbid
hypertension and diabetes, including lower salt, sugar, and fat intake, as well as
other

lifestyle changes.

<imgsrc="static/9.jpg">

<h1>HEALTHY FOODS</h1>

<h2> Tips for making healthy food choices:</h2>

Cook with a mix of spices instead of salt.

Choose veggie toppings such as spinach, broccoli, and peppers for your pizza.

Try baking or broiling meat, chicken, and fish instead of frying.

Serve foods without gravy or added fats.

Try to choose foods with little or no added sugar.

Gradually work your way down from whole milk to 2 percent milk until you’re drinking and cooking with fat-free (skim) or low-fat milk and milk products.

Eat foods made from whole grains—such as whole wheat, brown rice, oats, and whole-grain corn—every day.
Use whole-grain bread for toast

and sandwiches; substitute brown rice for white rice for home-cooked meals and when dining out.

Read food labels. Choose foods low in saturated fats, trans fats, cholesterol, salt (sodium), and added sugars.

Slow down at snack time. Eating a bag of low-fat popcorn takes longer than eating a slice of cake.
 Peel and eat an orange instead of drinking orange juice.

Try keeping a written record of what you eat for a week.
 It can help you see when you tend to overeat or eat foods high in fat or calories.</p>

</body>

</html>

Diagnosis.html:

<!DOCTYPE html>

<html lang="en">

<head>

<title>Diagnosis of test</title>

<link rel="stylesheet" href="static/diagnosis.css">

</head>

<body bgcolor="black">

<h1>What tests do doctors use to diagnose and monitor kidney disease?</h1>

<h2>To check for kidney disease, health care providers use:</h2>

A blood test that checks how well your kidneys are filtering your blood,
 called GFR. GFR stands for glomerular filtration rate.

A urine test to check for albumin.
 Albumin is a protein that can pass into the urine when the kidneys are damaged.

If you have kidney disease, your health care provider will use the same
two tests to help monitor your kidney disease and make sure your treatment plan is working.

<h1>Blood test for GFR</h1>

<h2>Your health care provider will use a blood test to check your kidney
function.
 The results of the test mean the following:
</h2>

A GFR of 60 or more is in the normal range.
 Ask your health care provider when your GFR should be checked again.

A GFR of less than 60 may mean you have kidney disease.
 Talk with your health care provider about how to keep your kidney health at
 this level.

A GFR of 15 or less is called kidney failure.
 Most people below this level need dialysis or a kidney transplant.
 Talk with your health care provider about your treatment options.</p>

<imgsrc="static/11.jpg" class="gfr">

<imgsrc="static/12.png" class="image">

</body>

</html>

Prediction.html:

<!doctype html>

<html lang="en">

<head>

<!-- Required meta tags -->

```
<meta charset="utf-8">
<meta name="viewport" content="width=device-width, initial-scale=1">
<!-- Bootstrap CSS -->
<link rel="stylesheet" href="static/prediction.css">
<link href="https://cdn.jsdelivr.net/npm/bootstrap@5.0.0-
beta3/dist/css/bootstrap.min.css" rel="stylesheet" integrity="sha384-
eOJMYsd53ii+scO/bJGFsiCZc+5NDVN2yr8+0RDqr0Ql0h+rP48ckxlpbzKgwr
a6" crossorigin="anonymous">
<link href="https://unpkg.com/tailwindcss@^2/dist/tailwind.min.css"
rel="stylesheet">
<title>prediction</title>
</head>
<body>
<section class="text-white-900 body-font">
<div class="container  py-5 mx-auto">
<div class="flex flex-col text-center w-full mb-20">
<h1 class="sm:text-6xl text-5xl font-medium title-font mb-4 text-white-
1000">Kidney Disese</h1>
<p class="sm:text-3xl mx-auto leading-relaxed text-base">Fill the form for
```

```
prediction</p>

<h2>{{prediction_text}}</h2>

</div>

<div>

</div>

<a class="btnbtn-primary" href="./" role="button">Back</a>

<form action='/predict' method='POST'>

<div class="mb-3">

<label for="exampleFormControlInput1" class="form-label">Enter blood

urea</label>

<input type="text" class="form-control"  name="blood_urea"

placeholder="blood_urea">

</div>

<div class="mb-3">

<label for="exampleFormControlInput1" class="form-label">Enter blood

glucose</label>

<input type="text" class="form-control" name="blood glucose random"

placeholder="blood glucose random">

</div>
```

```
<div class="mb-3">
<label for="exampleFormControlInput1" class="form-label">Enter
coronary_artery_disease</label>
<select class="form-select"  name="coronary_artery_disease" aria-
label="Default select example">
<option selected>-- select Enter coronary_artery_disease --</option>
<option value="0">0</option>
<option value="1">1</option>
</select>
</div>
<div class="mb-3">
<label for="exampleFormControlInput1" class="form-label">Enter
anemia</label>
<select class="form-select"  name="anemia" aria-label="Default select
example">
<option selected>-- select Enter anemia --</option>
<option value="0">0</option>
<option value="1">1</option>
</select>
```

```
</div>
<div class="mb-3">
<label for="exampleFormControlInput1" class="form-label">Enter
pus_cell</label>
<select class="form-select"  name="pus_cell" aria-label="Default select
example">
<option selected>-- select Enter pus_cell --</option>
<option value="0">0</option>
<option value="1">1</option>
</select>
</div>
<div class="mb-3">
<label for="exampleFormControlInput1" class="form-label">Enter
red_blood_cells</label>
<select class="form-select"  name="red_blood_cells" aria-label="Default select
example">
<option selected>-- select Enter red_blood_cells --</option>
<option value="0">0</option>
<option value="1">1</option>
```

```
</select>
</div>
<div class="mb-3">
<label for="exampleFormControlInput1" class="form-label">Enter
diabetesmellitus</label>
<select class="form-select"  name="diabetesmellitus" aria-label="Default select
example">
<option selected>-- select Enter diabetesmellitus --</option>
<option value="0">0</option>
<option value="1">1</option>
<option value="2">2</option>
</select>
</div>
<div class="mb-3">
<label for="exampleFormControlInput1" class="form-label">Enter
pedal_edema</label>
<select class="form-select"  name="pedal_edema" aria-label="Default select
example">
<option selected>-- select Enter pedal_edema --</option>
```

```
<option value="0">0</option>

<option value="1">1</option>

</select>

</div>

<button type="submit" class="btnbtn-primary">Predict</button>

</form>

</div>

</section>

</body>

</html>
```

