



Artificial Intelligence And Machine Learning

Parkinson's Disease Detection Python Project

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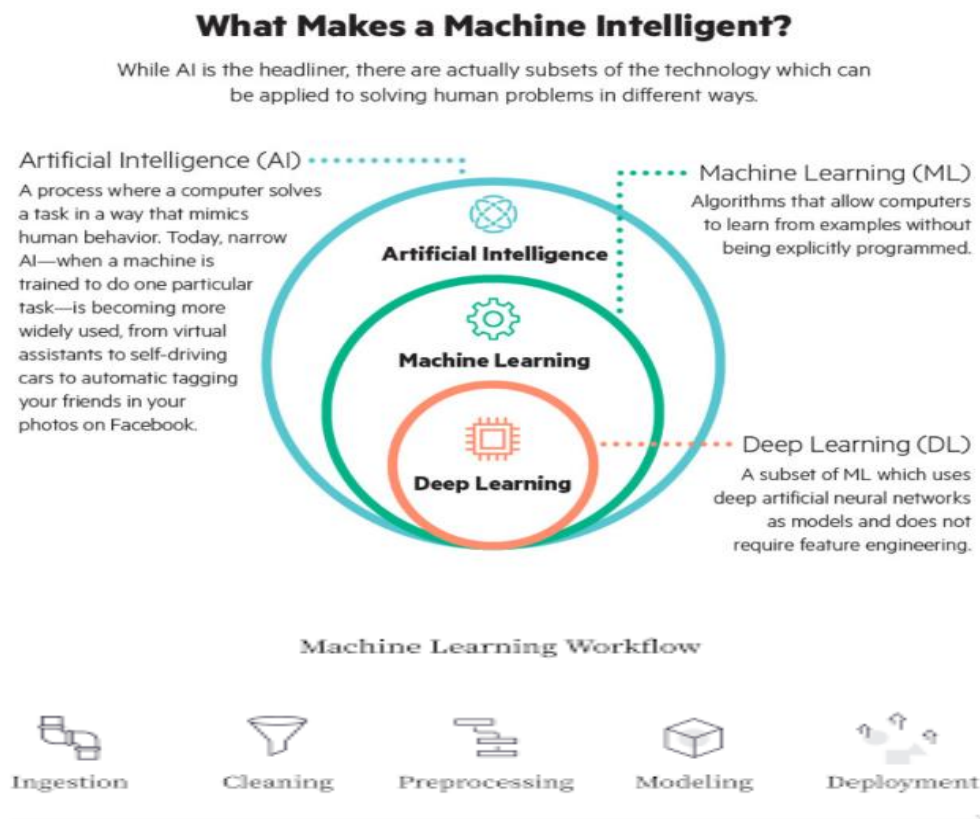
INTRODUCTION:

ARTIFICIAL INTELLIGENCE:

