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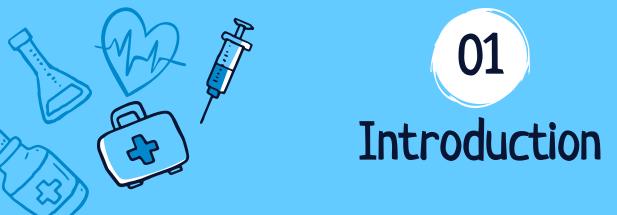
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- Primary End User



Introduction

- The healthcare industry collects a **vast amount** of **confidential data**; we can use advanced data mining techniques to provide relevant results and make effective decisions on data.
- In this project, we will establish a relationship and knowledge between various medical factors related to heart disease and the patterns.
- We will implement various Machine Learning techniques and algorithms to effectively predict heart disease risk.



Primary End Users

- **Insurance Companies:** To analyze insurance premiums and make recommendations.
- Reports & Published Journals: Detailed Analysis and predictions regarding the heart disease data.
- Private & Public Health Department: So that they can closely collaborate with neighborhood health care professionals to stop the spread of the disease and raise required awareness.







Exploratory Data Analytics

- Data Description & Transformation
 - Feature Analysis
- Data Visualization



Data Description & Transformation

- Heart disease is the leading cause of death; about half of Americans have at least one of the three major risk factors for heart disease: high blood pressure, high cholesterol, and smoking. Other indicators include diabetes status, obesity (high BMI), physical inactivity, or excessive alcohol consumption.
- Identifying and preventing the factors that have the most significant impact on heart disease is very important in healthcare.
- In turn, the development of computers allowed machine learning methods to identify "patterns" from data that could predict a patient's condition.



Data Description

The system uses **18 medical parameters** such as Age Category, Sex, Diabetic, Sleep Time, BMI, Smoking, Alcohol, Physical Health, Mental Health, and Race, to name a few.

Categorical / Boolean Variables in the Data:

Heart Disease, Smoking, Alcohol Drinking, Stroke, Difficulty in Walking, Diabetic, Physical Activity, Asthma, Kidney Disease, Skin Cancer.

Numerical Variables in the Data:

Age Category, BMI, Physical Health, Mental Health, Sleep Time.

Strings in the Data:

Race, General Health



Data Transformation

Creation of Dummy Variables:

We have used Pandas using get_dummies.

Handling Null Values:

We have removed all the null values before our analysis and model learning part.

Reducing Dimensionality & Variance using feature analysis:

We have analyzed and removed dimensionality by using correlation.

Removed Outliers for better model learning:

Identified Outliers and released them for better

Identified Outliers and released them for better analytics.



Feature Analysis

Analysis of the Correlation:

The dummy variables made have been used to do the feature analysis and their correlation values are calculated using the corr() function.

