

SRM INSTITUTE OF SCIENCE AND

TECHNOLOGY



SCHOOL OF COMPUTING

DEPARTMENT OF DATASCIENCE AND BUSINESS

SYSTEMS

18CSC305J ARTIFICIAL INTELLIGENCE

MINI PROJECT REPORT

Title

Name: Snehal Kumar Ketala

Register Number: RAI911027010104

Mail ID: kk3929@srmistedu.in

Department: CSE

Specialization: Big Data Analytics

Semester: VIth

Team Members

Name: Geetaanjali GNS Name: Kolisetty Yeshwanth Kumar Name: Gollapudi Vyshnavi Registration Number RA1911027010113
Registration Number RA1911027010111
Registration Number RA1911027010112

Content Page

Abstract

Chapter 1: Introduction and Motivation [Purpose of the problem statement (societal benefit)

Chapter 2: Review of Existing methods and their Limitations

Chapter 3: Proposed Method with System Architecture / Flow Diagram

Chapter 4: Modules Description

Chapter 5: Implementation requirements

Chapter 6: Output Screenshots

Conclusion

References

Appendix A – Source Code

Appendix B – GitHub Profile and Link for the Project

Chapter 01, Introduction and Motivation

- Introduction: The project involved analysis of the heart disease patient dataset with proper data processing. Then, different models were trained and and predictions are made with different algorithms KNN, Decision Tree, Random Forest, SVM, Logistic Regression etc This is the jupyter notebook code and dataset I've used for my Kaggle kernel 'Binary Classification with Sklearn and Keras'
 - We've used a variety of Machine Learning algorithms, implemented in Python, to predict the presence of heart disease in a patient. This is a classification problem, with input features as a variety of parameters, and the target variable as a binary variable, predicting whether heart disease is present or not.
- Motivation: preventing Heart diseases has become more than necessary. Good data-driven systems for predicting heart diseases can improve the entire research and prevention process, making sure that more people can live healthy lives. This is where Machine Learning comes into play. Machine Learning helps in predicting Heart diseases, and the predictions made are quite accurate.

Chapter 02, Review of Existing methods and their Limitations.

- Methods:
 - 1. Logistic Regression
 - 2. Naive Bayes
 - 3. SVM
 - 4. NNA
 - 5. Decision Tree
 - 6. Random Forest
 - 7. Neural Network
 - 8. XGBoost

Limitations

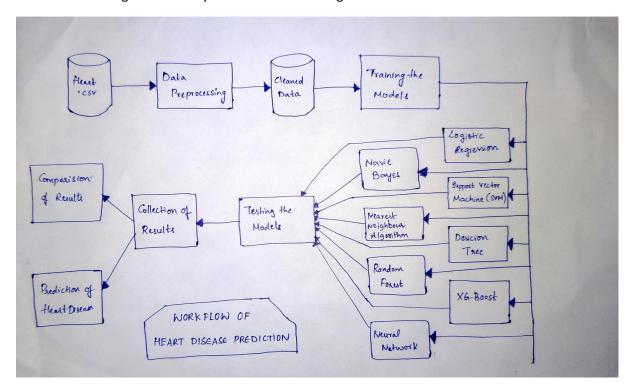
- 1. Logistic Regression: assumption of linearity between the dependent variable and the independent variables
- 2. Naive Bayes: assumption of independent predictors. In real life, it is almost impossible that we get a set of predictors which are completely independent.
- 3. SVM: does not perform very well when the data set has more noise i.e. target classes are overlapping
- 4. NNA:
- 5. Decision Tree: They are unstable, meaning that a small change in the data can lead to a large change in the structure of the optimal decision tree
- 6. Neural Network: Complex to Perform
- 7. XGBoost:

Chapter 03, Proposed Method with System Architecture / Flow Diagram.