Success.html:

```
<!DOCTYPE html>

<html lang="en">

<head>

<meta charset="UTF-8">

<meta http-equiv="X-UA-Compatible" content="IE=edge">

<meta name="viewport" content="width=device-width, initial-scale=1.0">

<title>Result Page</title>
```

```
<link rel="stylesheet" type="text/css" href="static/success.css">
</head>
<body>
<div class="image">
<h1>You are a healthy person!!!</h1>
<imgsrc="static/15.jpg">
</div>
</body>
</html>
```

Failure.html:

```
<!DOCTYPE html>
<html lang="en">
<head>
<meta charset="UTF-8">
<meta http-equiv="X-UA-Compatible" content="IE=edge">
<meta name="viewport" content="width=device-width, initial-scale=1.0">
<title>Result Page</title>
<link rel="stylesheet" type="text/css" href="static/failure.css">
```

```
</head>

<body>

<div class="image">

<h1>OOPS You have kidney disease!!!</h1>

<imgsrc="static/16.png">

</div>

</body>

</html>
```

CSS CODING:

Style.css:

```
*{

    margin: 0;

    padding: 0;

    box-sizing: border-box;

}

a,a:hover{

    text-decoration: none;

}
```

```
ul{
    list-style: none;
}
.navbar{
    background: #000;
    height: 80px;
    width: 100%;
    font-size: 20px;
}
.header .nav-bar{
    width: 90%;
    height: 80px;
    line-height: 80px;
    margin: auto;
    color: white;
    display: flex;
justify-content: space-between;
    font-size: 30;
}
```

```
.header .nav-bar .logo{
    z-index: 1;
    position: relative;
    left: 25px;
    font-size: 20px;
    color: transparent;
    -webkit-text-stroke: 1px #fff;
    background: url(static/1.jpg);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.header .nav-bar .logo a{
    font-size: 20px;
```

```
    font-weight: bold;
    padding: 10px;
    line-height: 10px;
    text-transform: uppercase;
}
.header .nav-bar .menu{
    z-index: 2;
}
.header .nav-bar .menu ul li{
    display: inline-block;
    line-height: 20px;
    margin-right: 50px;
}
.header .nav-bar .menu ul li a{
    font-size: 20px;
    font-weight: bold;
    padding: 5px;
    text-transform: capitalize;
    transition: all 0.5s;
```

```
    opacity: 0.8;
    color: black;
    -webkit-text-stroke: #fff;
}
.header .nav-bar .menu ul li a.active{
    background-color: #fff;
    color: #000;
    font-weight: bold;
}
.header .nav-bar .menu ul li a:hover{
    opacity: 1;
    background-color: #fff;
    color: #000;
}
.content{
    width: 100%;
    position: absolute;
    top: 50%;
    transform: translateY(-50%);
```

```
    text-align: center;
    color: black;
}
.content h1{
    font-size: 100px;
    margin-top: 80px;
    font-style: italic;
    color: white;
    font-weight: bolder;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/3.jpg);
    -webkit-background-clip: text;
}
.content p{
    margin: 20px auto;
    font-weight: 200;
    line-height: 25px;
    font-size: 40px;
```

```
    color: #fff;
}
video{
    position: fixed;
    right: 0;
    bottom: 0;
    width: 100%;
    height: 100vh;
}
```