[] EDA_HD.corr()																					
	OMT DI	enclosius ith Ma	entellesite d		and the same in the		Carlelina No.	faction Voc. 83	coholDrinking No Alcohol	Dafakiaa Vas	Continuately Endon Co	ulian lith Good (lantiavleh Book Gantie	alth Name and	Acthus No.	lether Voc V	Identificance No. VI.	landicasca Vac El	defense No. 6		
RMI	1,000000	0.109788	0.064131		=0.051803	0.051803		0.023118	0.038816	-0.038816	. 0.127364	0.118047	0.062501		-0.092345		0.050768	0.050768	0.033644	-0.033544	
PhysicalHealth	0.109788	1.000000	0.287987		-0.051803	0.170721	-0.023118 -0.115352	0.115352	0.036616	-0.036616	. 0.127364	-0.037663	0.062501	-0.196462		0.117907	-0.050768	0.000766	-0.041700	0.041700	
MenfalHealth	0.064131	0.287987	1.000000		-0.028591	0.028591	-0.085157	0.085157	-0.051282	0.051282	0.151321	0.013353	0.192079	-0.089956		0.114008	-0.037281	0.037281	0.033412	-0.033412	
SleepTime	-0.051822	-0.061387	-0.119717		-0.008327	0.008327	0.030336	-0.030336	0.005065	0.005065	.0.040923	-0.013725	.0.033074	0.019379		-0.048245	-0.00623B	0.006238	-0.041266	0.041266	
HeartDiceace_No							0.107764				-0.147954		-0.174662	0.101886							
HeartDicease_Yes	0.051803	0.170721	0.028591	0.008327	-1.000000	1.000000	-0.107764	0.107764	0.032080	-0.032080	. 0.147954	0.039033	0.174662	-0.101886	-0.041444	0.041444	-0.145197	0.145197	-0.093317	0.093317	
8moking_No		-0.115352	-0.085157	0.030336			1.000000	-1.000000			-0.095620	-0.059651	-0.086520	0.052305	0.024149	-0.024149	0.034920	-0.034920			
8moking_Yes	0.023118	0.115352	0.085157	-0.030336	-0.107764	0.107764	-1.000000	1.000000	-0.111768	0.111768	. 0.095620	0.059651	0.086520	-0.052305	-0.024149	0.024149	-0.034920	0.034920	-0.033977	0.033977	
AlcoholDrinking_No	0.038816			0.005065	-0.032080	0.032080			1.000000	-1.000000	. 0.018859	0.007808		-0.013005	-0.002202	0.002202	-0.028280	0.028280			
AlooholDrinking_Yes	-0.038816	-0.017254	0.051282	-0.005065	0.032080	-0.032080	-0.111768	0.111768	-1.000000	1.000000	-0.018859	-0.007808	-0.017058	0.013005	0.002202	-0.002202	0.028280	-0.028280	0.005702	-0.005702	
8troke_No			-0.046467		0.196835	-0.196835			-0.019858	0.019858	-0.104983			0.069395	0.038866	-0.038866				-0.048116	
8troke_Yes	0.019733	0.137014	0.046467	0.011900	-0.196835	0.196835	-0.061226	0.061226	0.019858	-0.019858	. 0.104983	0.013159	0.133641	-0.069395	-0.038866	0.038866	-0.091167	0.091167	-0.048116	0.048116	
DiffWalking_No		-0.428373							-0.035328	0.035328				0.184986	0.103222		0.153064	-0.153064	0.064840	-0.064840	
DiffWalking_Yes	0.181678	0.428373	0.152235	-0.022216	-0.201258	0.201258	-0.120074	0.120074	0.035328	-0.035328	. 0.282517	0.031570	0.308767	-0.184986	-0.103222	0.103222	-0.153064	0.153064	-0.064840	0.064840	
8ex_Female	-0.026940	0.040904	0.100058		0.070040	-0.070040	0.085052	-0.085052	0.004200	-0.004200	. 0.022456	-0.003642	0.010667		-0.069191	0.069191	-0.009084	0.009084		-0.013434	
Sex_Male	0.026940	-0.040904	-0.100058		-0.070040	0.070040	-0.085052	0.085052	-0.004200	0.004200	-0.022456	0.003642	-0.010667	-0.003239		-0.069191	0.009084	-0.009084	-0.013434	0.013434	
AgeCategory_18-24		-0.055866	0.075243						-0.004334	0.004334			-0.040685	0.018047				-0.043093		-0.082247	
AgeCategory_25-29	-0.023705	-0.046707	0.054452		0.065759	-0.065759	0.052149	-0.052149	-0.023069	0.023069	-0.037142	-0.017925	-0.032709	0.013041		0.024371	0.037750	-0.037750	0.071893	-0.071893	
AgeCategory_39-34	0.004500	-0.042484			0.065611	-0.065611			-0.015902		-0.034864										
AgeCategory_\$5-\$8	0.021160	-0.037248	0.037929		0.066685	-0.066685	-0.004290	0.004290	-0.021545	0.021545	0.029789	-0.011409	-0.029053	0.006085		0.004643	0.033914	-0.033914	0.072501	-0.072501	
AgeCategory_40-44	0.036475	-0.026575	0.025892		0.059196	-0.059196	-0.010680	0.010680	-0.018818	0.018818	-0.021467	-0.008870			-0.009222		0.027323	-0.027323	0.067055	-0.067055	
AgeCategory_45-49	0.049427	-0.011934 0.008711	0.016566		0.049733	-0.049733 -0.032648	0.006637	-0.006637 -0.011667	-0.009857 -0.010588	0.009857	0.011371 -0.005852	-0.008464 -0.005515	-0.011851 0.004513	0.001805	-0.007782 -0.002352	0.007782	0.023167	-0.023167 -0.014427	0.063725	-0.053725 -0.042876	
AgeCategory_60-64 AgeCategory_66-69	0.038984	0.008/11	0.006345		0.032648	-0.032648	-0.008701	0.008701	-0.010588 -0.008189	0.010588	. 0.010708	-0.008782	0.004613	-0.001180		-0.001461	0.005603	-0.014427	0.042876	-0.042876	
AgeCategory_66-68	0.026797	0.026416	-0.015002		-0.016152	0.016152	-0.008701	0.008/01	-0.006189	0.001621	. 0.024054	0.002738	0.017940	-0.010070		-0.001461	-0.007098	0.007098	-0.006900	0.006900	
AgeCategory_65-89	0.019006	0.021009	-0.010002		-0.016162	0.042626	-0.031892	0.030367	0.009026	-0.009026	. 0.024054	0.002736	0.027401	0.000424		-0.000837	-0.007098	0.007096	-0.049571	0.049571	
AgeCategory_70-74	-0.007720	0.022623	-0.055078		-0.082578	0.082578	-0.045288	0.045288	0.021730	-0.021730	. 0.026606	0.015349	0.021192		0.015297		-0.046293	0.046293	-0.096897	0.096897	
AgeCategory_76-79	-0.030726	0.027203		0.058898	-0.098690	0.098690	-0.048040	0.048040	0.027861	-0.027861	. 0.028147	0.024777	0.021436	-0.007261		-0.017448	-0.053572	0.063572	-0.121747	0.121747	
AgeCategory_80 or older	-0.094780	0.039621	-0.071718		-0.143041	0.143041	-0.013569	0.013569	0.045226	-0.045226	. 0.050048	0.038115	0.037282	-0.027169			-0.067691	0.067691			
Race_American Indian/Alackan Nativ	0.026347	0.022955	0.018394	-0.003615	-0.008547	0.008547	-0.035667	0.035667	0.004243	-0.004243	. 0.022334	0.015839	0.022017	-0.025611		0.013757	-0.007401	0.007401	0.025784	-0.026784	
Race Aslan	-0.078643	-0.035229	-0.023113	-0.019985	0.030262		0.060308	-0.060308	0.022275	-0.022275	-0.024360	0.003752	-0.018024	-0.003102	0.017007	-0.017007	0.016957	-0.016957	0.047749	-0.047749	
[] sns.heatmap(heartdisease.corr())																					