Method:

1. Collected the dataset named heart.csv from Kaggle to implement our M.L. algorithms.

- 2. Imported the dataset into google colab and then understood the data and the columns.
- 3. performed the exploratory data analysis on target variables and independent variables.
- 4. Cleaned the dataset.
- 5. Split the data for training and testing.
- 6. Trained the train dataset with models, namely, logistic regression, Naive Bayesian, Support Vector Machine, KNN, Random Forest, XG-boost, Neural Networks.
- 7. Tested the models on the train dataset.
- 8. Got the highest accuracy for Random Forest algorithm.



Chapter 04, Modules Description.

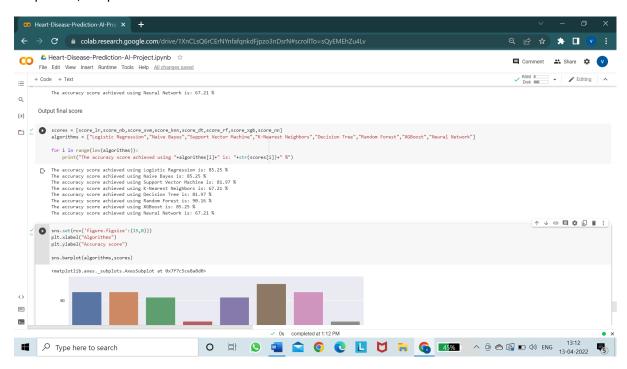
- Logistic Regression (Scikit-learn)
- Naive Bayes (Scikit-learn)
- Support Vector Machine (Linear) (Scikit-learn)
- K-Nearest Neighbours (Scikit-learn)
- Decision Tree (Scikit-learn)
- Random Forest (Scikit-learn)
- XGBoost (Scikit-learn)
- Artificial Neural Network with 1 Hidden layer (Keras)
- Accuracy achieved: 95% (Random Forest)

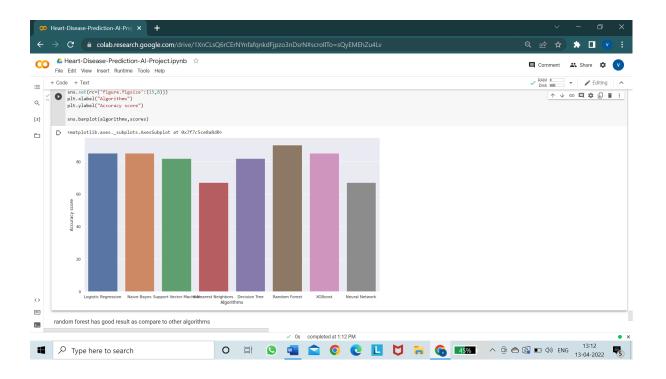
Chapter 05, Implementation Requirements.

- Google Collaboratory
- Heart.csv dataset

- Python Libraries
 - 1. Numpy
 - 2. Pandas
 - 3. Matplotlib
 - 4. seaborn
 - 5. OS
 - 6. warnings
 - 7. sklearn.metrics
 - 8. sklearn.linear_models
 - 9. sklearn.naive_bayes
 - 10. sklearn.neighbors
 - 11. sklearn.tree
 - 12. sklearn.ensemble
 - 13. XGBoost
 - 14. Keras.models
 - 15. Keras.layers

Chapter 06, Output Screenshots.





Conclusion

On Implementing multiple models per say, Logistic Regression (Scikit-learn), Naive Bayes (Scikit-learn), Support Vector Machine (Linear) (Scikit-learn), K-Nearest Neighbours (Scikit-learn), Decision Tree (Scikit-learn), Random Forest (Scikit-learn), XGBoost (Scikit-learn), Artificial Neural Network with 1 Hidden layer (Keras), Accuracy achieved: 95% (Random Forest), We found that Random Forest is the best algorithm to predict if a person has a Heart Disease.

