

Untitled

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In [18]: #Team members
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Use read_csv from pandas to load this file as DataFrame and head() to visualise its features

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In [15]: import pandas as pd
         from sklearn.model_selection import train_test_split
         from sklearn.metrics import accuracy_score
         from sklearn.metrics import confusion_matrix
         from sklearn.ensemble import RandomForestClassifier
```

```
In [3]: df = pd.read_csv(f'parkinsons.data')
         print(df.head())
```

	parkinsons.dataname	MDVP:F0(Hz)	MDVP:Fhi(Hz)	MDVP:Flo(Hz)	\
0	phon_R01_S01_1	119.992	157.302	74.997	
1	phon_R01_S01_2	122.400	148.650	113.819	
2	phon_R01_S01_3	116.682	131.111	111.555	
3	phon_R01_S01_4	116.676	137.871	111.366	
4	phon_R01_S01_5	116.014	141.781	110.655	

	MDVP:Jitter(%)	MDVP:Jitter(Abs)	MDVP:RAP	MDVP:PPQ	Jitter:DDP	\
0	0.00784	0.00007	0.00370	0.00554	0.01109	
1	0.00968	0.00008	0.00465	0.00696	0.01394	
2	0.01050	0.00009	0.00544	0.00781	0.01633	
3	0.00997	0.00009	0.00502	0.00698	0.01505	
4	0.01284	0.00011	0.00655	0.00908	0.01966	

	MDVP:Shimmer	...	Shimmer:DDA	NHR	HNR	status	RPDE	\
0	0.04374	...	0.06545	0.02211	21.033	1	0.414783	
1	0.06134	...	0.09403	0.01929	19.085	1	0.458359	
2	0.05233	...	0.08270	0.01309	20.651	1	0.429895	
3	0.05492	...	0.08771	0.01353	20.644	1	0.434969	
4	0.06425	...	0.10470	0.01767	19.649	1	0.417356	

	DFA	spread1	spread2	D2	PPE
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0  0.815285 -4.813031  0.266482  2.301442  0.284654
1  0.819521 -4.075192  0.335590  2.486855  0.368674
2  0.825288 -4.443179  0.311173  2.342259  0.332634
3  0.819235 -4.117501  0.334147  2.405554  0.368975
4  0.823484 -3.747787  0.234513  2.332180  0.410335

```

[5 rows x 24 columns]

Remove the "parkinsons.dataname" feature in the DataFrame so we'll drop this (drop('parkinsons.dataname', axis=1)) . Split the data into features and labels: the feature "status" contains labels therefore you need to drop this feature from DataFrame but create a variable y to which assign the values of status.

```

In [5]: X = df.drop('status', axis=1)
        X = X.drop('parkinsons.dataname', axis=1)
        y = df['status']

```

Split the data into a training and test set of data. Use "from sklearn.model_selection import train_test_split" function for this

```

In [8]: X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=1)

```

Create and train the model. The number of estimators (n_estimators) determines how # dense our decision forest is and the random_state is given for reproducibility.

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In [16]: random_forest = RandomForestClassifier(n_estimators=30, max_depth=10, random_state=1)
        random_forest.fit(X_train, y_train)

```

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Out[16]: RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                                max_depth=10, max_features='auto', max_leaf_nodes=None,
                                min_impurity_decrease=0.0, min_impurity_split=None,
                                min_samples_leaf=1, min_samples_split=2,
                                min_weight_fraction_leaf=0.0, n_estimators=30, n_jobs=1,
                                oob_score=False, random_state=1, verbose=0, warm_start=False)

```

Evaluate our model on our test set.

```

In [17]: y_predict = random_forest.predict(X_test)
        accuracy_score(y_test, y_predict)
        pd.DataFrame( confusion_matrix(y_test, y_predict),
                        columns=['Predicted Healthy', 'Predicted Parkinsons'],
                        index=['True Healthy', 'True Parkinsons'])

```

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Out[17]:

```

	Predicted Healthy	Predicted Parkinsons
True Healthy	11	1
True Parkinsons	2	35