Feature Analysis

In feature analysis, we will study the relationship between variables. The idea is to find interesting relationships that show the influence of one variable on the other, preferably on the target variable (Heart disease).

The most significant variables which have a strong relationship with Heart Disease are:

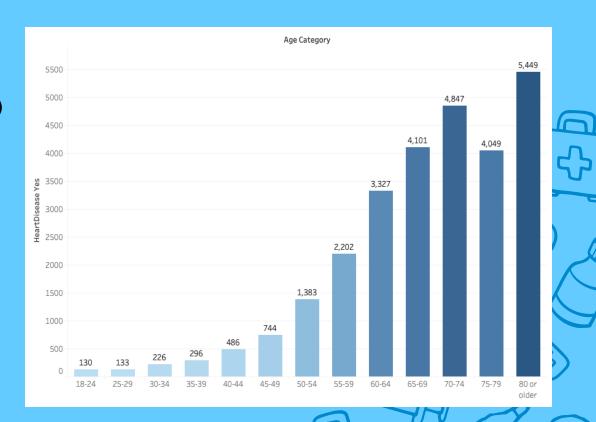
- Stroke Yes
- Difficulty in Walking Yes
- Diabetic Yes
- Physical Health
- General Health Poor
- Kidney Disease Yes

These variables will be used for Model Learning analysis and data visualization purposes.