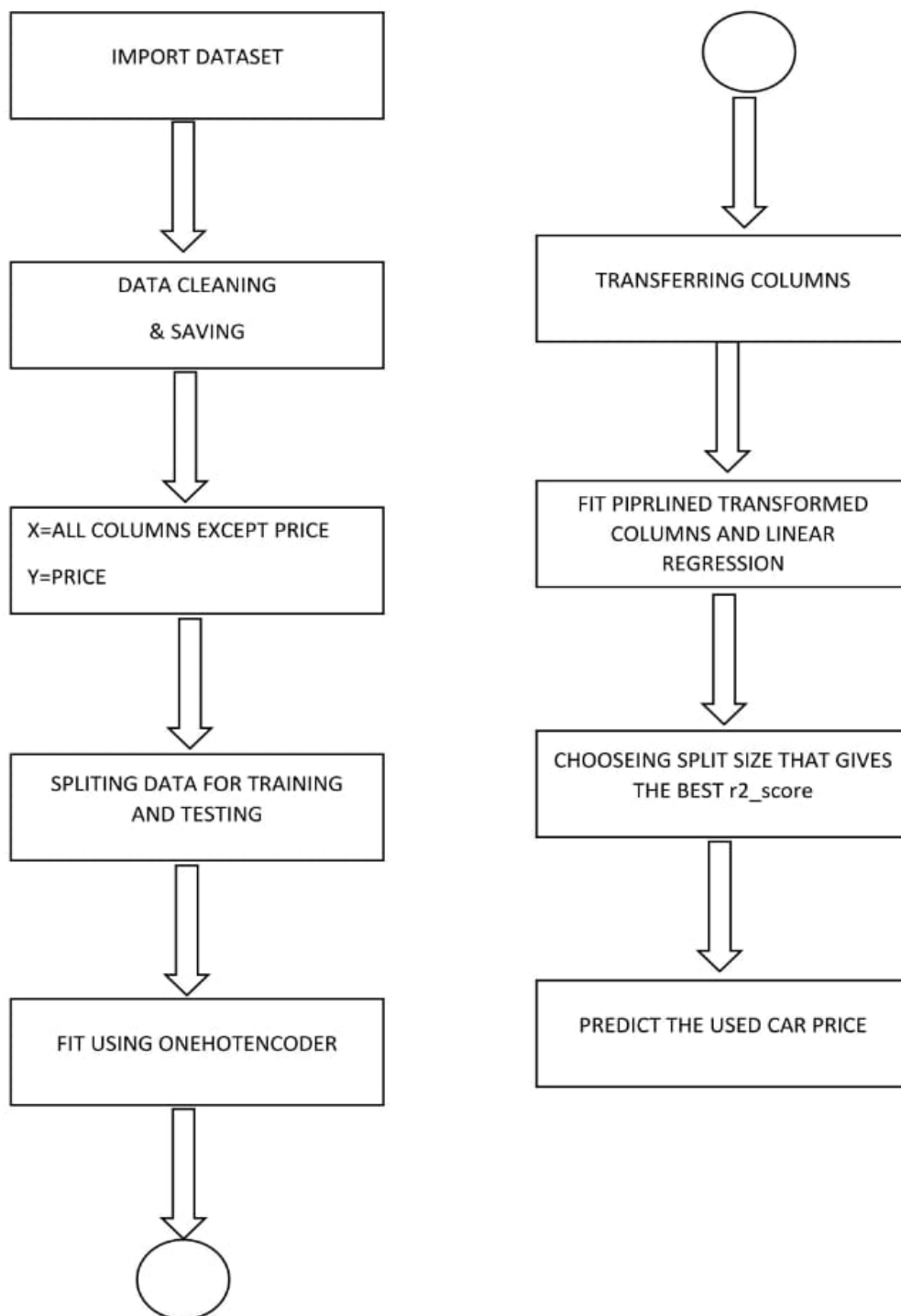
Artificial intelligence (AI) refers to the simulation of human intelligence in machines that are programmed to think like humans and mimic their actions. The term may also be applied to any machine that exhibits traits associated with a human mind such as learning and problem solving.



The ideal characteristic of artificial intelligence is its ability to rationalize and take actions that have the best chance of achieving a specific goal. A subset of artificial intelligence is machine learning, which refers to the concept that computer program can automatically learn from and adapt to new data without being assisted by humans. Deep learning techniques enable this automatic learning through the absorption of huge amount of unstructured data such as text, images, or video.



ALGORITHM:



ABSTRACT:

Parkinson is a nervous system disorder that affects movement. The dataset contains 195 records of people with 23 different attributes which contain biomedical measurements. The data is used to separate healthy people from people with Parkinson's disease.

THINGS TO IMPLEMENT:

The model can be used to differentiate healthy people from people having Parkinson's disease.

INTRODUCTION:

Parkinson's disease is a progressive disorder of the central nervous system affecting movement and inducing tremors and stiffness. It has 5 stages to it and affects more than 1 million individuals every year in India. This is chronic and has no cure yet. It is a neurodegenerative disorder affecting dopamine-producing neurons in the brain.

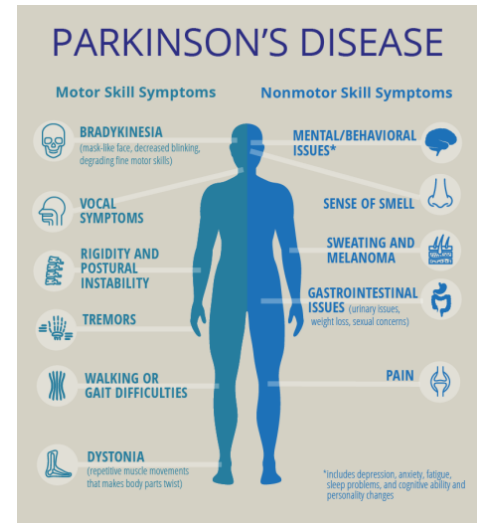
THE DATA:

The data used in this project was downloaded from kaggle.

