	Maternal Health Risk Data Set FINAL PROJECT ON MACHINE LEARNING MODEL PROGRAMMING IN PYTHON Project Overview
	The main purpose of doing this project was to get a proper idea about the machine learning and the model in machine learning. This project mainly focuses on the data set of Maternal health risk. Firstly, diffrent types of python libraries are used for a good output. while working with the datasets a target was selected to focus and predict accuracy. The models used for the accuracy of the dataset can become a important asset to understand the maternal health risk in the rural areas. Lastly, it can be said that the project's main target was to increase knowledge about diffrent algorithms and outcomes. #importing necessary python libraries import numpy as np import pandas as pd
	<pre>import matplotlib.pyplot as plt import seaborn as sns from sklearn import metrics %matplotlib inline Dataset reference https://archive.ics.uci.edu/ml/datasets/Maternal+Health+Risk+Data+Set</pre>
	Total Dataset Information The data was colleted from Maternal Health Risk Data Set . Data has been collected from different hospitals, community clinics, maternal health cares from the rural areas of Bangladesh through the IoT based risk monitoring system. The Age, Systolic Blood Pressure as SystolicBP, Diastolic BP as DiastolicBP, Blood Sugar as BS, Body Temperature as BodyTemp, HeartRate and RiskLevel. All these are the responsible and significant risk factors for maternal mortality, that is one of the main concern of SDG of UN. Attribute information: 1. AGE: Age-Any ages in years when a women during pregnant. 2.SBP: SystolicBP-Upper value of Blood Pressure in mmHg, another significant attribute during pregnancy. 3.DBP: DiastolicBP-Lower value of Blood Pressure in mmHg, another significant attribute during pregnancy. 4.BS: Blood glucose levels is in terms of a molar concentration, mmol/L. 5.BTM:
Tn [2].	BodyTemprature troughout the pregnancy. 6.HRT: HeartRate-A normal resting heart rate in beats per minute. 7.RLV: Risk Level: Predicted Risk Intensity Level during pregnancy considering the previous attribute. Relevant Papers: 1. Ahmed M., Kashem M.A., Rahman M., Khatun S. (2020) Review and Analysis of Risk Factor of Maternal Health in Remote Area Using the Internet of Things (IoT). In: Kasruddin Nasir A. et al. (eds) InECCE2019. Lecture Notes in Electrical Engineering, vol 632. Springer, Singapore. [Web Link] 2. IoT based Risk Level Prediction Model for Maternal Health Care in the Context of Bangladesh, STI-2020, [under publication in IEEE]
In [2]: Out[2]:	#importing the datasets from csv file in a dataframe and showing dF=pd.read_csv('r_dataset.csv') dF AGE SBP DBP BS BTM HRT RLV 0 25 130 80 15.0 98.0 86 high risk 1 35 140 90 13.0 98.0 70 high risk 2 29 90 70 8.0 100.0 80 high risk
	3 30 140 85 7.0 98.0 70 high risk 4 35 120 60 6.1 98.0 76 low risk 1009 22 120 60 15.0 98.0 80 high risk 1011 35 85 60 19.0 98.0 86 high risk 1012 43 120 90 18.0 98.0 70 high risk
	1013 32 120 65 6.0 101.0 76 mid risk 1014 rows × 7 columns Data Preprocessing Data preprocessing is one of the most important task for making a machine learing model. Because, with dirty data it most probability will generate inaccurate outputs. There are several techniques of data cleaning like checking the null value, checking if one of the columns of data set has most empty values. In our data set the gone through a cleaning process to make
In [3]:	#Exploring the Details of the dataset dF.info() <class 'pandas.core.frame.dataframe'=""> RangeIndex: 1014 entries, 0 to 1013 Data columns (total 7 columns): # Column Non-Null Count Dtype</class>
In [4]:	1 SBP 1014 non-null int64 2 DBP 1014 non-null int64 3 BS 1014 non-null float64 4 BTM 1014 non-null float64 5 HRT 1014 non-null int64 6 RLV 1014 non-null object dtypes: float64(2), int64(4), object(1) memory usage: 55.6+ KB #Replace null values with 'null'