References:

https://www.javatpoint.com/logistic-regression-in-machine-learning

https://www.upgrad.com/blog/naive-bayes-explained/

https://www.javatpoint.com/machine-learning-support-vector-machine-algorithm

https://www.geeksforgeeks.org/k-nearest-neighbours/

https://en.wikipedia.org/wiki/Decision_tree

https://www.ibm.com/cloud/learn/random-forest

https://www.geeksforgeeks.org/xgboost/

Appendix A – Source Code

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
import os
print(os.listdir())
import warnings
warnings.filterwarnings('ignore')
dataset = pd.read csv('heart.csv')
type (dataset)
dataset.shape
dataset.head(5)
dataset.sample(5)
dataset.describe()
dataset.info()
info = ["age","1: male, 0: female","chest pain type, 1: typical
angina, 2: atypical angina, 3: non-anginal pain, 4:
asymptomatic", "resting blood pressure", " serum cholestoral in
mg/dl", "fasting blood sugar > 120 mg/dl", "resting
electrocardiographic results (values 0,1,2)"," maximum heart rate
achieved", "exercise induced angina", "oldpeak = ST depression
induced by exercise relative to rest", "the slope of the peak
exercise ST segment", "number of major vessels (0-3) colored by
flourosopy","thal: 3 = normal; 6 = fixed defect; 7 = reversable
defect"]
```

```
for i in range(len(info)):
    print (dataset.columns[i]+":\t\t\t"+info[i])
dataset["target"].describe()
dataset["target"].unique()
print(dataset.corr()["target"].abs().sort values(ascending=False)
y = dataset["target"]
sns.countplot(y)
target temp = dataset.target.value counts()
print(target temp)
print("Percentage of patience without heart problems:
"+str(round(target temp[0]*100/303,2)))
print("Percentage of patience with heart problems:
"+str(round(target temp[1]*100/303,2)))
dataset["sex"].unique()
sns.barplot(dataset["sex"],y)
dataset["cp"].unique()
sns.barplot(dataset["cp"],y)
dataset["fbs"].describe()
dataset["fbs"].unique()
sns.barplot(dataset["fbs"],y)
dataset["restecg"].unique()
sns.barplot(dataset["restecg"],y)
dataset["exang"].unique()
sns.barplot(dataset["exang"],y)
```

```
dataset["slope"].unique()
dataset["ca"].unique()
sns.countplot(dataset["ca"])
sns.barplot(dataset["ca"],y)
dataset["thal"].unique()
sns.barplot(dataset["thal"],y)
sns.distplot(dataset["thal"])
from sklearn.model selection import train test split
predictors = dataset.drop("target",axis=1)
target = dataset["target"]
X_train, X_test, Y_train, Y_test =
train test split(predictors, target, test size=0.20, random state=0)
X train.shape
X test.shape
Y train.shape
Y test.shape
from sklearn.metrics import accuracy score
fromklearn.linear model s import LogisticRegression
lr = LogisticRegression()
lr.fit(X train, Y train)
Y pred lr = lr.predict(X test)
Y_pred_lr.shape
score_lr = round(accuracy_score(Y_pred_lr,Y_test)*100,2)
```

```
print("The accuracy score achieved using Logistic Regression is:
"+str(score_lr)+" %")
from sklearn.naive bayes import GaussianNB
nb = GaussianNB()
nb.fit(X_train,Y_train)
Y pred nb = nb.predict(X test)
Y pred nb.shape
score nb = round(accuracy score(Y pred nb,Y test)*100,2)
print ("The accuracy score achieved using Naive Bayes is:
"+str(score nb)+" %")
from sklearn import svm
sv = svm.SVC(kernel='linear')
sv.fit(X train, Y train)
Y_pred_svm = sv.predict(X_test)
Y_pred_svm.shape
score svm = round(accuracy score(Y pred svm, Y test)*100,2)
print("The accuracy score achieved using Linear SVM is:
"+str(score svm)+" %")
from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n neighbors=7)
```

