DATE	14-11-2022
TEAM ID	PNT2022TMID50020
PROJECT NAME	EARLY DETECTION OF CHRONIC KIDNEY DISEASE USING MACHINE LEARNING

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## **CHAPTER 1**

## INTRODUCTION

#### 1.1 PROJECT OVERVIEW

Every year, an increasing number of patients are diagnosed with late stages of renal disease. Chronic Kidney Disease, also known as Chronic Renal Disease, is characterized by abnormal kidney function or a breakdown of renal function that progresses over months or years. Chronic kidney disease is often found during screening of persons who are known to be at risk for kidney issues, such as those with high blood pressure or diabetes, and those with a blood family who has chronic kidney disease (CKD). As a result, early prognosis is critical in battling the disease and providing effective therapy. Only early identification and continuous monitoring can avoid serious kidney damage or renal failure. Machine Learning (ML) plays a significant part in the healthcare system, and it may efficiently aid and help with decision support in medical institutions. The primary goals of this research are to design and suggest a machine learning method for predicting CKD. Support Vector Machine (SVR), Random Forest (LR), Artificial Neural Network (ANN), and Decision Tree are four master teaching methodologies investigated (DT). The components are built using chronic kidney disease datasets, and the outcomes of these models are compared to select the optimal model for prediction.

**Keywords**: Chronic Kidney Disease (CKD), Machine Learning (ML), Support Vector Machine (SVR), Random Forest (LR), Artificial Neural Network (ANN), Decision Tree (DT).

#### 1.2 PURPOSE

Every year, an increasing number of patients are diagnosed with late stages of renal disease. Chronic Kidney Disease, also known as Chronic Kidney Disease, is characterized by abnormal kidney function or a breakdown of renal function that progresses over months or years. Chronic kidney disease is often found during screening of persons who are known to be at risk for kidney issues, such as those with high blood pressure or diabetes, and those with a blood family who has chronic kidney disease (CKD). As a result, early prognosis is critical in battling the disease and providing effective therapy. Only early identification and continuous monitoring can avoid serious kidney damage or kidney failure. Machine Learning plays a significant part in the healthcare system, and it may efficiently aid and help with decision support in medical institutions. The primary goals of this research are to design and suggest a machine learning method for predicting CKD. Support Vector Machine (SVR), Random Forest (LR), Artificial Neural Network (ANN), and Decision Tree are four master teaching methodologies investigated (DT). The components are built using chronic kidney disease datasets.

## **CHAPTER 2**

#### LITERATURE SURVEY

#### 2.1 EXISTING PROBLEM

Non communicable illnesses are the leading cause of early death, and CKD is the leading non communicable disease. Chronic Kidney Disease is a major concern for the global health care system. People with CKD must focus on implementing proven, cost-effective therapies to as many people as possible while taking into consideration restricted needs, human and financial resources. Chronic kidney disease (CKD) 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 chronic disease. Recent research suggests that some of the negative outcomes can be avoided with early identification and treatment.

https://www.kidney.org/phi/form?version=health

#### 2.2 REFERENCES

**1. Title:** Prediction of Chronic Kidney disease using adaptive hybridized deep convolutional neural network on the internet of medical things platform

Source: IEEE access Author: G. chenetal Date: October 2020

**2. Title:** Clinical practice guideline for diabetes management in chronic kidney disease

Source: Kidney Int Author: P.T. Coatesetal Date: October 2020

**3.Title:** Overview of clinical prediction models

Source: Ann Transt.Med

Author: L.Chen

Date: November 2019

4. Title: Prediction of chronic kidney diseases using deep artificial neural network technique

**Source:** Springer International Publishing

**Author:** H. kriplani **Date:**June 2019

5.Title: Three types of Machine Learning Algorithms List of Common Machine Learning

Algorithms

Source: IEEE Access

Author: Abdi

Date: November 2016

**6. Title:** Risk Prediction of Chronic Kidney Disease Using Machine Learning Algorithms

Source: ICCCNT

**Author:** N. A. Almansourteal

Date: October 2018

7. Title: Computer-Aided Diagnosis of Chronic Kidney Disease in Developing Countries

**Source:** IEEE Access **Author:** M. Eliete Pinheiro

Date: October 2020

8. Title: Neural network and support vector machine for the prediction of chronic kidney disease

**Source:** Computer Bio Medical **Author:** N. A. Almansouretal

Date: October 2018

9. Title: Prediction of chronic kidney disease (CKD) using Data Science

Source: Conf. Intelligent Computer Control System

Author: N. V. Ganapathi Raju

Date: November 2019

10. Title: Prediction of chronic kidney disease using machine learning

**Source:** Advanced Science Technology

Author: M. V. Maheshwar redy

Date: December 2019

## 2.3 PROBLEM STATEMENT DEFINITION

Every year, an increasing number of patients are diagnosed with late stages of renal disease. Chronic Kidney Disease, also known as Chronic Renal Disease, is characterized by abnormal kidney function or a breakdown of renal function that progresses over months or years. Chronic kidney disease is often found during screening of persons who are known to be at risk for kidney issues, such as those with high blood pressure or diabetes, and those with a blood family who has chronic kidney disease (CKD). As a result, early prognosis is critical in battling the disease and providing effective therapy. Only early identification and continuous monitoring can avoid serious kidney damage or renal failure.