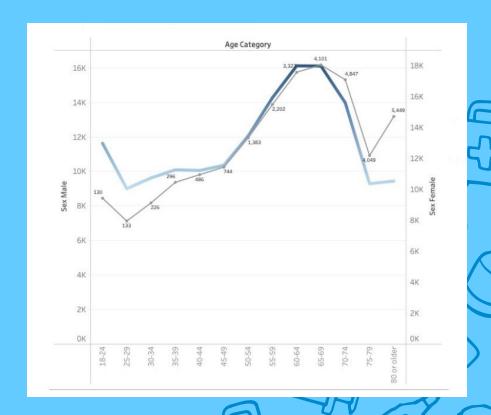
Age category v Heart Disease (Yes)

- Overall, there is an upward trend of heart disease as the age increases
- As age increases, heart disease in people increases accordingly
- However, there is a drop in heart disease at ages 75-79



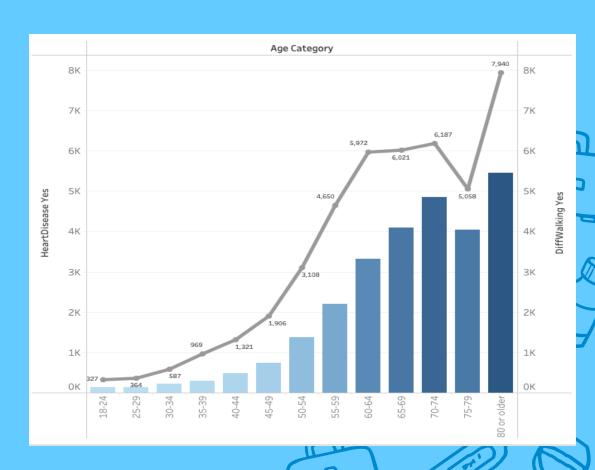
Age Category v Male v Female

- As people get older, heart disease affects both females and males in the same way
- Overall, males have a higher risk of heart disease as compared to females
- However, there is a sudden drop in heart disease in both females and males at ages 25-29 and 75-79



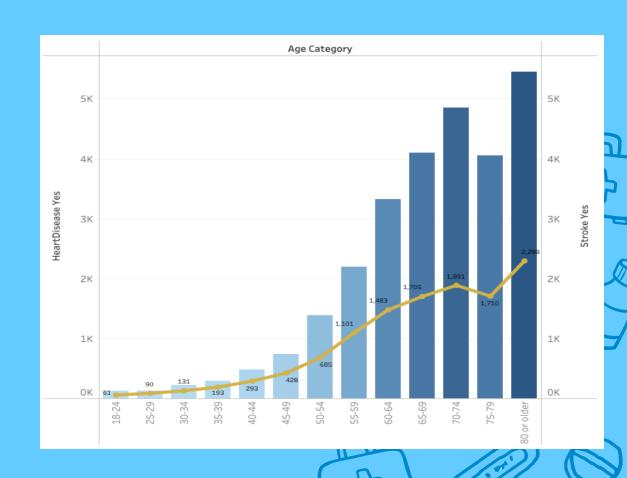
Age category v Heart Disease v Difficulty in walking

- Overall, there is an increasing trend among all the variables
- As the difficulty in walking increases age-wise, the chances of heart disease increases
- However, there is a drop in the number of people with both heart disease and difficulty in walking at ages 75-79.
- Hence, we can say that heart disease and difficulty in walking are going in sync



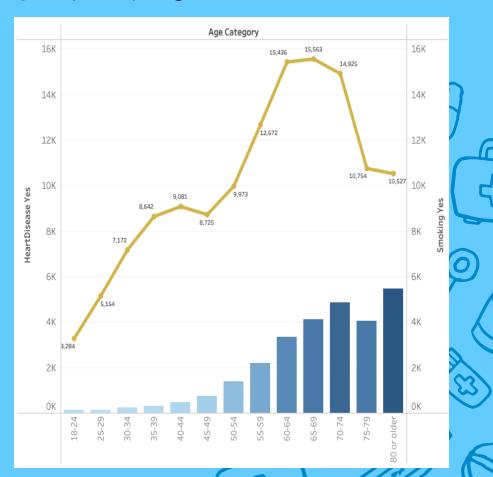
Age category v Heart Disease v Stroke

- Overall, there is an increasing trend among all the variables
- As the strokes increases age-wise, the chances of heart disease increases
- However, there is a drop in the number of people with both heart disease and strokes at ages 75-79.
- Hence, we can say that heart disease and strokes are going in sync