Out[4]:	new_dF = dF.fillna('null') AGE SBP DBP BS BTM HRT RLV 0 25 130 80 15.0 98.0 86 high risk 1 35 140 90 13.0 98.0 70 high risk 2 29 90 70 8.0 100.0 80 high risk 3 30 140 85 7.0 98.0 70 high risk 4 35 120 60 6.1 98.0 76 low risk
	III
In [5]:	<pre>new_dF.info() <class 'pandas.core.frame.dataframe'=""> RangeIndex: 1014 entries, 0 to 1013 Data columns (total 7 columns): # Column Non-Null Count Dtype 0 AGE 1014 non-null int64</class></pre>
	1 SBP 1014 non-null int64 2 DBP 1014 non-null int64 3 BS 1014 non-null float64 4 BTM 1014 non-null float64 5 HRT 1014 non-null int64 6 RLV 1014 non-null object dtypes: float64(2), int64(4), object(1) memory usage: 55.6+ KB Exploratory Data Analysis
	Exploritory Data analysis is also one of the important part for machine learning model creation. Because with the help of data analysis we can get an initial idea of the prediction of the dataset and chose a target. In the selected dataset the risk level was selected as target. Diffrent plotting with bar grap, scatter graph were discribe the co-relation between the target variable to form a relation with the variables. Overall, the data analysis part helped us a little bit to choose the feature matrix for accuracy models. #Returning All the keys from the dataset new_dF.keys() Index(['AGE', 'SBP', 'DBP', 'BS', 'BTM', 'HRT', 'RLV'], dtype='object')
<pre>In [7]: Out[7]:</pre>	#Replacing the Target Column True/False value with high/low/mild blood suger levels information new_dF['RLV']=new_dF['RLV'].replace(['high risk','mid risk','low risk'],['True','True or False','False']) new_dF AGE SBP DBP BS BTM HRT RLV 1 35 140 90 13.0 98.0 86 True 2 29 90 70 8.0 100.0 80 True
	3 30 140 85 7.0 98.0 70 True 4 35 120 60 6.1 98.0 76 False 1009 22 120 60 15.0 98.0 80 True 1010 55 120 90 18.0 98.0 60 True 1011 35 85 60 19.0 98.0 86 True 1012 43 120 90 18.0 98.0 70 True
	1013 32 120 65 6.0 101.0 76 True or False 1014 rows × 7 columns The risk factors are determined by BS(blood glucose) and changing high risk, low risk and mild risk to a true, false output, in maternal health cares from the rural areas of Bangladesh. # showing the number of woman at diffrent risk levels # data distribution of the target variable(RLV) plt.figure(figsize=(10,5))
Out[8]:	sns.countplot(x='RLV', data=dF, palette='rainbow') plt.title('Number of women at maternal health risk') Text(0.5, 1.0, 'Number of women at maternal health risk') Number of women at maternal health risk 400- 350-
	300 - 250 - 150 - 100 - 50 -
	Data exploration Models The target varible which is risk level and the corelation between the age and blood glucouse were determined by diffrent graps to understand co-relations. #plotting the Data of the blood glucouse levels and the level of health risk
Out[9]:	sns.set_style("white") plt.figure(figsize=(10,5)) sns.stripplot(x='RLV', y='BS', data=dF) plt.title("Swarmplot on diffrent glucouse level and health risk") Text(0.5, 1.0, 'Swarmplot on diffrent glucouse level and health risk') Swarmplot on diffrent glucouse level and health risk 18
	16 14 8 12 10 8
In [10]: Out[10]:	high risk low risk mid risk #plotting the Data if women with high glucouse level and their risk level. plt.figure(figsize=(10,5)) plt.scatter(dF['AGE'],dF['BS']) plt.title("Scatter plot on the age based on suger levels") Text(0.5, 1.0, 'Scatter plot on the age based on suger levels')
our[10]:	Scatter plot on the age based on suger levels 18 16 14
In [11]:	#plotting data to compare the ages and risk rate
Out[11]:	plt.figure(figsize=(10,5)) plt.scatter (dF['AGE'],dF['RLV']) plt.title("Scatter plot on the age and risk level during maternity") Text(0.5, 1.0, 'Scatter plot on the age and risk level during maternity') Scatter plot on the age and risk level during maternity mid risk **Bosons** **Bosons
	low risk
	Outcome from the plotting Here, thought the ploting of the data based on the risk levels and the glucous level in the blood it can be determined that women with higher blood glucose level have a higher risk than a woman with lower blood glucouse. Then the age and risk level of the womens were taken ito considation for a generating a idea with the help of the plotting. Model Development