Who does the problem affect?	Those who have a history of kidney failure in their families, diabetes, high blood pressure, or
	heart disease .
What are the boundaries	Diagnosing kidney diseases using parameters
of the problem?	like blood pressure and albumin .
What is the issue?	Kidney function is compromised by a disease or
	condition, leading to its damage over time
When does the issue	When a sickness or illness compromises kidney
occur?	function, causing kidney damage to worsen over
	several months or years
Where is the issue	The small blood veins in the kidneys might
occurring?	become strained by high blood pressure,
	thereby preventing normal functioning of the
	kidney.
	Blood glucose levels that are too high can harm
	the kidneys' small filters.
Why is it important that we fix the problem	The progression of chronic kidney disease to
	an advanced state may be slowed or stopped
	with early detection and treatment

## **CHAPTER 3**

## **IDEATION & PROPOSED SOLUTION**

#### 3.1 EMPATHY MAP CANVAS

#### THINK AND FEEL

- Tell physical Feeling to Doctor.
- They want to relief from chronic kidney disease.
- They feel tierd and weak of our body.

#### HEAR

- To Visit Doctor.
- To Avoid Smoke and Drinks.
- To Give their Own Personal Suggestion.

## SAY AND DO

- Muscle Cramps.
- High Blood Pressure.
- Urinating Less Or More.
- Loss Of Appetite.

#### SEE

- Blood Test Creatinine More Than 2.0 Cause Kidney Disease.
- EGFR:(Estimated Glomerular Filteration Rate) Below 15 Cause Kidney Disease.
- Blood Test :Urea Morethan 30 Cause Kidney Disease.
- Biopsy: It Checks their Kidney Present Condition.
- UrineTest: Albumin More than +++ Causes in Urine Cause Kidney Disease

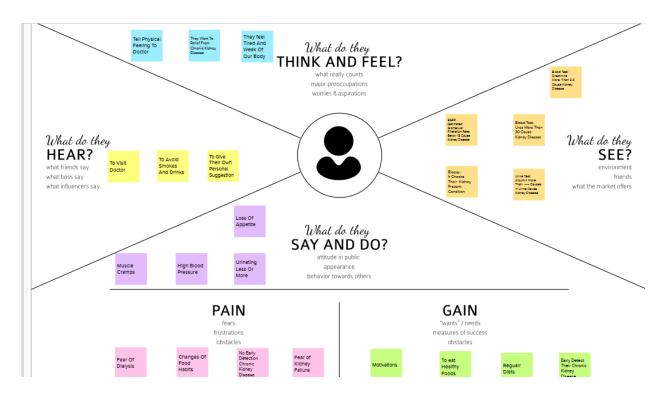
#### GAIN

- Motivations.
- Eat healthy foods.
- Regular diets.

■ Early detect their chronic kidney disease.

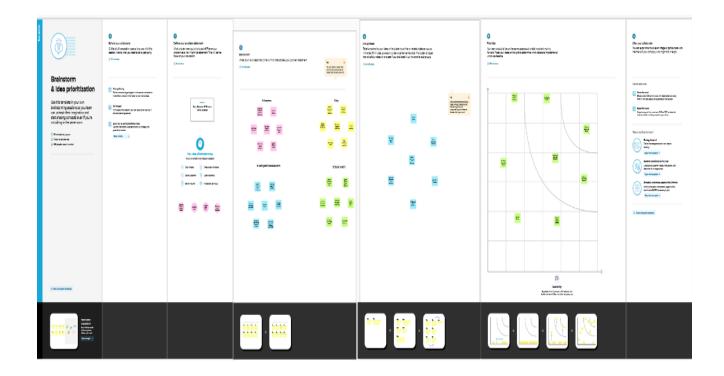
## **PAIN**

- Fear of dialysis.
- Change of food habits.
- Fear of kidney failure.
- No early detection of chronic kidney disease.



## **3.2 IDEATION & BRAINSTORMING**

- They feel tierd and weak .
- Collect information from the user.
- Visit doctor who has Chronic kidney disease.
- Test tha samples.
- To change their food habits.
- Eat healthy foods.



## **3.3 PROPOSED SOLUTION**

S.NO	Parameter	Description
1	Problem Statemeot(Problem to be solved)	Now a days, maoy people
		afected by kidneydisease it
		will be detected io severe
		stage causedeath. Our
		project detects the kidney
		diseasevery accurate using
		web application.
2	Ideas/solution description	To detect the early stage of
		kidney disease in the form of
		web application.
3	Novelty/Uniqueness	User cao detect their disease
		in early stage.
4	Social Impact/Customer Satsifaction	1. To reduce the death rate.
		2. To avoiding progression of
		kidney failure.
		3. To give awareness about
		kidney disease.
5	Business Model(Revenue Model)	1. Can generate revenue
		through direct
		customers
		2. Can collaborate with health
		care sector
		and generate revenue from
		their
		customers
6	Scalability of the solution	Our web application has a
		high security database.

## 3.4 PROBLEM SOLUTION FIT

## **CUSTOMER SEGMENTS**

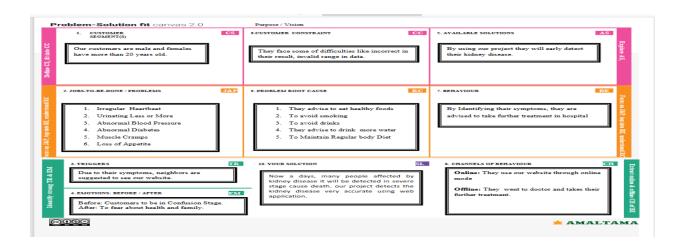
Our customer is male or female more than 20 years old.

