

**HEART DISEASE DETECTION USING MACHINE LEARNING
ALGORITHM**

Project report submitted in partial fulfillment of the requirement for the degree of
Bachelor of Technology

in

Computer Science and Engineering/Information Technology

By

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Under the supervision of

Mr.Praveen Modi

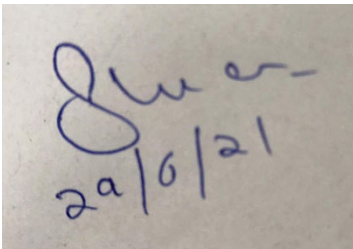
To



Department of Computer Science & Engineering and Information Technology
**Jaypee University of Information Technology Waknaghat, Solan-173234,
Himachal Pradesh**

DECLARATION

We hereby declare that the work reported in the B.Tech Project Report entitled “<Heart Disease Detection Using Machine Learning Algorithm>” submitted at **Jaypee University of Information Technology, Waknaghat, India** is an authentic record of our work carried out under the supervision of <**DR. Parveen Modi**>. We have not submitted this work elsewhere for any other degree or diploma.

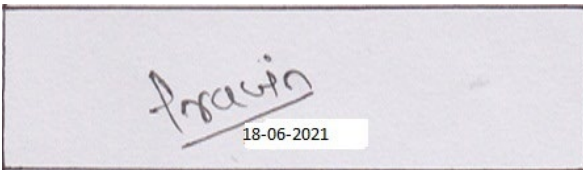


<Signature>

Shivam Mittal

171276

This is to certify that the above statement made by the candidates is correct to the best of my knowledge.



<Signature of the Supervisor>

Dr. Parveen Modi

Associate Professor

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Date:18-05-21

Head of the Department/Project Coordinator

DEPARTMENT OF COMPUTER SCIENCE ENGINEERING

&

INFORMATION TECHNOLOGY

CERTIFICATE

This is to certify that the work in this Project title as “**Heart Disease Detection Using Machine Learning Algorithm**” is entirely written, successfully completed and demonstrated by the following students themselves as a fulfillment of requirement for Bachelor’s of Engineering in Computer Science.

Shivam Mittal(171276)

ACKNOWLEDGEMENTS

Thanks to Almighty GOD Who gave us courage and understanding to start and later on finish this work of project. Thanks to **HOD** Sir and our Supervisor **DR. PARVEEN MODI** who guided and helped us at the different phases of the project.

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ABSTRACT

Heart disease is one of the biggest cause of death in this era today. With Regular Physical work and decent eating habit can may be prevent it to some extent. In this paper we have tried to implement the important Machine Learning algorithm to predict the heart disease in a patent. In this paper we used five different algorithm of Machine Learning such as Logistic Regression, K-Nearest Neighbor, Naïve Bayes, Random Forest and decision tree but in which Logistic Regression gives the higher accuracy of approximately 91.80%.

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INTRODUCTION

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1.1 INTRODUCTION

It is very difficult to predict the heart disease because of some risky factors such as diabetes, high blood pressure, high cholesterol and many other serious factors. Machine Learning is the most efficient technology in the today's era that provides systems the ability to automatically learn and improved from its own experiences. The heart disease is classified based on some methods such as- Logistic Regression, K-Nearest Neighbor (KNN), Naive Bayes (NB), Decision Tree [Abstract](DT) and Random Forest algorithm. After using these algorithm we find the accuracy mean of each algorithm that predicts which algorithm best measure the Heart disease efficiently. The data of heart disease patients collected from Kaggle.