Details.css:

```
*{
    margin: 0;
    padding: 0;
    text-decoration: none;
    list-style: none;
    box-sizing: border-box;
}
.body{
    font-family: 'Courier New', Courier, monospace;
```

```
}
.tests h1{
    font-size: 50px;
    font-weight: bolder;
    margin-left: 0px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/13.png);
    -webkit-background-clip:text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.tests{
    color: #fff;
```

```
}
.tests h2{
    font-size: 40px;
    font-weight: bold;
    margin-left: 40px;
    margin-top: 40px;
    color: #fff;
}
.tests p{
    margin-top: 50px;
    margin-bottom: 100px;
    margin-left: 40px;
    line-height: 70px;
    font-size: 40px;
    color: #fff;
}
.gfrimg{
    width: 20%;
    height: 50%;
```

```
    position: absolute;
    bottom: 0;
    right: 0px;
    top: 100px;
}
.gfr h1{
    font-size: 50px;
    top: 0px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/13.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
```

```
}
.gfr h2{
    font-size: 40px;
    font-weight: bold;
    line-height: 70px;
    top: 50px;
    color: #fff;
}
.gfr p{
    font-size: 40px;
    margin-left: 40px;
    margin-top: 40px;
    line-height: 70px;
    color: #fff;
}
.imageimg{
    width: 25%;
    height: 50%;
    position: absolute;
```

```
    bottom:   0;
    right:   0px;
    top: 800px;
}
```

Awareness.css:

```
*{
    padding: 0;
    margin: 0;
    box-sizing: border-box;
    font-family: 'Josefin Sans, sans-serif';
}
.header{
    margin-left: 30px;
    margin-top: 30px;
    color: white;
}
.header img{
    width: 40%;
```

```
    height: 100%;
    position: absolute;
    bottom: 0;
    right: 0px;
    top: 0px;
}
.content h1{
    font-size: 40px;
    font-weight: bolder;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
```

```
    }
}
.content h3{
    font-size: 30px;
    font-weight: bold;
    margin-top: 20px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.content p{
```

```
    margin-top: 2px;
    margin-bottom: 100px;
    margin-left: 40px;
    line-height: 40px;
    font-size: 30px;
    color: #fff;
}
.health{
    margin-right: 0;
    margin-top: 100px;
}
.health img{
    margin-right: 0px;
    float: right;
}
.foods{
    margin-left: 0px;
    margin-top: 30px;
    color: white;
```

```
}
.foods h1{
    margin-left: 0px;
    margin-top: 50px;
    font-size: 40px;
    font-weight: bolder;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
.foods h2{
    font-size: 30px;
    font-weight: bold;
    margin-top: 20px;
    color: white;
    -webkit-text-stroke: 1px #fff;
```

```
    background: url(static/8.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.foods p{
    margin-top: 30px;
    margin-bottom: 100px;
    margin-left: 40px;
    line-height: 60px;
    font-size: 30px;
    color: #fff;
}
```

Diagnosis.css:

```
*{
    margin: 0;
    padding: 0;
    text-decoration: none;
    list-style: none;
    box-sizing: border-box;
}
.body{
    font-family: 'Courier New', Courier, monospace;
}
.tests h1{
    font-size: 50px;
    font-weight: bolder;
    margin-left: 0px;
    color: white;
    -webkit-text-stroke: 1px #fff;
    background: url(static/13.png);
    -webkit-background-clip:text;
```

```
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.tests{
    color: #fff;
}
.tests h2{
    font-size: 40px;
    font-weight: bold;
    margin-left: 40px;
    margin-top: 40px;
    color: #fff;
}
.tests p{
```

```
    margin-top: 50px;
    margin-bottom: 100px;
    margin-left: 40px;
    line-height: 70px;
    font-size: 40px;
    color: #fff;
}
.gfrimg{
    width: 20%;
    height: 50%;
    position: absolute;
    bottom: 0;
    right: 0px;
    top: 100px;
}
.gfr h1{
    font-size: 50px;
    top: 0px;
    color: white;
```

```
    -webkit-text-stroke: 1px #fff;
    background: url(static/13.png);
    -webkit-background-clip: text;
    background-position: 0 0;
    animation: back 20s linear infinite;
}
@keyframes back{
    100%{
        background-position: 2000px 0;
    }
}
.gfr h2{
    font-size: 40px;
    font-weight: bold;
    line-height: 70px;
    top: 50px;
    color: #fff;
}
.gfr p{
```

```
    font-size: 40px;
    margin-left: 40px;
    margin-top: 40px;
    line-height: 70px;
    color: #fff;
}
.imageimg{
    width: 25%;
    height: 50%;
    position: absolute;
    bottom: 0;
    right: 0px;
    top: 800px;
}
```