## PROBLEM ROOT CAUSE

- Advised to eat healthy foods
- To avoid smoking.
- To avoid drinks.
- Advised to drink more water.
- Follow regular diets.

#### YOUR SOLUTION

■ Now a days, many people affected by kidney disease it will detected in severe stage cause death. our project detects the disease accurately using web application.



## **CHAPTER 4**

## **REQUIREMENT ANALYSIS**

## **4.1 FUNCTIONAL REQUIREMENTS**

Following are the Functional Requirements of proposed solution.

FR	Functional Requirements Sub requirements(Story/subtask)				
NO	(Epic)				
FR-1	User Registration	Registration through Form			
		2. Registration through Gmail			
		3. Registration through LinkedIn.			
FR-2	User Confirmation	1. Confirmation via Email			
		2. Confirmation via OTP			
FR-3	Data Entry	Collect The data from the user			
		2. Fill the Form			
FR-4	Analysing the Data	1. Analyse the data			
		2. Getting their results			
FR-5	Machine Learning	1. Decision Tree			
	Algorithms	2. Rainforest			
		3. Support Vector Machine			
		4. Artificial Neural Network			
FR-6	Phases	1. Testing Phase			
FK-0	Filases	_			
		2. Training Phase			

# **4.2 NON-FUNCTIONAL REQUIREMENTS**

# Following are the Non Functional Requirements of the Proposed Solutions.

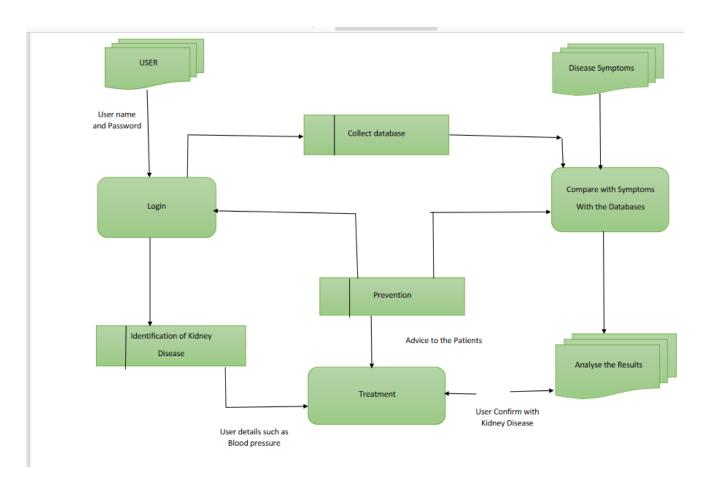
NRF NO	Non-Functional Requiremnts	Description
NFR-1	Usability	Preventing loss of kidney disease.
		2. Delaying oravoiding progression to kidney
		failure.
NFR-2	Security	1. Encrypt your data.
		2. Focus the hosting service measure
		3. Avoid security misconfigurations.
NFR-3	Reliability	Result should be 99.99% accurate.
NFR-4	Performance	Compare the data with symptoms to give
		results.
NFR-5	Availability	It access at any time.
NFR-6	Scalability	1. Memory utilization.
		2. CPU usage.
		3. Networkinput/output.
		4. Disk input/output.

## **CHAPTER 5**

## **PROJECT DESIGN**

## **5.1 DATA FLOW DIAGRAMS**

A Dataflow Diagrams(DFD) is a traditional visual representation of the information flows within a system .A neat and clear DFDcan depict the right amount of the system requirements graphically . it shows how data enters and leave the system, what changesthe information, and where data is stored.