Age Category v Heart Disease v Smoking

- Overall, there is an increasing trend among all the variables
- People who smoke are more in number than people who have heart disease. On average, ¼ people who smoke have heart disease
- Furthermore, there is a sudden drop in heart disease and smoking in people at ages 75-79







- Logistic Regression
 - KNN
 - Decision Tree
 - Random Forest



Heart Disease RMI Smoking Alc

Sample Data

HeartDisease	ВМІ	Smoking	AlcoholDrinking	Stroke	PhysicalHealth	MentalHealth	DiffWalking	Sex	AgeCategory	Race
No	16.6	Yes	No	No	3.0	30.0	No	Female	55-59	White
No	20.34	No	No	Yes	0.0	0.0	No	Female	80 or older	White
No	26.58	Yes	No	No	20.0	30.0	No	Male	65-69	White
No	24.21	No	No	No	0.0	0.0	No	Female	75-79	White
No	23.71	No	No	No	28.0	0.0	Yes	Female	40-44	White

Diabetic	PhysicalActivity	GenHealth	SleepTime	Asthma	KidneyDisease	SkinCancer
Yes	Yes	Very good	5.0	Yes	No	Yes
No	Yes	Very good	7.0	No	No	No
Yes	Yes	Fair	8.0	Yes	No	No
No	No	Good	6.0	No	No	Yes
No	Yes	Very good	8.0	No	No	No



data.info()

Data Insights

<class 'pandas.core.frame.DataFrame'> RangeIndex: 319795 entries, 0 to 319794 Data columns (total 18 columns): Column Non-Null Count Dtype **HeartDisease** 319795 non-null object BMI 319795 non-null float64 Smoking 319795 non-null object AlcoholDrinking 319795 non-null object Stroke 319795 non-null obiect PhysicalHealth 319795 non-null float64 MentalHealth 319795 non-null float64 DiffWalking 319795 non-null object 319795 non-null object Sex **AgeCategory** 319795 non-null object Race 319795 non-null object Diabetic 319795 non-null object 12 PhysicalActivity 319795 non-null object GenHealth 319795 non-null obiect SleepTime 319795 non-null float64 Asthma object 319795 non-null KidnevDisease 319795 non-null object 17 SkinCancer 319795 non-null object

data.shape

Checking null values in data

```
data.isnull().sum()
HeartDisease
BMI
Smoking
AlcoholDrinking
Stroke
PhysicalHealth
MentalHealth
DiffWalking
Sex
AgeCategory
Race
Diabetic
PhysicalActivity
GenHealth
SleepTime
Asthma
KidneyDisease
SkinCancer
```



Data Balancing

For balancing data, We have used SMOTE method to do oversampling the data

```
from imblearn.over_sampling import SMOTENC
smote = SMOTENC([1,2,3,6,7,8,9,10,11,12,14,15,16],random_state = 101)
X_oversample, y_oversample = smote.fit_resample(X, y)
```

```
print('X sample :',X_oversample.shape)
print('Y_sample :',y_oversample.shape)

X sample : (584844, 17)
Y_sample : (584844,)
```

```
y_oversample.value_counts(normalize=True)*100

No 50.0
Yes 50.0
Name: HeartDisease, dtype: float64
```

0



For numeric data

For Categorical data

Full pipeline

Building Pipeline

```
from sklearn.preprocessing import OneHotEncoder, StandardScaler
from sklearn.compose import ColumnTransformer
from sklearn.pipeline import make pipeline, Pipeline
from sklearn.impute import SimpleImputer
cat columns = [i for i in X oversample.columns if data[i].dtype == 'object']
num columns = [i for i in data.columns if data[i].dtype != 'object' and i not in ['BMI']]
num pipe = make pipeline(SimpleImputer(missing values=np.nan,strategy = 'mean'),
    StandardScaler(), SimpleImputer(missing values=np.nan,strategy = 'mean')
cat pipe = make pipeline(SimpleImputer(missing values=np.nan,strategy = 'most frequent'),
    OneHotEncoder(handle unknown='ignore',drop='if binary'),
    SimpleImputer(missing values=np.nan,strategy = 'most frequent')
full pipe = ColumnTransformer([('num', num pipe, num columns),
    ('cat', cat pipe, cat columns)
```