Flask Integrate With Scoring End Pints:

```
import pickle
import numpy as np
import pandas as pd
```

```
from flask import Flask, render_template, request
app=Flask(__name__, template_folder='templates')
model=pickle.load(open('CKD.pkl','rb'))

@app.route('/')
def index():
    return render_template('index.html')

@app.route('/details.html')
def details():
    return render_template('details.html')

@app.route('/awareness.html')
def awareness():
    return render_template('awareness.html')

@app.route('/diagnosis.html')
def diagnosis():
    return render_template('diagnosis.html')
```

```
@app.route('/prediction.html')
def prediction():
    return render_template('prediction.html')


@app.route('/predict',methods=['POST','GET'])
def predict():
input_features=[float(x) for x in request.form.values()]
    features_value=[np.array(input_features)]
features_name=['blood_urea','blood glucose
random','coronary_artery_disease','anemia','pus_cell','red_blood_cells','diabetes
mellitus','pedal_edema']
df=pd.DataFrame(features_value, columns=features_name)
output=model.predict(df)
 if output == 1:
        return render_template('success.html')
    else:
        return render_template('failure.html')
```

if__name__== '__main__':

app.run(debug=True)

NOTEBOOK .ipynb

Import Dataframe:

```
In [87]: import os, types
         import pandas as pd
         from botocore.client import Config
         import ibm_boto3

         def __iter__(self): return 0

         # @hidden_cell
         # The following code accesses a file in your IBM Cloud Object Storage. It includes your credentials.
         # You might want to remove those credentials before you share the notebook.
         cos_client = ibm_boto3.client(service_name='s3',
             ibm_api_key_id='GenImWOC7pmDNtYSNujVb89Ra_uBfV7HW1eYmqCch1cr',
             ibm_auth_endpoint="https://iam.cloud.ibm.com/oidc/token",
             config=Config(signature_version='oauth'),
             endpoint_url='https://s3.private.us.cloud-object-storage.appdomain.cloud')

         bucket = 'ckd-donotdelete-pr-llwzpw51re9nxp'
         object_key = 'chronickidneydisease.csv'

         body = cos_client.get_object(Bucket=bucket,Key=object_key)['Body']
         # add missing __iter__ method, so pandas accepts body as file-like object
         if not hasattr(body, "__iter__"): body.__iter__ = types.MethodType( __iter__, body )
```

Import Libraries:

```
In [131]: import numpy as np
          import pandas as pd
          import matplotlib.pyplot as plt
          import seaborn as sns
          from sklearn import preprocessing
          import scipy.stats as stats
          from sklearn.model_selection import train_test_split
          from collections import Counter
          from sklearn.metrics import accuracy_score,confusion_matrix,classification_report
          from sklearn import metrics
          from sklearn.ensemble import RandomForestClassifier
          from sklearn.svm import SVC
          from sklearn.linear_model import LogisticRegression
          from sklearn.naive_bayes import GaussianNB
          from sklearn.neighbors import KNeighborsClassifier
          import joblib
```

Load The Dataset:

LOAD DATASET

```
In [15]: df=pd.read_csv("chronickidneydisease.csv")
         pd.set_option("display.max_column",None)
         df
```

```
In [38]: df.shape
Out[38]: (400, 27)
```

```
In [39]: df.info
```

```
In [42]: df.sod.unique()
Out[42]: array([139. , 136. , 140. , 111. , 142. , 104. , 147. , 131. , 114. ,
                124. , 138. , 135. , 130. , 141. , 145. ,   4.5, 128. , 129. ,
                144. , 132. , 133. , 150. , 134. , 125. , 137. , 163. , 143. ,
                127. , 146. , 126. , 122. , 115. , 113. , 120. ])
```

```
In [43]: df.pcc.value_counts()
Out[43]: notpresent    354
         present        42
         Name: pcc, dtype: int64
```

```
In [18]: print(numerical_feature)
         ['id', 'age', 'bp', 'sg', 'al', 'su', 'bgr', 'bu', 'sc', 'sod', 'pot', 'hemo']
```

```
In [104]: df.isnull().any()
Out[104]: id               False
```

Visualization:

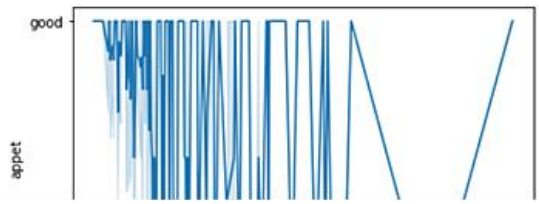
VISUALIZATION

```
In [44]: sns.displot(df.age)
Out[44]: <seaborn.axisgrid.FacetGrid at 0x28be2b13d30>
```

In [45]:
```
sns.lineplot(df.bu,df.appet)
```

```
C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn\_decorators.py:36: FutureWarning: Pass the followi
ng variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `da
ta`, and passing other arguments without an explicit keyword will result in an error or misinterpreta
tion.
  warnings.warn(
```