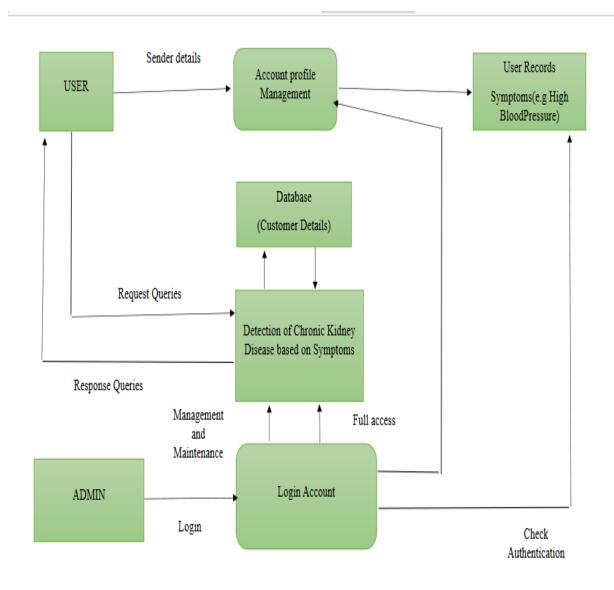


## **USER STORIES**

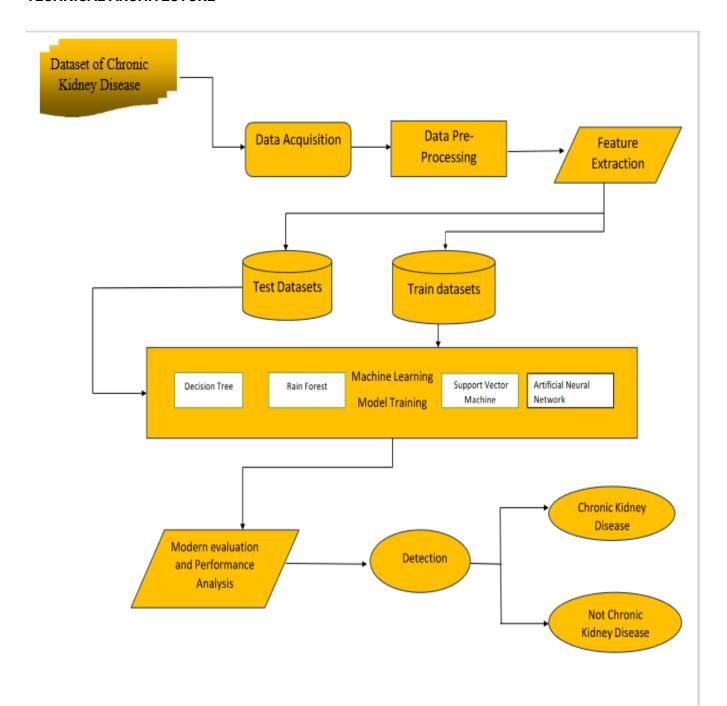
User Type	Functional User Story User story/Task Acceptance		priority	Release		
	Requirements	Number		criteria		
Customer	Registration	USN-1	As a user, I can register for	access account	High	Sprint 1
			the website by mail id,			
			password.			
	Validation	USN-2	As a user will receive a	Receive mail	High	Sprint 2
			conformation mail.			
	Login	USN-3	User will login to the	Create and login	High	Sprint 1
			website by email id,			
			password.			
	Collect	USN-4	User have to answer for 10	Choose answers	High	Sprint 1
	database		-20 questions.			
	Output to user	USN-5	User can view the result.	View results	High	Sprint 2
	Remedies	USN-6	Suggest some advices to	Accept the	Medium	Sprint 2
			the users	remedies		
Administrator	Prediction	USN-7	Admin will build machine	Deploy in the	High	Sprint 1
			learning	websites		
			models to predict chronic			
			kidney disease.			

## **5.2 SOLUTION & TECHNICAL ARCHITECTURE**

## **SOLUTION ARCHITECTURE**



## **TECHNICAL ARCHITECTURE**



**TABLE 1: COMPONENTS & TECHNOLOGIES:** 

SI.NO	COMPONENTS	DESCRIPTION	TECHNOLOGY
1	User Registration	Registration through Form	HTML,
		2. Registration through Gmail	CSS and
		3. Registration through LinkedIn.	Python flask.
2	User Confirmation	1. Confirmation via Email	HTML,
		2. Confirmation via OTP	CSS and
			Python flask.
3	Data Entry	1. Collect The data from the user	HTML, CSS,
		2. Fill the Form	My Sql, Python flask
			Backend Python.
4	Analysing the Data	1. Analyse the data	Machine Learning
		2. Getting their results	Models.
5	Machine Learning	1. Decision Tree	Based on Machine
	Algorithms	2. Rainforest	Learning Concepts.
		3. Support Vector Machine	
		4. Artificial Neural Network	
6	Phases	1. Testing Phase	Machine Learning
		2. Training Phase	and Deep learning
			concepts.

**TABLE 2: APPLICATION & CHARACTERISTICS** 

SI NO	CHARACTERISTICS	DESCRIPTION	TECHNOLOGY
1	Usability	1. Preventing loss of kidney	Cloud(Open Source
		disease.	Platform).
		2. Delaying or	
		avoiding	
		progression to	
		kidney failure.	
2	Security	1. Encrypt your data.	Encryption and
		2. Focus the hosting service	Authentication
		measure	
		3. Avoid security	
		misconfigurations.	
3	Reliability	Result should be 99.99%accurate	Web Development.

4	Performance	Compare the data with	Machine Learning
		symptoms to give	and Deep Learning
		results.	Neural Networks.
5	Availability	It access at any time.	Machine Learning.
6	Scalability	1. Memory utilization.	Performance
		2. CPU usage.	Optimization.
		3. Network input/output.	
		4. Disk input/output.	

## **5.3 USER STORIES**

User Type	Functional	<b>User Story</b>	User story/Task	Acceptance	priority	Release
	Requirements	Number		criteria		
Customer	Registration	USN-1	As a user, I can register for	access account	High	Sprint 1
			the website by mail id,			
			password.			
	Validation	USN-2	As a user will receive a	Receive mail	High	Sprint 2
			conformation mail.			
	Login	USN-3	User will login to the	Create and login	High	Sprint 1
			website by email id,			
			password.			
	Collect	USN-4	User have to answer for 10	Choose answers	High	Sprint 1
	database		-20 questions.			
	Output to user	USN-5	User can view the result.	View results	High	Sprint 2
	Remedies	USN-6	Suggest some advices to	Accept the	Medium	Sprint 2
			the users	remedies		
Administrator	Prediction	USN-7	Admin will build machine	Deploy in the	High	Sprint 1
			learning	websites		
			models to predict chronic			
			kidney disease.			