Splitting Data

```
from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test = train_test_split(X_oversample,y_dummy,test_size = 0.4,random_state=2)
X_validation,X_test,y_validation,y_test = train_test_split(X_test,y_test,test_size = 0.5,random_state=2)
```

```
y_train.value_counts(True)*100

0 50.110571

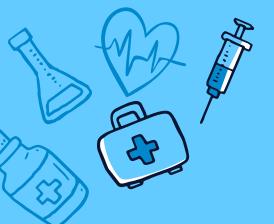
1 49.889429

Name: HeartDisease, dtype: float64
```

We have split the data into 3 parts:

1 Training dataset,2 Validation dataset3 Test dataset





Logistic Regression

- Logistic regression is a supervised classification algorithm.
- Logistic Regression accomplishes binary classification tasks by predicting an outcome, event, or observation probability.
- It uses logistic functions to predict the probability of a binary outcome.
- This classification model delivers a binary outcome limited to two possible effects: yes/no, 0/1, or true/false.



Data Preparation

```
LR_data = X_oversample.copy()
LR_data['HeartDisease'] = y_dummy.copy()
```

Removing Outliers

```
def outliers(features,df):
 Q1 = df[[features]].quantile(q = 0.25)[0]
  Q3 = df[[features]].quantile(q = 0.75)[0]
  iqr = Q3 - Q1
  min iqr = Q1 - 1.5*iqr
  max igr = Q3 + 1.5*igr
 return min iqr,max iqr,df[[features]].min()[0],df[[features]].max()[0]
def convert_nan(x,min_iqr = min_iqr,max_iqr = max_iqr):
    if (x > max_iqr):
      x=np.nan
    else:
      x = x
    return x
for i in ['BMI', 'SleepTime']:
  c=0
 min igr, max igr, min value, max value = outliers(i, LR data)
  if min value < min igr:
    c+=1
    print('Column ->',i)
    print('--->low bound Outliers at',min igr)
  if max_value > max_iqr:
    if c==0:
      print('Column ->',i)
    print('---->uppar bound Outliers at',max iqr)
  print()
  print('Conver outliers to nan\n')
  LR_data[i] = LR_data[i].map(lambda x: np.nan if x<min_iqr or x >max_iqr else x
```



The accuracy of the Logistic Regression model is 76.8%

```
LR = make_pipeline(full_pipe, LogisticRegression(max_iter=10000))
LR.fit(X_train,y_train)
y_lr_predict = LR.predict(X_validation)
print('LogisticRegression accuracy:',(accuracy_score(y_validation,y_lr_predict))*100)
LogisticRegression accuracy: 76.80240063606597
```

Confusion Matrix

	Predicted Yes	Predicted No	c
Actual Yes	43295	11716	
Actual No	14855	47103	



K- Nearest Neighbours

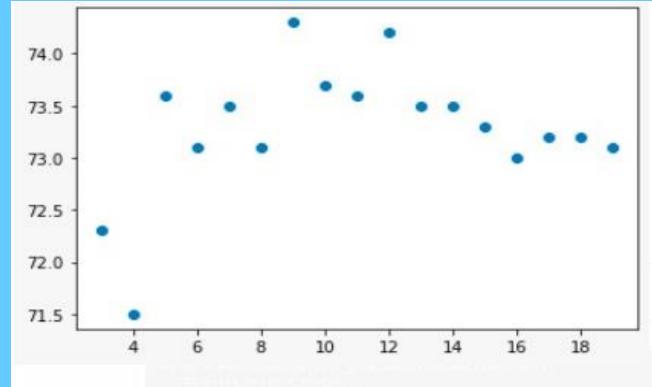
- KNN algorithm is a **supervised machine learning algorithm**.
- It's a classification algorithm that predicts a class of a target variable based on a defined number of nearest neighbors.
- Choosing the number of nearest neighbors, i.e., the value of k, significantly determines the model's efficacy.