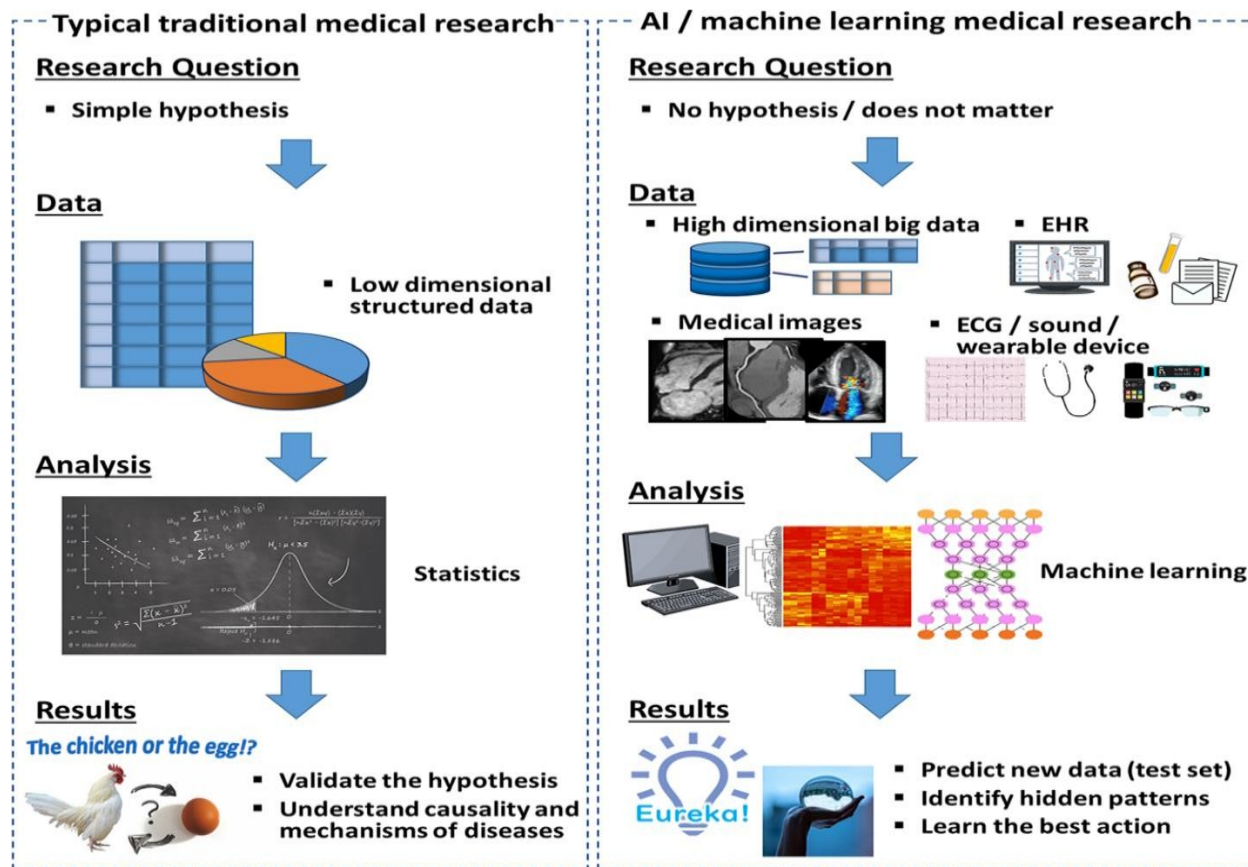


Fig: 1 (Research Overview)

1.2 **OBJECTIVES**

Advancement in the Medical Technologies. Detecting a person is suffering from any Heart disease or not? Its implementation as a desktop application

1.3 **PROBLEM STATEMENT**

Now understanding Heart disease so any kind of disorder in heart is refers as heart disease

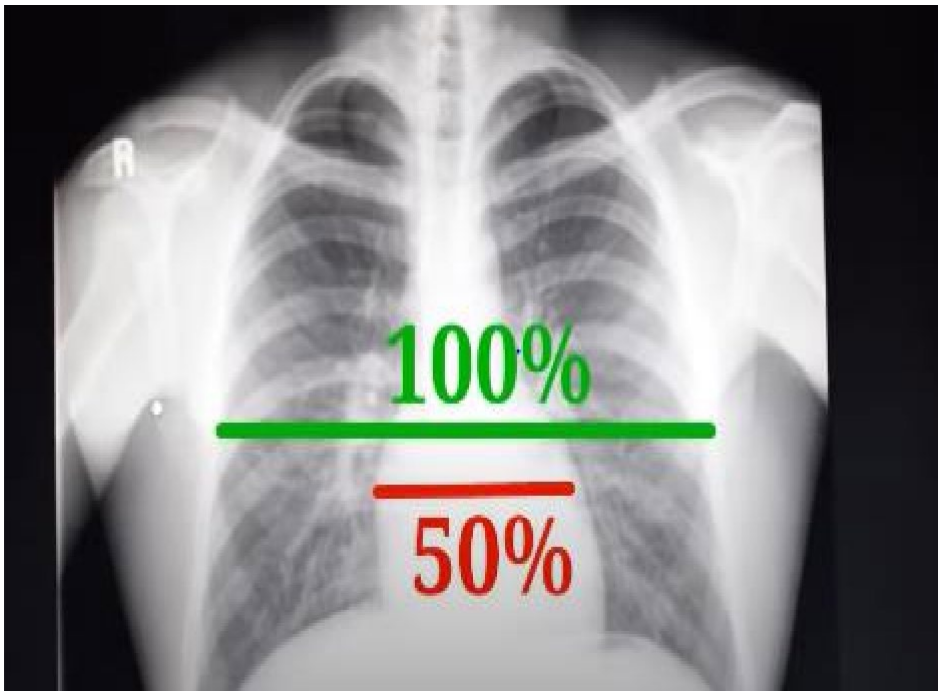


Fig: 2 (X-Ray Scan Image)

So if the shown area 50% increase while a little bit then there is a definitely chances of heart disease and the problem is to detect these heart disease so for this purpose we have a collected dataset which have records of 303 patients having 14 attributes so using these attributes and applying Machine learning model for detecting heart disease is our task.

1.4 METHODOLOGY

For this task as we defined earlier, we have record of 303 patients having 14 attributes so

- first task is testing and training for this task we use 80% of our data as training purpose and rest 20% is for testing purpose because this is an efficient way for study features.
- Now second task is data wrangling it means reduce the irrelevant components.