Out[45]:
```
<AxesSubplot:xlabel='bu', ylabel='appet'>
```



In [48]:
```
sns.scatterplot(df.sg,df.bp)
```

```
C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn\_decorators.py:36: FutureWarning: Pass the followi
ng variables as keyword args: x, y. From version 0.12, the only valid positional argument will be `da
ta`, and passing other arguments without an explicit keyword will result in an error or misinterpreta
tion.
  warnings.warn(
```

Out[48]:
```
<AxesSubplot:xlabel='sg', ylabel='bp'>
```

In [55]:
```
df.hist(figsize=(8,8))
```

Out[55]:
```
array([[<AxesSubplot:title={'center':'id'}>,
```

Outlier Detection:

OUTLIER DETECTION ¶

In [57]:
```
sns.boxplot(df.pot)
```

```
C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn\_decorators.py:36: FutureWarning: Pass the followi
ng variable as a keyword arg: x. From version 0.12, the only valid positional argument will be `data
`, and passing other arguments without an explicit keyword will result in an error or misinterpretati
on.
  warnings.warn(
```

Out[57]:
```
<AxesSubplot:xlabel='pot'>
```

Outlier Using IQR:

OUTLIER REMOVAL USING IQR

```
In [58]: q1=df.hemo.quantile(0.25)  #(Q1)
         q3=df.hemo.quantile(0.75)  #(Q3)

In [65]: IQR=q3-q1

In [66]: upper_limit= q3 + 1.5*IQR

In [68]: uppper_limit= q3 + 1.5*IQR
         lower_limit=q1 - 1.5*IQR

In [69]: sns.boxplot(df.bu)
```

Outlier removal with percentile:

OUTLIER REMOVAL WITH PERCENTILE

```
In [71]: p99=df.hemo.quantile(0.99)
         p99
```

Z-Score:

Z-SCORE

```
In [73]: from scipy import stats

In [76]: hemo_zscore=stats.zscore(df.hemo)

In [77]: hemo_zscore
```

Encoding Techniques:

ENCODING TECHNIQUES

1. One hot encoding

```
In [78]: from sklearn.preprocessing import LabelEncoder

In [79]: le=LabelEncoder()

In [80]: df.age=le.fit_transform(df.age)
         df.bp=le.fit_transform(df.bp)
```

2. One hot encoding

```
In [83]: df_main=pd.get_dummies(df,columns=['hemo'])
         df_main.head()
```

X and Y Split:

X and Y Split

```
In [116]: # dependent variable

          y=df_main['pcc']
          y
```

```
In [92]: # independent variable

         x=df_main.drop(columns=['pcc'],axis=1)
         x.head()
```

Linear Regression:

LINEAR REGRESSION

1.Simple Linear Regression

In [120]:
```
sns.displot(df.wc)
```

```
C:\Users\ELCOT\anaconda3\lib\site-packages\seaborn\axisgrid.py:88: UserWarning: Glyph 9 (       ) missing from current font.
  self._figure.tight_layout(*args, **kwargs)
```

In [141]:
```
x=df.drop(columns=['hemo'],axis=1)
x
```

Multi Linear Regression:

2. Multi Linear Regression

In [126]:
```
from sklearn.linear_model import LinearRegression
```

In [151]:
```
from sklearn.preprocessing import LabelEncoder
le=LabelEncoder()
```

In [153]:
```
df.State=le.fit_transform(df.hemo)
df
```

In [155]:
```
df.corr().age.sort_values(ascending=False)
```

X ana Y split

```
In [157]: X=df.drop(columns=['age'],axis=1)
          X.head()  # independent variables
```

```
In [158]: y= df.age
          y.head()
```

Polynomial Regression:

Polynomial Regression

```
In [163]: x=df.iloc[:,1:2]
          x
```

```
In [166]: from sklearn.linear_model import LinearRegression
          lr=LinearRegression()
```

```
In [167]: lr.fit(xp,y)
```


Convert Normal Feature to Polynomial feature

```
In [164]: from sklearn.preprocessing import PolynomialFeatures
          pr=PolynomialFeatures(degree=4)
```

```
In [165]: xp=pr.fit_transform(x)
          xp
```

```
In [168]: # checking the prediction on a random value

          id=lr.predict(pr.fit_transform([[6]]))
          id
```

Logistic Regression:

LOGISTIC REGRESSION

```
In [170]: from sklearn.preprocessing import LabelEncoder

          le=LabelEncoder()

          df.age=le.fit_transform(df.age)

          df.head()
```

```
In [171]: df.id.value_counts()
```

```
In [237]: sns.displot(df.age)
```

Model Building:

```
In [243]: from sklearn.preprocessing import MinMaxScaler
          scale = MinMaxScaler()
```

Model Building

```
In [242]: ## Logistic Regression

          from sklearn.linear_model import LogisticRegression
```

```
In [244]: model = LogisticRegression()
```

```
In [245]: model.fit(x_train,y_train)
```

Evaluating Model:

Evaluating the Model

In [182]:
```
# accuracy score

from sklearn.metrics import accuracy_score,classification_report,confusion_matrix,roc_auc_score,roc_curve
```