## **CHAPTER 6**

## **PROJECT PLANNING & SCHEDULING**

## **6.1 SPRINT PLANNING & ESTIMATION**

Sprint	Functional Requirement (Epic)	User Story Numb er	User Story / Task	Story Points	Priority	Team Members
Sprint-1	User Registration	USN-1	As a user, I can register for the application by entering my name, mobile number, email, password, and confirming my password.	10	High	S.Rubashree S.Sudarmathi S.Sathya C.Raju
Sprint-2		USN-2	As a user, I can register for the application through Gmail	5	Medium	S.Rubashree S.Sudarmathi S.Sathya C.Raju
Sprint-1	User Confirmation	USN-3	As a user, I will receive confirmation email once I have registered for the application	10	High	S.Rubashree S.Sudarmathi S.Sathya C.Raju
Sprint-2		USN-4	As a user, I will receive confirmation otp to verify the identity.	5	High	S.Rubashree S.Sudarmathi S.Sathya C.Raju
Sprint-2	Data Collection	USN-5	As a user, I will enter the input data for disease prediction in the form	10	High	S.Rubashree S.Sudarmathi S.Sathya C.Raju
Sprint-3	Provide output to the user	USN-6	As a user, I will get the result of disease prediction in the dashboard.	10	High	S.Rubashree S.Sudarmathi S.Sathya C.Raju

Sprint-3	Data Analysis	USN-7	As the admin, I will develop modules to	10	High	S.Rubashree
			preprocess and store the data.			S.Sudarmathi
						S.Sathya
						C.Raju
Sprint-4	Prediction of	USN-8	As the admin, I will build a Machine	10	High	S.Rubashree
	disease		Learning model to predict the disease			S.Sudarmathi
						S.Sathya
						C.Raju
Sprint-4	Final Delivery	USN-9	Deploy the application in IBM cloud and	10	High	S.Rubashree
			make it available for use.			S.Sudarmathi
						S.Sathya
						C.Raju

## **6.2 SPRINT DELIVARY SCHEDULE**

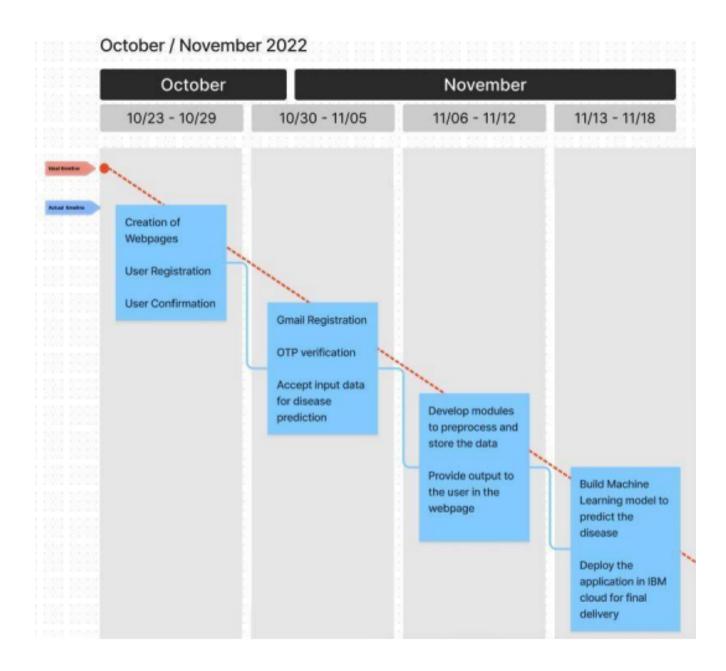
Sprint	Total Story Points	Duration	Sprint Start Date	Sprint End Date (Planned)	Story Points Completed (as on Planned End Date)	Sprint Release Date (Actual)
Sprint -1	20	6 Day	24 Oct 2022	29 Oct 2022	20	29 Oct 2022
Sprint -2	20	6 Day	31 Oct 2022	05 Nov 2022	20	05 Nov 2022
Sprint -3	20	6 Day	07 Nov 2022	12 Nov 2022	20	12 Nov 2022
Sprint -4	20	6 Day	14 Nov 2022	19 Nov 2022	20	19 Nov 2022

## Velocity:

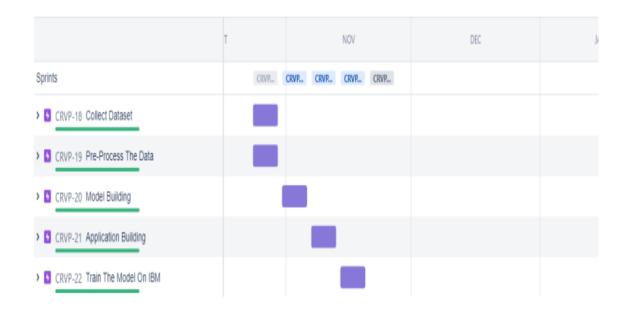
We have a 6-day sprint duration, and the velocity of the team is 20 (points per sprint). Theteam's average velocity (AV) per iteration unit (story points per day)