1.5 SCOPE

As the future of machine learning is very wide so scope of this project is very clear that this software is helpful for detecting a person is suffering from any kind of Heart disease or not ? and according to the latest study in the area of medical domain related to machine learning which is related to present scenario of COVID-19 epidemic Dr. Kamlesh Jain Professor in the department of Civil Engineering at Indian Institute of Technology, Roorkee develops a software that detects COVID-19 so in future we can also apply the similar task for Heart disease prediction.

LITERATURE REVIEW

2.1 HISTORY

Many researchers work in this field Dr. John Moraque first develops the first machine learning model but after this many researchers work in that field and contribute in the medical domain.

2.2 ANALYSIS

Today medical services have come a long way to treat patients with various diseases. Among the most lethal one is the heart disease problem which cannot be seen with a naked eye and comes instantly when its limitations are reached. Today diagnosing patients correctly and administering effective treatments have become quite a challenge.

2.3 RECENT STUDY

The appreciable work of Dr. Deepak Ranjan Nayak (Currently working as an Assistant Professor in the Department of Computer Science & Engineering at SVNIT Surat) in the field of medical domain pathology and gives different-different ideas related to prediction of Medical disease through Computer vision and Machine learning Model.

2.4 APPLICATION

2.4.1 From Medical Perspective

SYSTEM DEVELOPMENT

3.1 OVERVIEW OF METHOD

Machine Learning process starts from preprocessing data phase followed by feature selection, classification.

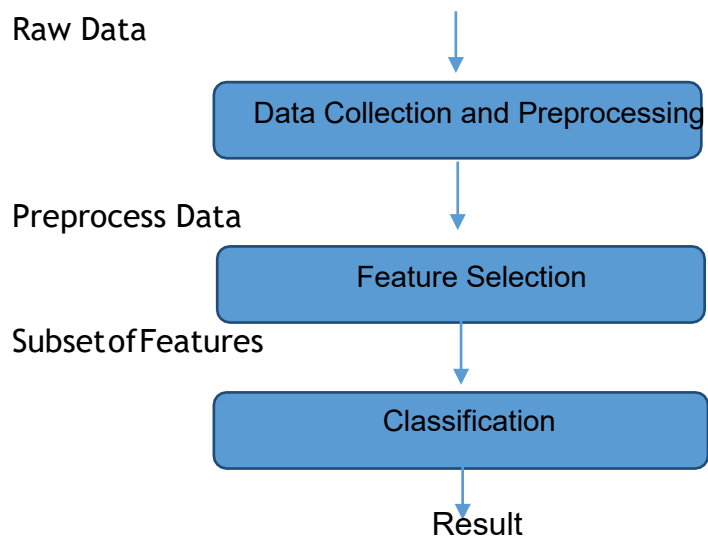


Fig: 3 (Overview Of Method)

3.2 DATA COLLECTION

Data is pre-processed after collection of various records. This dataset contains the record of 303 patients. The patient having heart disease or not is set to 1 or 0. There are 14 columns in our dataset which is described as:

ATTRIBUTE	DESCRIPTION
Age	Displays the age of the individual

Sex	Display the sex (0=female and 1=male)
Cp	Chest pain type (0=typical angina, 1=atypical angina, 2= non-angina, 3= asymptotic)
Trestbps	Level of blood pressure at resting mode (in mm/Hg at time of admitting in hospital)
Chol	Serum cholesterol (in mg/dl)
Fbs	Fasting blood sugar of individual with 120mg/dl (if fbs>120mg/dl then: 1(true) else: 0(false))
Restecg	Resting ECG (0=normal, 1=having ST-T wave abnormality,2=left ventricular hypertrophy)
Thalach	Maximum heart rate achieved
Exang	Exercise induced angina (1=yes and 0=no)
Oldpeak	ST depression induced by exercise relative to rest
Slope	Peak exercise ST segment (0=upsloping, 1=flat, 2= downsloping)
Ca	Fluoroscopy colored major vessels numbered from 0 to 4
Thal	Thalassemia (1=normal, 2=fixed defect, 3=reversible defect)
Target	Diagnosis of heart disease (0=absence, 1=present)

3.3 FEATURE SELECTION

among the 14 attributes there may be some columns which may be irrelevant can be dropped but for this there is need to train the data and the rest coming attributes is considered as an important factors for predicting the Heart disease.

Feature selection means after dividing our data into training and testing (here we reserve data for training for 80% and testing for 20% because this is an efficient way) then apply the tools of machine learning

The correlation between these attributes as shown below:

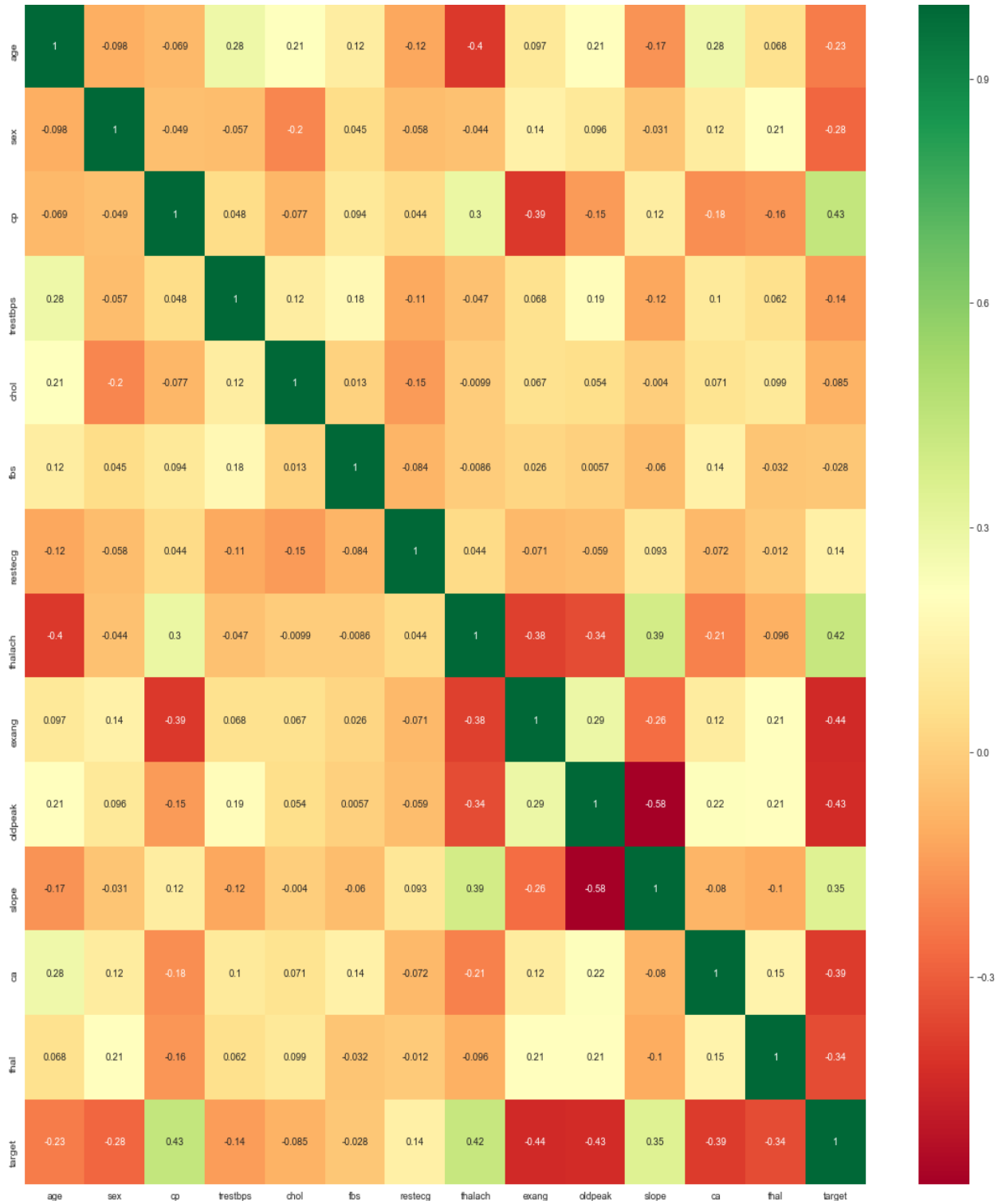
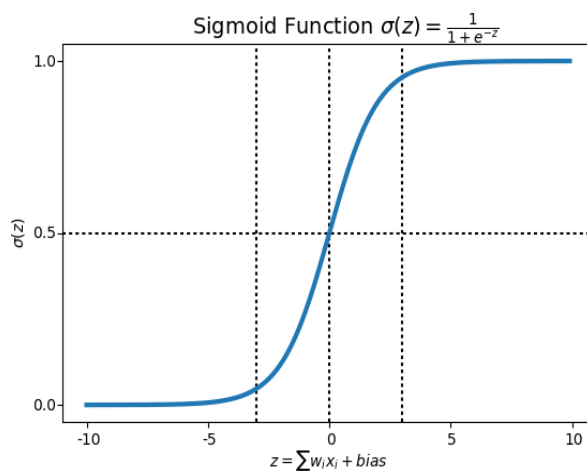


Fig: 4 (Correlation Matrix of Data)

3.4 CLASSIFICATION

Apply the different-different Machine Learning algorithm and Measure the best one.

- i) **LOGISTIC REGRESSION**: It is used for classification problem. It is a predictive analysis algorithm and based on the concept of probability. The hypothesis of logistic regression is to limit the cost function between 1 and 0. In machine learning, we use sigmoid function to map predictions to probabilities which is represented as:



Graph: 1 (Logistic Regression)

It will first Analyzing the data followed by Collecting Data, Data Wrangling, Train & Test, Accuracy Check.

- ii) **K-Nearest Neighbor**: It extract the knowledge based on the samples Euclidean distance function $d(X_i, Y_i)$ and the majority of K-nearest neighbors:-
 $d(X_i, Y_i) = [(X_1 - Y_1)^2 + (X_2 - Y_2)^2 + \dots + (X_i - Y_i)^2]^{0.5}$

- iii) **NAIVE BAYES:** It is a probabilistic classifier. It is based on probability models that incorporate assumptions. It assumes that the value of a particular feature is independent of the value of any other feature, given the class variable.