```
knn.fit(X_train,Y_train)
Y pred knn=knn.predict(X test)
score knn = round(accuracy score(Y pred knn,Y test)*100,2)
print("The accuracy score achieved using KNN is:
"+str(score knn)+" %")
from sklearn.tree import DecisionTreeClassifier
\max \ accuracy = 0
for x in range (200):
    dt = DecisionTreeClassifier(random state=x)
    dt.fit(X train, Y train)
    Y pred dt = dt.predict(X test)
    current accuracy =
round(accuracy_score(Y_pred_dt,Y_test)*100,2)
    if(current_accuracy>max_accuracy):
        max accuracy = current accuracy
        best x = x
#print(max accuracy)
#print(best_x)
dt = DecisionTreeClassifier(random state=best x)
dt.fit(X train, Y train)
Y_pred_dt = dt.predict(X_test)
print(Y pred dt.shape)
score dt = round(accuracy score(Y pred dt, Y test) *100,2)
```

```
print("The accuracy score achieved using Decision Tree is:
"+str(score_dt)+" %")
from sklearn.ensemble import RandomForestClassifier
\max \ accuracy = 0
for x in range (2000):
    rf = RandomForestClassifier(random state=x)
    rf.fit(X train, Y train)
    Y pred rf = rf.predict(X test)
    current accuracy =
round(accuracy score(Y pred rf,Y test)*100,2)
    if(current accuracy>max accuracy):
        max _accuracy = current_accuracy
        best x = x
#print(max accuracy)
#print(best x)
rf = RandomForestClassifier(random state=best x)
rf.fit(X_train,Y_train)
Y pred rf = rf.predict(X test)
Y pred rf.shape
score_rf = round(accuracy_score(Y_pred_rf,Y_test)*100,2)
print("The accuracy score achieved using Decision Tree is:
"+str(score rf)+" %")
import xgboost as xgb
```

```
xgb model = xgb.XGBClassifier(objective="binary:logistic",
random state=42)
xgb model.fit(X train, Y train)
Y pred xgb = xgb model.predict(X test)
Y pred xgb.shape
score_xgb = round(accuracy_score(Y_pred_xgb,Y_test)*100,2)
print("The accuracy score achieved using XGBoost is:
"+str(score xgb)+" %")
from keras.models import Sequential
from keras.layers import Dense
model = Sequential()
model.add(Dense(11,activation='relu',input dim=13))
model.add(Dense(1,activation='sigmoid'))
model.compile(loss='binary_crossentropy',optimizer='adam',metrics
=['accuracy'])
Y pred nn = model.predict(X test)
Y_pred_nn.shape
rounded = [round(x[0]) for x in Y_pred_nn]
Y pred nn = rounded
score nn = round(accuracy score(Y pred nn,Y test)*100,2)
print ("The accuracy score achieved using Neural Network is:
"+str(score nn)+" %")
```

```
scores =
[score_lr,score_nb,score_svm,score_knn,score_dt,score_rf,score_xg
b,score_nn]
algorithms = ["Logistic Regression","Naive Bayes","Support Vector
Machine","K-Nearest Neighbors","Decision Tree","Random
Forest","XGBoost","Neural Network"]

for i in range(len(algorithms)):
    print("The accuracy score achieved using "+algorithms[i]+"
is: "+str(scores[i])+" %")
sns.set(rc={'figure.figsize':(15,8)})
plt.xlabel("Algorithms")
plt.ylabel("Accuracy score")
sns.barplot(algorithms,scores)
```

Appendix B – GitHub Profile and Link for the Project

https://github.com/Gollapudi-vyshnavi1104/Heart-Disease-Prediction - Vyshnavi

https://github.com/GeetaanjaliGNS/8086-add-two-arrays/blob/main/Heart_Disease_Prediction_Al_Project.ipynb -Geetaanjali

https://github.com/snehalketala/Heart_Disease_Prediction_Al_Project - Snehal Kumar Ketala