AV = Sprint duration / velocity = 20 / 6 = 3.33

## **Burndown Chart**



## **6.3 REPORTS FROM JIRA**



## **CHAPTER-7**

## **CODING & SOLUTIONING**

## **7.1 FEATURE 1**

## index.html

```
rgba(20,20,20, .75),
           rgba(20,20,20,.75)),
           url(
'https://pharmanewsintel.com/images/site/article_headers/_normal/Chronic_Disease_Manage.
png');
.container {
 border: 2px solid #ccc;
 padding: 10px;
 width: 20em;
height:21em;
background-color:white;
}
.hello{
opacity: 0.5;
</style>
</head>
<body>
<marquee bgcolor="white">IBM NALAIYA THIRAN</marquee>
<center>Chronic Disease Prediction</center>
<form action="/val" method="post"><center>
 <label for="age">Age:</label><br>
 <input type="number" id="age" name="age"><br><br><br>
 <label for="bp">Blood Pressure:</label><br>
 <input type="number" id="bp" name="bp">
<br>
<br>
<br>
 <label for="sg">Urinary Specific Gravity:</label><br>
 <input type="number" id="sg" name="sg">
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<br>
 <label for="al">al:</label><br>
 <input type="number" id="al" name="al">
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<br>
<label for="su">su:</label><br>
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 <label for="rbc">rbc:</label><br>
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<label for="pc">pc:</label><br>
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 <label for="sod">sod:</label><br>
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<label for="pot">pot:</label><br>
<input type="number" id="pot" name="pot">
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<label for="pcv">pcv:</label><br>
<input type="text" id="pcv" name="pcv">
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<br>
<label for="pe">pe:</label><br>
 <input type="text" id="pe" name="pe">
<br>
<br>
<br>
 <label for="ane">ane:</label><br>
 <input type="text" id="ane" name="ane">
<br>
<br/>center>
 <center><button type="submit">Check</button></center>
</form>
</body>
</html>
```

## rename.html

```
<html>
<head>
<style>
body { background-color: #E6E6FA;
}
</style>
</head>
<body >
<br>
<br>
<br>
<br>
<br>
<br>
<center><h1>{{answer1}}</h1></center>
```

```
<br/>
<br/>
<center><h1>{{answer2}}</h1></center><br/>
<center><img alt="Qries" src="https://image.shutterstock.com/image-vector/cute-illustration-sick-kidney-characters-260nw-1529381825.jpg width=500" height="500"></center><br/>
</body><br/>
</html>
```

#### rename2.html

```
<html>
<head>
<style>
body {
background-color: #E6E6FA;
}
</style>
</head>
<body >
<br>
<br>
<br>
<center><h1>{{answer1}}</h1></center>
<br>
<center><h1>{{answer2}}</h1></center>
<center><img alt="Qries" src="https://media.istockphoto.com/vectors/vector-kidney-cartoon-</pre>
human-body-health-organ-smiling-mascot-on-vector-
id1136533246?k=20&m=1136533246&s=612x612&w=0&h=eWjhtoVOfX3lBF4fcsDD4ZLLzjeJ_A
```

```
x4cgdgdUexG1Q="
width=500" height="500"></center>
</body>
</html>
```

#### **7.2 FEATURE 2**

```
import pickle
loaded_class = pickle. load(open('randomclass_chronic', 'rb'))
loaded_reg = pickle. load(open('randomreg_chronic', 'rb'))
import numpy as np
import pandas as pd
from flask import Flask, request, redirect, render_template
app = Flask(__name__)
@app.route("/",methods=['GET', 'POST'])
def index():
  return render_template('index.html')
@app.route("/val",methods=['POST'])
def val():
  test=[]
  if request.method == 'POST':
    test.append(request.form.get("age"))
    test.append(request.form.get("bp"))
    test.append(request.form.get("sg"))
    test.append(request.form.get("al"))
    test.append(request.form.get("su"))
    rb=request.form.get("rbc")
    if rb=='abnormal':
      test.append(1)
    else:
      test.append(0)
    pc=request.form.get("pc")
    if pc=='abnormal':
      test.append(1)
    else:
      test.append(0)
    pcc=request.form.get("pcc")
```

```
if pcc=='present':
  test.append(1)
else:
  test.append(0)
ba=request.form.get("ba")
if ba=='present':
  test.append(1)
else:
  test.append(0)
test.append(request.form.get("bgr"))
test.append(request.form.get("bu"))
test.append(request.form.get("sc"))
test.append(request.form.get("sod"))
test.append(request.form.get("pot"))
test.append(request.form.get("hemo"))
test.append(request.form.get("pcv"))
test.append(request.form.get("wc"))
test.append(request.form.get("rc"))
ht=request.form.get("htn")
if ht=='yes':
  test.append(1)
else:
  test.append(0)
d=request.form.get("dm")
if d=='yes':
  test.append(1)
else:
  test.append(0)
ca=request.form.get("cad")
if ca=='yes':
  test.append(1)
else:
  test.append(0)
ap=request.form.get("appet")
if ap=='good':
  test.append(1)
elif ap=='poor':
  test.append(0)
else:
  test.append(np.nan)
p=request.form.get("pe")
```

```
if p=='yes':
      test.append(1)
    else:
      test.append(0)
    an=request.form.get("ane")
    if an=='yes':
      test.append(1)
    else:
      test.append(0)
  print(test)
  test_df=pd.DataFrame(test)
  test_df=np.array(test_df).reshape(1, -1)
  ans1=loaded_class.predict(test_df)
  ans2=loaded_reg.predict(test_df)
  if int(ans1)==1:
    answer1="Sorry to say!! You have CHRONIC DISEASE!!!"
    return render_template('rename.html',answer1=answer1,answer2=ans2)
  else:
    answer1="Happy to say that you don't have CHRONIC DISEASE"
    return render_template('rename2.html',answer1=answer1,answer2=ans2)
if __name__ == "__main__":
  app.debug=True
  app.run(debug=False)
```