- iv) **DECISION TREE:** For training samples of data D, the trees are constructed based on high entropy inputs. These trees are simple and fast constructed in a top down recursive divide and conquer approach. Tree pruning is performed to remove the irrelevant samples on D.

- v) **RANDOM FOREST:** In this technique, a set of decision trees are grown and each tree votes for the most popular class, then the votes of different trees are integrated and a class is predicted for each sample. This approach is designed to increase the accuracy of the decision tree, more trees are produced to vote for class prediction. This approach is ensemble classifier composed of some decision trees and the final result is the mean of individual trees results.

PERFORMANCE ANALYSIS

4.1 **System Requirement:** Python Editor as an essential such as Spyder (Anaconda). You can also use inPycharm, Python IDE 3.7.4 or many more editors for python development and you can also build this project using Matlab (Octave) depends on Software Developer but for this project we use Spyder (Anaconda) platform.

4.1.1 Minimum Requirements:

- a. 200 MHz Processor
- b. 64 MB of RAM

4.1.2 Best Requirements:

- a. 1.6 GHz processor
- b. 128 MB or more of RAM

4.2 **Library Requirement:**

4.2.1 NumPy :

NumPy stands for Numeric Python. It is used in adding support for large multi-dimensional arrays and matrices along with large collection of high level mathematical functions to operate on arrays.

4.2.2 Pandas:

It is a software library written for python programming language for data manipulation and analysis. It offers data structures and operations for manipulating numerical tables and time series.

4.2.3 Matplotlib:

It is a plotting library in python. It provides an object-oriented API for embedding plots into applications using general purpose GUI toolkits like Tkinter, rcParams, rainbow etc.

4.2.4 skLearn:

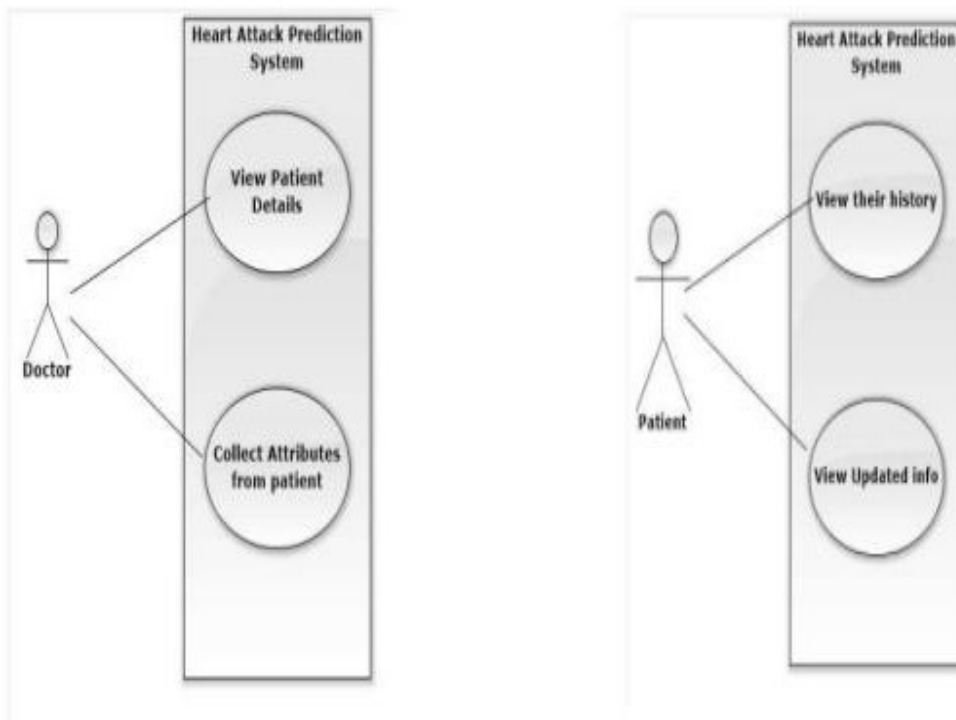
skLearn stands for scikit-learn. It is a open source library that has powerful tools for data analysis and data mining. It supports Python numerical and scientific libraries like NumPy and SciPy.

4.2.5 Seaborn:

It is a Python data visualization library based on matplotlib. It provides a high level interface for drawing attractive and informative statistical graphics.