Choosing hyper-parameter

We have tried multiple K values to check which K value is the best for the model.

We have selected K value as 9 based on this graph.





Model Evaluation

The accuracy of the KNN model is 74.4%

```
knn = make_pipeline(full_pipe, KNeighborsClassifier(n_neighbors=9))
knn.fit(X_train,y_train)
y_knn_predict = knn.predict(X_validation)
print('knn accuracy:',(accuracy_score(y_validation,y_knn_predict))*100)
knn accuracy: 74.3

predict = knn.predict(X_test)
print('knn accuracy:',(accuracy_score(y_test,predict))*100)
knn accuracy: 74.4
```

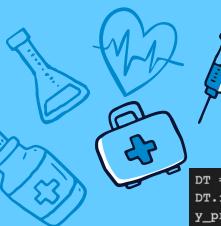
Confusion Matrix

cm(confusion_matrix(y_test,predict))					
	Predicted	Yes	Predicted	No	
Actual Yes		354		92	
Actual No		164	:	390	



Decision Tree

- A decision tree is one of the supervised machine learning algorithms. This algorithm can be used for regression and classification problems
- A decision tree follows a set of if-else conditions to visualize the data and classify it according to the requirements.



Model Evaluation

The accuracy of the Decision Tree is 75.6%

```
DT = make_pipeline(full_pipe,DecisionTreeClassifier())
DT.fit(X_train,y_train)
y_predict = DT.predict(X_test)

print('Decision Tree accuracy:',(accuracy_score(y_test,y_predict))*100)

cm(confusion_matrix(y_test,y_predict))

Decision Tree accuracy: 75.6
```

Confusion Matrix

	Predicted	Yes	Predicted	No
Actual Yes		406		132
Actual No		112	;	350



Random Forest

- Random forest is a Supervised Machine Learning Algorithm that is used widely in Classification and Regression problems.
- The "forest" it builds is an ensemble of decision trees, usually trained with the "bagging" method. The general idea of the bagging method is that a combination of learning models increases the overall result.
 - Random forest adds additional randomness to the model while growing trees. Whenever the tree splits, it only has access to a random sample predictors.



Model Evaluation

The accuracy of the Random Forest model is 77.1%

```
RF = make_pipeline(full_pipe,RandomForestClassifier())
RF.fit(X_train,y_train)
y_predict = RF.predict(X_test)

print('Random Forest Tree accuracy:',(accuracy_score(y_test,y_predict))*1
cm(confusion_matrix(y_test,y_predict))

Random Forest Tree accuracy: 77.1000000000001
```

Confusion Matrix

	Predicted	Yes	Predicted	No	1
Actual Yes		381		92	
Actual No		137		390	





Null Value Test

```
[ ] import random
    null test_data = X_test.copy()
    column_list = null_test_data.columns
    for i in range(1000):
      r_index = random.randint(0,null_test_data.shape[0]-1)
      column = column_list[random.randint(0,len(column_list)-1)]
      null_test_data.loc(null_test_data.index == r_index,column) = np.nan
    null_test_data.isna().sum()
    Smoking
    AlcoholDrinking
    Stroke
    PhysicalHealth
    MentalHealth
    DiffWalking
    Sex
    AgeCategory
    Race
    Diabetic
    PhysicalActivity
    GenHealth
    SleepTime
    Asthma
    KidneyDisease
    SkinCancer
    dtype: int64
    RF.predict(null_test_data)
    array([0, 1, 1, ..., 0, 0, 0])
```





- The most significant variable is Stroke Yes, for the complete classification process.
- We can observe that Males have higher chances of heart disease than females.
- People older than 65 years have a very high chance of heart disease.
- The age group between 25 to 40 who have a smoking habit have a higher chance of having heart disease.
- Random Forrest is the best-performing model, with an accuracy of 77%.

MODE	ACCURACY
Logistic	76.8
KNN	74.4
Decision Tree	75.6
Random Forest	77





Thanks