#### 7.3 DATABASE SCHEMA

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
from sklearn.model_selection import train_test_split, GridSearchCV
from sklearn.metrics import roc_curve, auc, confusion_matrix,
classification_report,accuracy_score
from sklearn.ensemble import RandomForestClassifier
import warnings
warnings.filterwarnings('ignore')
%matplotlib inline
def auc_scorer(clf, X, y, model): # Helper function to plot the ROC curve
  if model=='RF':
    fpr, tpr, _ = roc_curve(y, clf.predict_proba(X)[:,1])
  elif model=='SVM':
    fpr, tpr, _ = roc_curve(y, clf.decision_function(X))
  roc_auc = auc(fpr, tpr)
  plt.figure() # Plot the ROC curve
  plt.plot(fpr, tpr, label='ROC curve from '+model+' model (area = %0.3f)' % roc_auc)
  plt.plot([0, 1], [0, 1], 'k--')
  plt.xlim([0.0, 1.0])
  plt.ylim([0.0, 1.05])
  plt.xlabel('False Positive Rate')
  plt.ylabel('True Positive Rate')
  plt.title('ROC Curve')
  plt.legend(loc="lower right")
  plt.show()
  return fpr,tpr,roc_auc
```

```
df = pd.read_csv("C:/Users/Sinegalatha/Desktop/2nd year online class/nalaiya
thiran/dataset/kidney_disease.csv")
df.head()
df['wc']
df.info()
df.describe()
df[df.duplicated()]
df.isna().sum()
df2 = df.dropna(axis=0)
df.head()
df2['class'].value_counts()
corr_df = df2.corr()
mask = np.zeros_like(corr_df, dtype=np.bool)
mask[np.triu_indices_from(mask)] = True
# Set up the matplotlib figure
f, ax = plt.subplots(figsize=(11, 9))
# Generate a custom diverging colormap
cmap = sns.diverging_palette(220, 10, as_cmap=True)
# Draw the heatmap with the mask and correct aspect ratio
sns.heatmap(corr_df, mask=mask, cmap=cmap, vmax=.3, center=0,
      square=True, linewidths=.5, cbar_kws={"shrink": .5})
plt.title('Correlations between different predictors')
plt.show()
X_train, X_test, y_train, y_test = train_test_split(df2.iloc[:,:-1], df2['class'],
                             test_size = 0.33, random_state=44,
                            stratify= df2['class'])
X_train.head()
print(X_train.shape)
print(X_test.shape)
```

```
y_train.value_counts()
tuned_parameters = [{'n_estimators':[7,8,9,10,11,12,13,14,15,16],'max_depth':[2,3,4,5,6,None],
            'class_weight':[None,{0: 0.33,1:0.67},'balanced'],'random_state':[42]}]
clf = GridSearchCV(RandomForestClassifier(), tuned_parameters, cv=10,scoring='f1')
clf.fit(X_train, y_train)
print("Detailed classification report:")
y_true, Ir_pred = y_test, clf.predict(X_test)
print(classification_report(y_true, lr_pred))
confusion = confusion_matrix(y_test, lr_pred)
print('Confusion Matrix:')
print(confusion)
fpr,tpr,roc_auc = auc_scorer(clf, X_test, y_test, 'RF')
print('Best parameters:')
print(clf.best_params_)
clf_best = clf.best_estimator
plt.figure(figsize=(12,3))
features = X_test.columns.values.tolist()
importance = clf_best.feature_importances_.tolist()
feature_series = pd.Series(data=importance,index=features)
feature_series.plot.bar()
plt.title('Feature Importance')
list_to_fill = X_test.columns[feature_series>0]
print(list_to_fill)
corr_df = pd.isnull(df).corr()
mask = np.zeros_like(corr_df, dtype=np.bool)
mask[np.triu_indices_from(mask)] = True
f, ax = plt.subplots(figsize=(11, 9))
cmap = sns.diverging_palette(220, 10, as_cmap=True)
sns.heatmap(corr_df, mask=mask, cmap=cmap, vmax=.3, center=0,
      square=True, linewidths=.5, cbar_kws={"shrink": .5})
```