43 System Development Approach :

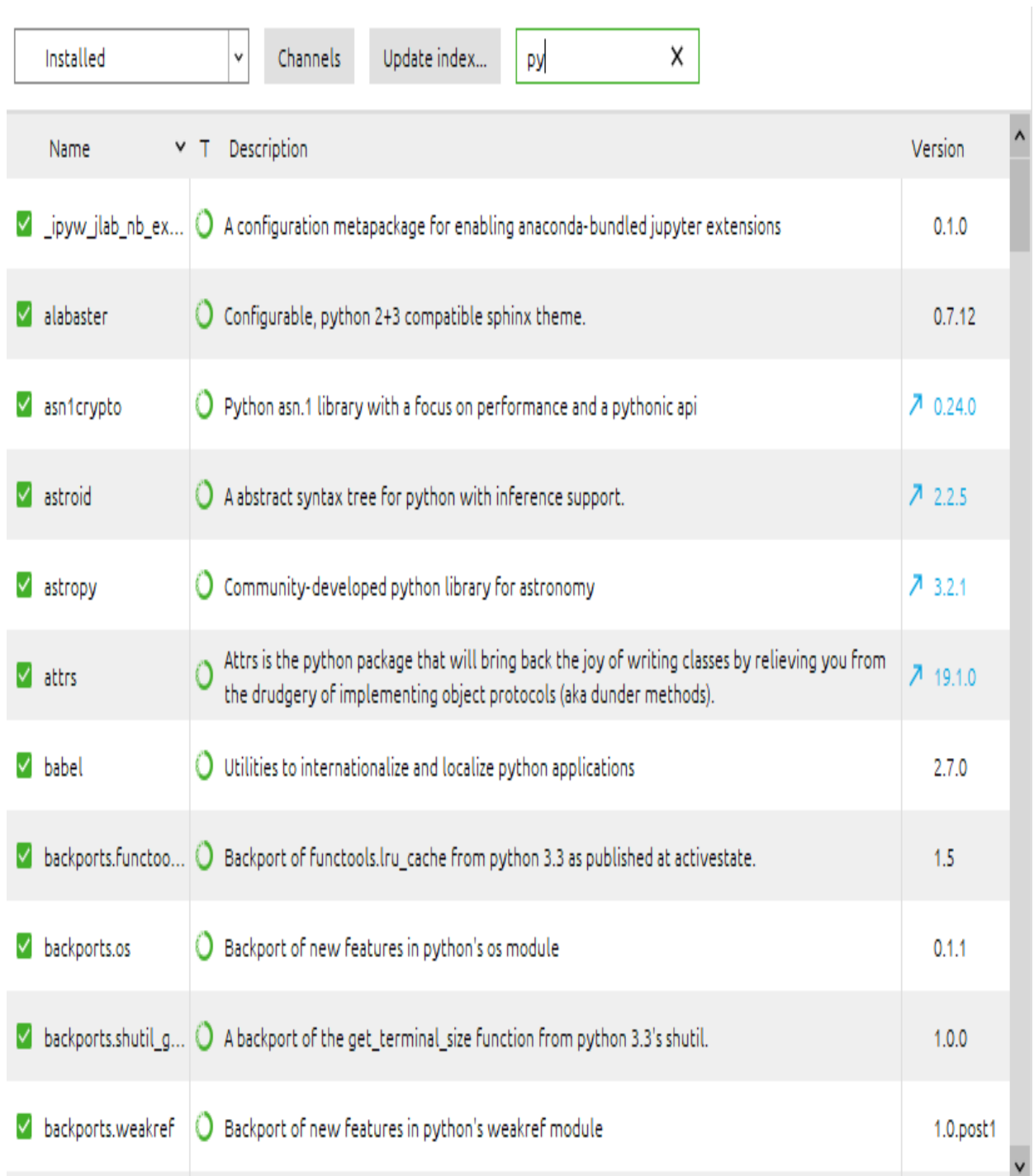
Use Case Diagrams



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Fig: 5 (Use Case Diagram)

4.4 **Installation of Packages:**



The screenshot shows the Anaconda Package Environment Manager interface. At the top, there is a search bar containing 'py|' and a dropdown menu set to 'Installed'. Below the search bar are buttons for 'Channels', 'Update index...', and a search input field. The main area displays a table of installed packages with columns for Name, Description, and Version. Each row includes a green checkmark in the Name column and a green circle icon in the Description column. The Version column shows the current version, with some entries having a blue arrow icon indicating an update is available.

Name	Description	Version
<input checked="" type="checkbox"/> _ipyw_jlab_nb_ex...	<input type="radio"/> A configuration metapackage for enabling anaconda-bundled jupyter extensions	0.1.0
<input checked="" type="checkbox"/> alabaster	<input type="radio"/> Configurable, python 2+3 compatible sphinx theme.	0.7.12
<input checked="" type="checkbox"/> asn1crypto	<input type="radio"/> Python asn.1 library with a focus on performance and a pythonic api	↗ 0.24.0
<input checked="" type="checkbox"/> astroid	<input type="radio"/> A abstract syntax tree for python with inference support.	↗ 2.2.5
<input checked="" type="checkbox"/> astropy	<input type="radio"/> Community-developed python library for astronomy	↗ 3.2.1
<input checked="" type="checkbox"/> attrs	<input type="radio"/> Attrs is the python package that will bring back the joy of writing classes by relieving you from the drudgery of implementing object protocols (aka dunder methods).	↗ 19.1.0
<input checked="" type="checkbox"/> babel	<input type="radio"/> Utilities to internationalize and localize python applications	2.7.0
<input checked="" type="checkbox"/> backports.functoo...	<input type="radio"/> Backport of functools.lru_cache from python 3.3 as published at activestate.	1.5
<input checked="" type="checkbox"/> backports.os	<input type="radio"/> Backport of new features in python's os module	0.1.1
<input checked="" type="checkbox"/> backports.shutil_g...	<input type="radio"/> A backport of the get_terminal_size function from python 3.3's shutil.	1.0.0
<input checked="" type="checkbox"/> backports.weakref	<input type="radio"/> Backport of new features in python's weakref module	1.0.post1