	After the initial data preprocessing and analysis also with the creation of feature matrix and separating the target variable it is time for creating the machine learning model. Total 5 algorithms have been used to test the accuracy of this model. They are listed below: naive bayes knn decision tree logistic regression sym
	All of them were used to test the model. #creating the correlation Matrix for the whole Dataset from sklearn.model_selection import train_test_split from sklearn.preprocessing import LabelEncoder from sklearn.metrics import classification_report, confusion_matrix, precision_recall_curve, auc, roc_curve from sklearn.tree import DecisionTreeClassifier, export_graphviz
	<pre>labelencoder=LabelEncoder() #it is used to convert the categorical data into machine readable form for column in dF.columns: dF[column] = labelencoder.fit_transform(dF[column]) plt.figure(figsize=(10,5)) sns.heatmap(dF.corr(),linewidths=.1,cmap="YlGnBu", annot=True) plt.yticks(rotation=0);</pre>
	AGE 1 0.43 0.41 0.39 -0.27 0.058 -0.22 -0.8 SBP 0.43 1 0.79 0.28 -0.28 -0.017 -0.19 DBP 0.41 0.79 1 0.35 -0.3 -0.056 0.25 -0.4 BS 0.39 0.28 0.35 1 -0.026 0.14 -0.48 -0.2 BTM -0.27 -0.28 -0.3 -0.026 1 0.085 0.0068 -0.0
	RLV 0.058 0.017 0.056 0.14 0.085 1 0.11 1 0.11 AGE SBP DBP BS BTM HRT RLV Total 2 columns have been selected from the dataset based on correlation SBP(Systolic blood pressure)-systolic blood pressure of 160 mm or higher can produce complication in maternal health. so, The women are at higher risk. This column is important
	0 15 10 1 17 13 2 7 7 3 17 11 4 13 2 1009 13 2 1010 13 13
In [15]:	1011 6 2 1012 13 13 1013 13 4 1014 rows × 2 columns #seprating the target variable y=dF['RLV']
Out[15]:	y 0
In [16]:	Name: RLV, Length: 1014, dtype: int64 #using data splitting functions from scikit-learn from sklearn.model_selection import train_test_split X_train, X_test, y_train, y_test = train_test_split(up_df, y, test_size = 0.3, random_state = 16) print("X_training shape: ", X_train.shape) print("X_testing shape: ", X_test.shape) perc = {} X_training shape: (709, 2)
	Traing and Testing Dataset Training- 709 out of 1014 data has been choosen for training the model Testing- 305 out of 1014 data has been choosen for the testing model. ploting where necessary and data spliting
	Data splitting is very important because without it the model cannot be generated. Data needs to be spitted into two parts one is for training the dataset and another part is for testing the dataset. In our project there were total 1014 instances of data. where 709 of 1014 is used for training the model. the rest 305 data were used for testing the model's accuracy. The spitting mostly depends on user as both training and testing is equally important for the model to be accurate. #Naive bayes # importing the necessary package to use the naive bayes algorithm from sklearn.naive_bayes import GaussianNB model_nb = GaussianNB() model_nb.fit(X_train, y_train) y_prediction_nb = model_nb.predict(X_test)
In [18]:	<pre>score_nb = metrics.accuracy_score(y_prediction_nb, y_test).round(4) print('The accuracy of the model based on Naive Bayes Algorithm is {}%'.format(score_nb*100)) perc['Naive-bayes'] = score_nb The accuracy of the model based on Naive Bayes Algorithm is 50.82% #Knn # importing the necessary package to use the knn algorithm from sklearn.neighbors import KNeighborsClassifier model_knn = KNeighborsClassifier(n_neighbors=3) model_knn.fit(X_train, y_train)</pre>
In [19]:	<pre>y_prediction_knn = model_knn.predict(X_test) score_knn = metrics.accuracy_score(y_prediction_knn, y_test).round(4) print('The accuracy of the model based on K Nearest Neighbour Algorithm is {}%'.format(score_knn*100)) perc['Knn'] = score_knn The accuracy of the model based on K Nearest Neighbour Algorithm is 54.75% # importing the necessary package to use the decision tree algorithm from sklearn.tree import DecisionTreeClassifier model_dt = DecisionTreeClassifier(random_state=4) model_dt.fit(X_train, y_train)</pre>
