	PARKINSON'S DISEASE DETECTION
	Objective:
	To identify whether a person is suffering from Parkinson.s disease or not.  Importing Dependencies
In [1]:	<pre>import numpy as np import pandas as pd from sklearn.model_selection import train_test_split from sklearn.preprocessing import StandardScaler from sklearn import svm from sklearn.metrics import accuracy_score</pre> The Data
In [2]: In [3]:	<pre>parkinsons_data=pd.read_csv(r'C:\Users\HP\Downloads\archive\parkinsons.csv') parkinsons_data.head()</pre>
Out[3]:	name         MDVP:Fo(Hz)         MDVP:Fo(Hz)         MDVP:Flo(Hz)         MDVP:Itter(Mz)         MDVP:RAP         MDVP:RAP         MDVP:PDQ         Jitter:DDP         MDVP:Shimmer          Shimmer:DDA         NHR         HNR         status         RPDE         DFA         spread         spread         spread         policy           0         phon_R01_S01_1         119.992         157.302         74.997         0.00784         0.00007         0.0054         0.01109         0.04374          0.06545         0.0211         21.033         1         0.414783         0.815285         -4.813031         0.21           1         phon_R01_S01_2         122.400         148.650         113.819         0.00968         0.00008         0.00465         0.00696         0.01394         0.06134          0.09403         0.01909         19.085         1         0.458359         0.819521         -4.075192         0.33           2         phon_R01_S01_3         116.682         131.111         111.565         0.01050         0.00099         0.00502         0.00698         0.01505         0.05492          0.08771         0.01353         20.641         1         0.434969         0.819235         -4.117501         0.33
In [5]: Out[5]:	parkinsons_data.shape
In [6]:	This dataframe has 195 rows and 24 columns.  parkinsons. data.info() <class 'pandas.core.frame.dataframe'=""> RangeTodex: 195 cort.ise, 5 to 194  ### Column  #### Column  ### Column  #### Column  ##### Column  ###################################</class>
In [7]:	Getting some statistical insights from the dataset  parkinsons_data.describe()  MDVP:Fo(Hz) MDVP:Fhi(Hz) MDVP:Flo(Hz) MDVP:Jitter(%) MDVP:Jitter(Abs) MDVP:RAP MDVP:PPQ Jitter:DDP MDVP:Shimmer MDVP:Shimmer(dB) Shimmer:DDA NHR HNR status RPDE
Out[7]:	count 195.000000 195.00000 195.00000 195.00000 195.000000 195.000000 195.000000 195.000000 195.000000 195.000000 195.000000 195.00000 195.00000 195.000000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 195.000000 195.00000 195.00000 195.00000 195.00000 195.00000 195.00000 1
In [8]: Out[8]:	parkinsons_data['status'].value_counts()  1
In [9]:	Grouping the data based on the target variable  parkinsons_data.groupby('status').mean()  C:\Users\HP\AppData\Local\Temp\ipykernel_10264\3507134583.py:1: FutureWarning: The default value of numeric_only in DataFrameGroupBy.mean is deprecated. In a future version, numeric_only in DataFrameGroupBy.mean is deprecated. In a future version, numeric_only in DataFrameGroupBy.mean is deprecated. In a future version, numeric_only in DataFrameGroupBy.mean is deprecated. In a future version, numeric_only in DataFrameGroupBy.mean is deprecated. In a future version, numeric_only in DataFrameGroupBy.mean is deprecated. In a future version, numeric_only in DataFrameGroupBy.mean is deprecated.
Out[9]:	c_only will default to False. Either specify numeric_only or select only columns which should be valid for the function.  parkinsons_data.groupby('status').mean()  MDVP:Fo(Hz) MDVP:Fio(Hz) MDVP:Fio(Hz) MDVP:Fio(Hz) MDVP:Jitter(%) MDVP:Jitter(Abs) MDVP:RAP MDVP:PPQ Jitter:DDP MDVP:Shimmer MDVP:Shimmer MDVP:Shimmer(dB) MDVP:APQ Shimmer:DDA NHR HNR RPDE DFJ  status  0 181.937771 223.636750 145.207292 0.003866 0.000023 0.001925 0.002056 0.005776 0.017615 0.162958 0.013305 0.028511 0.011483 24.678750 0.442552 0.695711  1 145.180762 188.441463 106.893558 0.006989 0.000051 0.003757 0.003900 0.011273 0.033658 0.321204 0.027600 0.053027 0.029211 20.974048 0.516816 0.725401  2 rows × 22 columns  It can be clearly observed from these mean values that those who are healthy people have voice of higher frequency than those who are affected by Parkinson's disease.  Data Preprocessing
In [10]:	Seperating the features and target  X= parkinsons_data.drop(columns=['name', 'status'], axis=1) Y= parkinsons_data['status']
In [12]:	161
In [15]:	print(X.shape,X_train.shape,X_test.shape) (195, 22) (156, 22) (39, 22)
In [16]:	Data Standardization  scaler= StandardScaler()  scaler fit(( train)
In [18]: Out[18]:	<pre>scaler.fit(X_train)  * StandardScaler StandardScaler()</pre>
	print(X_train)  [[ 0.63239631 -0.02731081 -0.879850490.97586547 -0.55160318
In [23]:	Support Vector Machine Model  model=svm.SVC(kernel='linear')
Out[24]:	Model Evaluation  Accuracy Score
	Accuracy Score of training data  X_train_prediction=model.predict(X_train)  training_data_accuracy = accuracy_score(Y_train , X_train_prediction)  print('Accuracy Score of training data:',training_data_accuracy)  Accuracy Score of training data: 0.8846153846153846
	Accuracy Score of test data  X_test_prediction=model.predict(X_test) testing_data_accuracy = accuracy_score(Y_test , X_test_prediction) print('Accuracy score of testing data:', testing_data_accuracy)  Accuracy score of testing data: 0.8717948717948718
In [32]:	Building a Predictive System  input_data=(197.076, 206.896, 192.055, 0.00289, 0.00001, 0.00166, 0.00168, 0.00498, 0.01098, 0.0097, 0.00563, 0.0068, 0.00802, 0.01689, 0.00339, 26.775, 0.422229, 0.741367, -7.3483, 0.177551, 1.743867,  # changing input data to a numpy array input_data_as_numpy_array=np.asarray(input_data) # reshaping the numpy array input_data_reshaped =input_data_as_numpy_array.reshape(1,-1) # standardising the data std_data =scaler.transform(input_data_reshaped) prediction = model.predict(std_data) print(prediction)  if (prediction[0] == 0):     print("The person does not have Parkinson's disease.")  else:     print("The person is suffering from Parkinson's disease.")
	[0] The person does not have Parkinson's disease. C:\ProgramData\anaconda3\Lib\site-packages\sklearn\base.py:464: UserWarning: X does not have valid feature names, but StandardScaler was fitted with feature names warnings.warn(