Fig: 6 (Installation of Packages)

4.5 Analysis:

The image shows the Spyder Python IDE interface. The left pane displays a Python script named 'heart disease detction.py' with the following code:

```
5 """
6 import numpy as np
7 import pandas as pd
8 import matplotlib.pyplot as plt
9 from matplotlib import rcParams
10 from matplotlib.cm import rainbow
11 import warnings
12 warnings.filterwarnings('ignore')
13 from sklearn.neighbors import KNeighborsClassifier
14 from sklearn.tree import DecisionTreeClassifier
15 from sklearn.ensemble import RandomForestClassifier
16 from sklearn.svm import SVC
17 df = pd.read_csv('C:\Users\Akshit\Documents\5th sem\python\prog\heart disease detction.csv')
18 df
19 df.describe()
20 import seaborn as sns
21 #get correlations of each features in dataset
22 corrmat = df.corr()
23 top_corr_features = corrmat.index
24 plt.figure(figsize=(10,10))
25 #plot heat map
26 g=sns.heatmap(df[top_corr_features].corr(),annot=True,cmap="RdYlGn")
27 df.hist()
28
29 sns.set_style('whitegrid')
30 sns.countplot(x='target',data=df,palette='RdBu_r')
31 dataset = pd.get_dummies(df, columns = ['sex', 'cp', 'fbs', 'restecg', 'exang', 'slope', 'ca',
32 plt.figure(figsize=(10,10))
33
34 from sklearn.model_selection import train_test_split
35 from sklearn.preprocessing import StandardScaler
36 standardScaler = StandardScaler()
37 columns_to_scale = ['age', 'trestops', 'chol', 'thalach', 'oldpeak']
38 dataset[columns_to_scale] = standardScaler.fit_transform(dataset[columns_to_scale])
39 dataset.head()
40 y = dataset['target'] # target attribute
41 X = dataset.drop(['target'], axis = 1) #data wrangling
```

The right pane shows the IPython console output, which includes the execution of the script and the resulting accuracy scores for different kernels:

```
...: score=cross_val_score(knn_classifier,X,y,cv=10)
...: score.mean()
...:
...: # Naive Bayes Classifier
...: from sklearn.naive_bayes import GaussianNB
...: nb_classifier=GaussianNB()
...: nb_classifier.fit(X_train,y_train)
...: y_pred_nb=nb_classifier.predict(X_test)
...: accuracy_score(y_test,y_pred_nb)
...:
...: # Random Forest
...: from sklearn.ensemble import RandomForestClassifier
...: randomforest_classifier= RandomForestClassifier(n_estimators=10)
...: score=cross_val_score(randomforest_classifier,X,y,cv=10)
...: score.mean()
...:
...: # Decision Tree classifier
...: from sklearn.tree import DecisionTreeClassifier
...: dt_classifier=DecisionTreeClassifier(criterion='entropy',random_state=51)
...: dt_classifier.fit(X_train,y_train)
...: y_pred_dt=dt_classifier.predict(X_test)
...: accuracy_score(y_test,y_pred_dt)
...:
...: # Support Vector Machine
...: svc_scores = []
...: kernels = ['linear', 'poly', 'rbf', 'sigmoid']
...: for i in range(len(kernels)):
...:     svc_classifier = SVC(kernel = kernels[i])
...:     svc_classifier.fit(X_train, y_train)
...:     svc_scores.append(svc_classifier.score(X_test, y_test))
...:
...:
...: colors = rainbow(np.linspace(0, 1, len(kernels)))
...: plt.bar(kernels, svc_scores, color = colors)
...: for i in range(len(kernels)):
...:     plt.text(i, svc_scores[i], svc_scores[i])
...:
...:
...: plt.xlabel('Kernels')
...: plt.ylabel('Scores')
...: plt.title('Support Vector Classifier scores for different kernels')
...: accuracy_score(y_test,y_pred_dt)
...: score.mean()
Out[1]: 0.818197997753059
```

Fig: 7 (Result Analysis)

CONCLUSION

5.1 Advantages:

- User can search for doctor's help at any point.
- User can talk about their heart disease and get instant Diagnosis
- It helps to make the treatment process fast and time consuming

5.2 Disadvantages:

- The system is not fully automated, it needs data from user for fully diagnosis

5.3 Future Enhancement:

The Future of this project is very wide and at enhancement level we can also apply deep learning Neural Network and relate this concept with Image detection means through this model we can pass the images of our X-Ray scan which can detect the level of heart size if it increase or either decrease then it will definitely change the chances of heart disease and the related work in this field is also done by various eminent Professors of different NIT as well as IIT. 1. Dr. Deepak Ranjan Nayak (Currently working as an Assistant Professor in SVNIT Surat) 2. Dr. Kamal Jain (Recently submitted a report on COVID-19 Detection Software)

So, considering the appreciable work of above professor we can also apply the different model of neural networks.