In [20]:	<pre>y_prediction_dt = model_dt.predict(X_test) score_dt = metrics.accuracy_score(y_prediction_dt, y_test).round(4) print('The accuracy of the model based on Decision Tree Algorithm is {}%'.format(score_dt*100)) perc['Decision-tree'] = score_dt The accuracy of the model based on Decision Tree Algorithm is 58.36% #logistic regression algorithm # importing the necessary package to use the logistic regression algorithm from sklearn.linear_model import LogisticRegression model_lr = LogisticRegression()</pre>
In [21]:	<pre>model_lr.fit(X_train, y_train) y_prediction_lr = model_lr.predict(X_test) score_lr = metrics.accuracy_score(y_prediction_lr, y_test).round(4) print('The accuracy of the model based on Logistic Regression Algorithm is {}%'.format(score_lr*100)) perc['Logistic-regression'] = score_lr The accuracy of the model based on Logistic Regression Algorithm is 55.74% #svm # importing the necessary package to use the svm algorithm from sklearn import svm model_svm = svm.SVC()</pre>
In [51]:	<pre>model_svm = svm.SVC() model_svm.fit(X_train, y_train) y_prediction_svm = model_svm.predict(X_test) score_svm = metrics.accuracy_score(y_prediction_svm, y_test).round(4) print('The accuracy of the model based on Support Vector Machine Algorithm is {}%'.format(score_svm*100)) perc['SVM'] = score_svm The accuracy of the model based on Support Vector Machine Algorithm is 58.69% max_accuracy = max(perc.values()) max_model = max(perc, key=perc.get) print(f'The maximum accuracy from all these algorithms is {max_model} with {max_accuracy*100}%')</pre>
In [48]: Out[48]:	The maximum accuracy from all these algorithms is SVM with 58.69% score = pd.DataFrame(list(perc.keys()), columns=['names']) score['accuracy'] = list(perc.values()) score names accuracy Naive-bayes 0.5082
In [50]:	1 Knn 0.5475 2 Decision-tree 0.5836 3 Logistic-regression 0.5574 4 SVM 0.5869 plt.figure(figsize=(10,5)) sns.barplot(x='names', y='accuracy', data=score, palette='rainbow') plt.title('Different model accuracy')
Out[50]:	Text(0.5, 1.0, 'Different model accuracy') Different model accuracy 0.5 0.4
	0.2 0.1 0.0 Naive-bayes Knn Decision-tree names Logistic-regression SVM
	Comparison of Models The 5 models used for the accuracy prediction of the data sets were used to determine the percentage of accuracy of 50.82% based on Support Naive Bayes Algorithm again the higest accuracy percentage of 58.69% based on SVM algorithm, again the second highest decision percentage for the decision tree Algorithm was 58.36%. based on the model accuracy percentage were determined to produce accuracy value by plotting the 5 models and plotting them on a bar graph to understand more clearly that SVM has a little more accuracy from the other 4 models. SVM or Support Vector Machine is a linear model for classification and regression problems for the problem. The reason SVM can be useful for the datasets can solve linear and non-linear problems and work well for a more accurate result.
	Discussion This model is created based on the Maternal Health Risk Data Set which has the data of high, low and mild risk. This model has an accuracy of 50.82%. Which is not the best result or outcome in this data set which has 1014 numbers of instances. This desions where made by diffrent modeles chossing the most relevent data through diffrent plotting and graphs. still as the accuracy is not even near the 70% it might not be an ideal result. which might be the solved if their where more attributes and data sets. If we had more columns which had more correlatation with the other variable the prediction could have been more better. As, it was based on a real life dataset, the prediction can also be called a realistic if it predicts the