In [247]:
```
accuracy_score(y_test,pred_test) # test accuracy
```

Out[247]: 0.06060606060606061

In [248]:
```
# Confusion matrix

pd.crosstab(y_test,pred_test)
```

In [249]:
```
# Classification report

print(classification_report(y_test,pred_test))
```

In [186]:
```
# Precision

#  TP/(TP+FP)

24/30
```

In [187]:
```
# Recall

# TP /(TP+FN)

24/32
```

In [188]:
```
# F1 score

# 2*precision*Recall / (precision+Recall)

2*0.8*0.75/(0.8+0.75)
```

In [229]:
```
# performs the split

from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(x,y)
```

In [223]:
```
# Display the shape

print(x_train.shape)
print(x_test.shape)
print(y_train.shape)
print(y_test.shape)
```

```
(297, 1)
(99, 1)
(297,)
(99,)
```

In [231]:
```
x_train.shape
```

Out[231]: (297, 1)

In [232]:
```
x_test.shape
```

Out[232]: (99, 1)

In [252]:
```
# ROC - AUC Score

probability = model.predict_proba(x_test)[:,1]
probability
```


Ridge and Lasso Regression:

RIDGE AND LASSO REGRESSION

In [256]:
```
# l1 Regularization technique- Lasso Regression
# l2 Regularization technique- Ridge Regression
```

In [259]:
```
from sklearn.linear_model import Ridge
from sklearn.linear_model import Lasso
```

In [260]:
```
r=Ridge()
l=Lasso()
```

In [261]:
```
r.fit(x_train,y_train)
```

Out[261]: Ridge()

In [262]:
```
l.fit(x_train,y_train)
```

Out[262]: Lasso()

In [265]:
```
pred1=r.predict(x_test)
pred1
```

In [266]:
```
pred2=l.predict(x_test)
pred2
```

Evaluation Metrics for Regression:

1.Evaluation metrics for Regression problem

In [267]:
```
# R-Square
# testing accuracy for both model

print(metrics.r2_score(y_test,pred1))
print(metrics.r2_score(y_test,pred2))
```

```
0.9999999998368482
0.9999856081783125
```

In [268]:
```
profit=pd.DataFrame({'Actual':y_test,'ridge_pred':pred1,'lasso_pred':pred2})
profit.head(10)
```

In [269]:
```
## MSE(Mean square error)

print(metrics.mean_squared_error(y_test,pred1))
print(metrics.mean_squared_error(y_test,pred2))
```

```
3.7522833790478166e-08
0.0033099339693025715
```

Decision Tree Classifier:

DECISION TREE CLASSIFIER

In [270]:
```
# X and y split

X=df.iloc[:,:-1]
X.head()
```

1. Model Building

```
In [277]: from sklearn.tree import DecisionTreeClassifier

          model = DecisionTreeClassifier(max_depth=4,splitter='best',criterion='entropy')

In [278]: model.fit(x_train,y_train)

Out[278]: DecisionTreeClassifier(criterion='entropy', max_depth=4)

In [279]: y_predict= model.predict(x_test)
          y_predict
```

Random Forest Classifier:

RANDOM FOREST CLASSIFIER

1. Model Building

```
In [282]: from sklearn.ensemble import RandomForestClassifier
          model = RandomForestClassifier(n_estimators=10,criterion='entropy')

In [283]: model.fit(x_train,y_train)

Out[283]: RandomForestClassifier(criterion='entropy', n_estimators=10)

In [284]: y_predict = model.predict(x_test)

In [285]: y_predict_train = model.predict(x_train)

In [286]: from sklearn.metrics import accuracy_score,confusion_matrix,classification_report

In [287]: print('Training accuracy: ',accuracy_score(y_train,y_predict_train))
          print('Testing accuracy: ',accuracy_score(y_test,y_predict))

          Training accuracy:  0.98989898989899
          Testing accuracy:  0.9797979797979798
```


Naive Baye's Classifier:

NAIVE BAYE'S CLASSIFIER

```
In [288]: # using for classification problem.
          # Naive baye's based on Baye's theorem fro the classification.

In [289]: from sklearn.preprocessing import MinMaxScaler
          scale = MinMaxScaler()

In [394]: x_scaled = pd.DataFrame(scale.fit_transform(x),columns= x.columns)
          x_scaled.head()
```

1. Model Building

In [294]:
```
# model building
from sklearn.naive_bayes import GaussianNB
model =GaussianNB()
```

In [295]:
```
model.fit(x_train,y_train)
```

Out[295]:
```
GaussianNB()
```

In [296]:
```
# evaluating the model
y_pred = model.predict(x_test)
```

In [297]:
```
from sklearn.metrics import accuracy_score,confusion_matrix,classification_report
```

In [298]:
```
accuracy_score(y_test,y_pred)
```

Out[298]:
```
0.9797979797979798
```

In [299]:
```
print(classification_report(y_test,y_pred))
```

Unsupervised K-Means:

UNSUPERVISED K-MEANS

In [300]:
```
from sklearn import cluster
```

In [332]:
```
error =[]
for i in range(11,1):
    kmeans=cluster.KMeans(n_clusters=i,init="k-means++",random_state=3)
    kmeans.fit(new_df)
    error.append(kmeans.inertia_)
```

In [333]:
```
error
```

Out[333]:
```
[]
```

In [334]:
```
import matplotlib.pyplot as plt
plt.plot(range(11,1),error)
plt.title('Elbow method')
plt.xlabel('no of clus')
plt.ylabel('error')
plt.show()
```

Model Building

```
In [340]: # model building
          from sklearn.neighbors import KNeighborsClassifier
          model = KNeighborsClassifier()

In [341]: model.fit(x_train,y_train)

Out[341]: KNeighborsClassifier()

In [395]: y_pred=model.predict(x_test)

In [396]: y_pred1=model.predict(x_train)
```

Model Evaluation:

Model Evaluation

```
In [345]: # model evaluation
          from sklearn.metrics import accuracy_score,confusion_matrix,classification_report

In [385]: # testing with a random observation

          model.predict([[1.1]])
```

IBM CODING:

```
import numpy as np

import pandas as pd

from flask import Flask, render_template, request

import requests

import pickle


# NOTE: you must manually set API_KEY below using information retrieved
```

from your IBM Cloud account.

```
API_KEY = "QXCDaXxG__YG-
rHuYkTQwfeatR70SW8MTP9Lr29ORWNB"
token_response = requests.post('https://iam.cloud.ibm.com/identity/token',
data={"apikey":API_KEY, "grant_type": 'urn:ibm:params:oauth:grant-
type:apikey'})
mltoken = token_response.json()["access_token"]

header = {'Content-Type': 'application/json', 'Authorization': 'Bearer ' +
mltoken}

model=pickle.load(open('CKD.pkl','rb'))

# NOTE: manually define and pass the array(s) of values to be scored in the
next line

app=Flask(__name__, template_folder='templates')

@app.route('/')
```

```
def index():
    return render_template('index.html')

@app.route('/details.html')
def details():
    return render_template('details.html')

@app.route('/awareness.html')
def awareness():
    return render_template('awareness.html')

@app.route('/diagnosis.html')
def diagnosis():
    return render_template('diagnosis.html')


@app.route('/prediction.html')
def prediction():
    return render_template('prediction.html')
```

```
@app.route('/predict',methods=['POST','GET'])
def predict():

    input_features=[float(x) for x in request.form.values()]
    features_value=[np.array(input_features)]

    features_name=['blood_urea','blood glucose
random','coronary_artery_disease','anemia','pus_cell','red_blood_cells','diabetes
mellitus','pedal_edema']

    df=pd.DataFrame(features_value, columns=features_name)

    output=model.predict(df)

    #showing the prediction resultsin a UI# showing the prediction results in UI
    if output == 1:
        return render_template('success.html')
    else:
```

```
        return render_template('failure.html')

    payload_scoring = {"input_data":  [{"fields":
[['age','bp','sg','al','su','rbc','pc','pcc','ba','bgr','bu','sc','sod','pot','hemo','pcv','wc','rc
','htn','dm','cad','appet','pe','ane']], "values": features_name}]}
    response_scoring = requests.post('https://us-
south.ml.cloud.ibm.com/ml/v4/deployments/e0e99a97-2d3d-482b-bedf-
d1a5f97dfdc4/predictions?version=2022-11-12', json=payload_scoring,
    headers={'Authorization': 'Bearer ' + mltoken})
    print("Scoring response")
    print(response_scoring.json())
if__name__== '__main__':
    app.run(debug=True)
```

IBM DEPLOYMENT:

Building a Machine Learning model

In [34]:
```
from sklearn.linear_model import LogisticRegression
lgr=LogisticRegression()
lgr.fit(x_train,y_train)
```

```
C:\Users\DX\anaconda3\lib\site-packages\sklearn\utils\validation.py:993: DataConversionWarning: A column-vector y was passed wh
en a 1d array was expected. Please change the shape of y to (n_samples, ), for example using ravel().
  y = column_or_1d(y, warn=True)
```

Out[34]:
```
LogisticRegression()
```

Predicting our output with the model which we build

In [35]:
```
y_pred=lgr.predict(x_test)
```

In [37]:
```
y_pred1=lgr.predict([[129,99,1,0,0,1,0,1]])
print(y_pred)
c(y_pred)
```

```
[0 0 0 0 1 0 0 0 1 0 0 0 1 1 0 0 0 1 1 0 1 1 0 1 0 1 0 0 1 0 0 1 0 0 0 0 1
 0 0 1 0 1 0 0 0 1 0 1 1 1 0 0 0 1 0 1 0 1 1 1 0 1 1 0 0 0 0 1 0 1 1 0 0 1
 0 1 0 1 1 0]
```