5.4 **Result:**

Models	Accuracy
Logistic Regression	0.9180327868852459
K-Nearest Neighbor	0.8506637004078605
Naïve Bayes	0.8360655737704918
Decision Tree	0.8032786885245902
Random Forest	0.8049721913236929

Tab: 2 (Accuracy Result of Models) Since the logistic regression gives higher accuracy in this case so logistic regression algorithm in this dataset best depicts the person is suffering from heart disease or not basically there is an another way i.e cross validation through which we can also shows which algorithm is best.

5.5 **Conclusion:**

Knowing the heart information will help in long term saving of human lives and early detection of abnormalities in heart conditions. Heart disease prediction is very important in the sector of medical. The Heart diseases can be easily controlled if it is detected at early stage.

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CODE

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from matplotlib import rcParams
from matplotlib.cm import rainbow
import warnings
warnings.filterwarnings('ignore')
from sklearn.neighbors import KNeighborsClassifier
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
df = pd.read_csv('C:\\Users\\Akshit\\Documents\\5th sem\\python\\dataset.csv')
df.describe()

import seaborn as sns
# get correlations of each features in dataset
corrmat = df.corr()
top_corr_features = corrmat.index
plt.figure(figsize=(20,20))
# plot heatmap
mapg = sns.heatmap(df[top_corr_features].corr(), annot=True, cmap="RdYlGn")
df.hist()
sns.set_style('whitegrid')
sns.countplot(x='target', data=df, palette='RdBu_r')
dataset = pd.get_dummies(df, columns = ['sex', 'cp', 'fbs', 'restecg', 'exang', 'slope', 'ca', 'thal'])
plt.figure(figsize=(20,20))

from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
standardScaler = StandardScaler()
columns_to_scale = ['age', 'trestbps', 'chol', 'thalach', 'oldpeak']
dataset[columns_to_scale] = standardScaler.fit_transform(dataset[columns_to_scale])
dataset.head()
y = dataset['target']
X = dataset.drop(['target'], axis = 1) #data wrangling

from sklearn.model_selection import cross_val_score
knn_scores = []
for k in range(1,21):
    knn_classifier = KNeighborsClassifier(n_neighbors = k)
    score = cross_val_score(knn_classifier, X, y, cv=10)
    knn_scores.append(score.mean())
plt.plot([k for k in range(1, 21)], knn_scores, color = 'red')
for i in range(1,21):
    plt.text(i, knn_scores[i-1], (i, knn_scores[i-1]))
plt.xticks([i for i in range(1, 21)])
plt.xlabel('Number of Neighbors (K)')
plt.ylabel('Scores')
plt.title('K Neighbors Classifier scores for different K values')
knn_classifier = KNeighborsClassifier(n_neighbors = 12)
score = cross_val_score(knn_classifier, X, y, cv=10)
score.mean()
```

```
from sklearn.ensemble import RandomForestClassifier
randomforest_classifier=
RandomForestClassifier(n_estimators=10)
score=cross_val_score(randomforest_classifier,X,y,cv
=10)
score.mean()
```

```
from sklearn.model_selection import
train_test_split
X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.2,random_state=5)
X_trainy_trainy_test
```

```
# feature scaling
```

```
from sklearn.preprocessing import
StandardScaler
sc=StandardScaler()
X_train_sc=sc.fit_transform(X_train)
X_test_sc=sc.transform
(X_test)
from sklearn.metrics import accuracy_score
```

```
###logistic regression
```

```
from sklearn.linear_model import
LogisticRegression
lr_classifier=LogisticRegression(random_state=51,penalty='l1')
lr_classifier.fit(X_train,y_train)
y_pred_lr=lr_classifier.predict(X_test)
accuracy_score(y_test,y_pred_lr)
```

```
# Naive Bayes Classifier
```

```
from sklearn.naive_bayes import
GaussianNB
nb_classifier=GaussianNB()
nb_classifier.fit(X_train,y_train)
y_pred_nb=nb_classifier.predict(X_test)
accuracy_score(y_test,y_pred_nb)
```

```
# Decision Tree classifier
```

```
from sklearn.tree import
DecisionTreeClassifier
dt_classifier=DecisionTreeClassifier(criterion='entropy',random_state=51)
dt_classifier.fit(X_train,y_train)
y_pred_dt=dt_classifier.predict(X_test)
accuracy_score(y_test,y_pred_dt)
```

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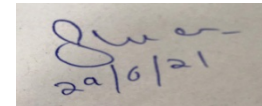
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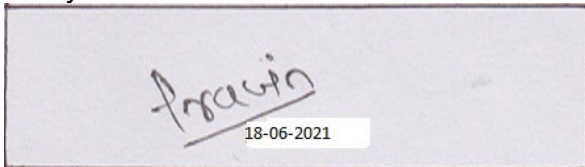
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