```
plt.show()
df2 = df.dropna(axis=0)
no_na = df2.index.tolist()
some_na = df.drop(no_na).apply(lambda x: pd.to_numeric(x,errors='coerce'))
some_na = some_na.fillna(0) # Fill up all Nan by zero.
X_test = some_na.iloc[:,:-1]
y_test = some_na['class']
y_true = y_test
lr_pred = clf_best.predict(X_test)
print(classification_report(y_true, lr_pred))
confusion = confusion_matrix(y_test, lr_pred)
print('Confusion Matrix:')
print(confusion)
print('Accuracy: %3f' % accuracy_score(y_true, lr_pred))
# Determine the false positive and true positive rates
fpr,tpr,roc_auc = auc_scorer(clf_best, X_test, y_test, 'RF')
import pickle
pickle. dump(clf_best, open('randomclass_chronic', 'wb')
from sklearn.ensemble import RandomForestRegressor
reg=RandomForestRegressor()
reg.fit(X_train,y_train)
RandomForestRegressor()
y_pred=reg.predict(X_test)
pickle. dump(reg, open('randomreg_chronic', 'wb'))
y_pred
1_pred=list(y_pred)
l_test=list(y_test)
d={'prob':l_pred,'out':y_test}
df_i=pd.DataFrame(d)
df_i.head()
df_i.to_csv('C:/Users/Sinegalatha/Desktop/2nd year online class/nalaiya
thiran/output/file1.csv')
```

# **CHAPTER - 8**

# **TESTING**

## **8.1 TEST CASES**

TEST CASE NO	TEST CASE SCENARIO	RESULT
1	Verify user is able to fill id and password and login to the website	Pass
2	Verify user has to fill the information	Pass
3	Verify user is able to navigate from homepage to other pages?	Pass
4	Verify user is able to get the expected results accurately	Pass
5	Verify information collected page elements	Pass

## **8.2 USER ACCEPTANCE TESTING**

Section	Total	Not Tested	Fail	Pass	
	Cases				
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	4	0	0	4	
Output					
Version Control	2	0	0	2	

## **CHAPTER - 9**

## **RESULT**

## 9.1 PERFORMANCE METRICES

Here we will be evaluating the model built. We will be using the test set for evaluation. The test set is given to the model for prediction and prediction values are stored in another variable called y\_pred. The score of the model is calculated and its performance is estimated.

Learning Algorithm			Training Time
	Test Data	Training Data	
Linear Regression	0.84	0.85	6 Minutes

## **CHAPTER - 10**

## **ADVANTAGES & DISADVANTAGES**

#### **ADVANTAGES**

- In early detection of CKD should be beneficial because it enables clinicians to initiate effective treatment of mild disease, preventing loss of kidney function and delaying or avoiding progression to kidney failure.
- Higher preexisting rate of kidney testing in at-risk groups
- A lower prevalence of disease in people with and without such risk factors
- A higher prevalence of other diseases that require similar treatment among people with CKD,and which could be detected instead
- A lower baseline likelihood of developing kidney failure in the reservoir of cases (e.g., slower progression or shorter life expectancy in the absence of early detection)
- Lower capacity to treat newly identified cases of CKD
- Overtesting in the target population upon initiation of early detection (e.g., testing more frequently than necessary, duplicate testing from multiple providers)

#### **DISADVANTAGES**

- Poverty related factors such as infectious diseases secondary to poor sanitation, inadequate supply of safe water, environmental pollutants and high concentrations of disease-transmitting vectors continue to play an important role in the development of CKD in low-income countries.
- Although rates of diabetic nephropathy are rising, chronic glomerulonephritis and interstitial nephritis are among the principal causes of CKD in many countries. Of note is the emergence of HIV-associated nephropathy as the major cause of CKD in Sub-Saharan Africa.
- A high prevalence of CKD of unknown etiology has been reported in rural agricultural communities from Central America, Egypt, India and Sri Lanka. Male farmworkers are affected disproportionately, and the clinical presentation is suggestive of interstitial nephritis, confirmed on renal biopsies.
- The strong association with farm work has led to suggestions that exposure to agrochemicals, dehydration, and consumption of contaminated water might be responsible. Additionally, the use of traditional herbal medications is common and frequently associated with CKD among the poor.
- In Mexico, CKD prevalence among the poor is 2-3fold higher than the general population, and the etiology is unknown in 30% of ESRD patients.

#### **CHAPTER -11**

#### CONCLUSION

This study developed an algorithm for predicting CKD at an early stage. The dataset contains input parameters obtained from CKD patients, and the models are trained and validated using the valid parameters. To diagnose CKD, decision tree, random forest, and support vector machine learning models are built. The accuracy of prediction is used to assess the performance of the models. The study's findings revealed that the Random Forest Classifier model outperforms Decision Trees and Support Vector Machines in predicting CKD. As an extension of this research, the comparison may also be done depending on the duration of execution and feature set selection.

## **CHAPTER - 12**

#### **FUTURE SCOPE**

The National Kidney Foundation's Kidney Disease Outcomes Quality Initiative (K/DOQI) Clinical Practice Guidelines on Chronic Kidney Disease: Evaluation, Classification and Stratification of Risk published in 2002 provided the first definition of CKD independent of cause, and classification of severity based on GFR level. The guidelines have been widely disseminated and generally accepted. However, concerns have been expressed about the definition and classification, methods to estimate GFR, and ascertainment of proteinuria.

The goals for the KDIGO Controversies Conference were to provide a clear understanding to both the nephrology and non nephrology communities of the evidence base for the K/DOQI definition and classification of severity of CKD; to develop global consensus for the adoption of a simple definition and classification system for CKD, clarifications and modifications to current guidelines to facilitate more widespread implementation of initiatives for patient care and physician and public education worldwide; and to identify a collaborative research agenda and plan that would improve the evidence base and facilitate the implementation of the definition and classification of CKD

## **CHAPTER - 13**

## **APPENDIX**

**GitHub & Project Demo Link** 

https://github.com/IBM-EPBL/IBM-Project-41116-166063955 https://drive.google.com/file/d/1o\_hUlkIA765i-7JuOjQ\_r8UC4t-AjAlX/view?usp=share\_link