Out[37]:
```
Counter({0: 47, 1: 33})
```

In [39]:
```
accuracy_score(y_test,y_pred)
```

Out[39]:
```
0.9125
```


Confusion Matrix of our model

In [40]:
```
conf_mat=confusion_matrix(y_test,y_pred)
conf_mat
```

Out[40]:
```
array([[47,  7],
       [ 0, 26]], dtype=int64)
```

IBM Deployment

In [2]:
```
!pip install -U ibm-watson-machine-learning
```

```
Requirement already satisfied: ibm-watson-machine-learning in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (1.0.257)
Requirement already satisfied: importlib-metadata in /opt/conda/envs/Python-3.9/lib/python3.9/site-packages (from ibm-watson-ma
chine-learning) (4.8.2)
```

Authenticate and Set Space

In [1]:
```
wml_credentials = {
    "apikey":"QXCDaXxG__YG-rHuYkTQwfeatR70SW8MTP9Lr29ORWNB",
    "url":"https://us-south.ml.cloud.ibm.com"
}
```

In [8]:
```
from ibm_watson_machine_learning import APIClient
import json
```

In [9]:
```
wml_client = APIClient(wml_credentials)
```

In [11]:
```
wml_client.spaces.list()
```

```
Note: 'limit' is not provided. Only first 50 records will be displayed if the number of records exceed 50
-----------------------------------  ------------  ------------------------
```

In [57]:
```
SPACE_ID= "d307bf7a-276c-4443-b82c-f850214871c8"
```

In [58]:
```
wml_client.set.default_space(SPACE_ID)
```

Out[58]:
```
'SUCCESS'
```

In [60]:
```
wml_client.software_specifications.list(500)
```

```
------------------------------  ------------------------------------  ----
NAME                            ASSET_ID                              TYPE
```

In [37]:
```
## Save and Deploy the model
```

In [15]:
```
import sklearn
sklearn.__version__
```

Out[15]:
```
'1.0.2'
```

In [ ]:
```
x = data.iloc[:, :-1]
y = data['id']
```

In [ ]:
```
x.head
```

In [34]:
```
y.tail
```

Out[34]:
```
<bound method NDFrame.tail of 0        0
```

In [ ]:
```
from sklearn.model_selection import train_test_split
x_train, x_test, y_train, y_test = train_test_split(x,y)
```

In [44]:
```
## Build a machine learning model
```

In [ ]:
```
from sklearn.linear_model import LogisticRegression
lgr=LogisticRegression()
lgr.fit(x_train,y_train)
```

In [46]:
```
MODEL_NAME = 'CKD'
DEPLOYMENT_NAME = 'CKD Notebook'
DEMO_MODEL = model
```

In [64]:
```
# Set Python Version
software_spec_uid = wml_client.software_specifications.get_id_by_name('runtime-22.1-py3.9')
```

In [65]:
```
# Setup model meta
model_props = {
    wml_client.repository.ModelMetaNames.NAME: MODEL_NAME,
    wml_client.repository.ModelMetaNames.TYPE: 'scikit-learn_1.0',
    wml_client.repository.ModelMetaNames.SOFTWARE_SPEC_UID: software_spec_uid
}
```

In [66]:
```
#Save model
model_details = wml_client.repository.store_model(
    model=DEMO_MODEL,
    meta_props=model_props,
    training_data=x_train,
    training_target=y_train
)
```

In [67]:
```
model_details
```

Out[67]:
```
{'entity': {'hybrid_pipeline_software_specs': [],
  'label_column': 'id',
  'schemas': {'input': [{'fields': [{'name': 'age', 'type': 'float64'},
     {'name': 'bp', 'type': 'float64'},
     {'name': 'sg', 'type': 'float64'},
```

[70]:
```
model_id = wml_client.repository.get_model_id(model_details)
model_id
```

t[70]:
```
'091254ff-fb60-4562-91c8-67b69e496e35'
```

[71]:
```
# Set meta
deployment_props = {
    wml_client.deployments.ConfigurationMetaNames.NAME:DEPLOYMENT_NAME,
    wml_client.deployments.ConfigurationMetaNames.ONLINE: {}
}
```

In [72]:
```
# Deploy
deployment = wml_client.deployments.create(
    artifact_uid=model_id,
    meta_props=deployment_props
)
```

```
#######################################################################################

Synchronous deployment creation for uid: '091254ff-fb60-4562-91c8-67b69e496e35' started

#######################################################################################


initializing
Note: online_url is deprecated and will be removed in a future release. Use serving_urls instead.

ready


------------------------------------------------------------------------------------------------
Successfully finished deployment creation, deployment_uid='e0e99a97-2d3d-482b-bedf-d1a5f97dfdc4'
------------------------------------------------------------------------------------------------
```

GitHub and Project Demo Link

GitHub Link: https://github.com/IBM-EPBL/IBM-Project-54113-1661597336

Demo Link : google drive