This data includes columns like name, 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:APQ3,Shimmer:APQ5,MDVP:APQ,Shimmer:DDA,NHR,HNR,status,RPDE,D2,DFA,spread1,spread2,PPE.

ROLES:

- DATA ACQUISITION AND CLEANING
- DATA VISUALIZATION
- DATA MODELLING
- TESTING
- COMPARISON AND MEASUREMENT



ARTIFICIAL INTELLIGENCE / MACHINE LEARNING CODE TO ANALYZE PARKINSON'S DISEASE:

First, import all the libraries/packages which are necessary to analyze the given dataset about Parkinson's disease.

```
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
```

Next step is to import the given dataset about Parkinson's disease.

```
parkinson_data=pd.read_csv('Parkinson disease.csv')
print(parkinson_data)
```

OUTPUT:

	name	MDVP:F0 (Hz)	MDVP:Fhi (Hz)	...	spread2	D2	PP
E							
0	phon_R01_S01_1	119.992	157.302	...	0.266482	2.301442	0.28465
4							
1	phon_R01_S01_2	122.400	148.650	...	0.335590	2.486855	0.36867
4							
2	phon_R01_S01_3	116.682	131.111	...	0.311173	2.342259	0.33263
4							
3	phon_R01_S01_4	116.676	137.871	...	0.334147	2.405554	0.36897
5							
4	phon_R01_S01_5	116.014	141.781	...	0.234513	2.332180	0.41033
5							
..
.							
190	phon_R01_S50_2	174.188	230.978	...	0.121952	2.657476	0.13305
0							
191	phon_R01_S50_3	209.516	253.017	...	0.129303	2.784312	0.16889
5							
192	phon_R01_S50_4	174.688	240.005	...	0.158453	2.679772	0.13172
8							
193	phon_R01_S50_5	198.764	396.961	...	0.207454	2.138608	0.12330
6							
194	phon_R01_S50_6	214.289	260.277	...	0.190667	2.555477	0.14856
9							

[195 rows x 24 columns]

Parkinson's Disease Detection

Get the features(x) and labels(y) from the DataFrame (dataset). The features are all the columns except 'names' and 'status', and the labels(y) are those in the 'status' column.

```
x=parkinson_data.drop(columns=['name','status'],axis=1)
y=parkinson_data['status']
print(x)
print(y)
```

OUTPUT:

```
-
   MDVP:F0 (Hz)  MDVP:F1 (Hz)  MDVP:F2 (Hz)  ...  spread2  D2  PPE
0      119.992      157.302      74.997  ...  0.266482  2.301442  0.284654
1      122.400      148.650     113.819  ...  0.335590  2.486855  0.368674
2      116.682     131.111     111.555  ...  0.311173  2.342259  0.332634
3      116.676     137.871     111.366  ...  0.334147  2.405554  0.368975
4      116.014     141.781     110.655  ...  0.234513  2.332180  0.410335
..      ...      ...      ...  ...  ...      ...      ...
190     174.188     230.978      94.261  ...  0.121952  2.657476  0.133050
191     209.516     253.017      89.488  ...  0.129303  2.784312  0.168895
192     174.688     240.005      74.287  ...  0.158453  2.679772  0.131728
193     198.764     396.961      74.904  ...  0.207454  2.138608  0.123306
194     214.289     260.277      77.973  ...  0.190667  2.555477  0.148569

[195 rows x 22 columns]
```

```
0      1
1      1
2      1
3      1
4      1
..
190    0
191    0
192    0
193    0
194    0
Name: status, Length: 195, dtype: int64
```

Now, split the dataset into training and testing sets keeping 20% of the data for testing and printing.

```
x_train,x_test,y_train,y_test = train_test_split(x,y,test_size=0.2,random_state=2)
print(x.shape,x_train.shape,x_test.shape)
```

OUTPUT:

```
(195, 22) (156, 22) (39, 22)
```

Initialize an XGBClassifier and train the model. This classifies using eXtreme Gradient Boosting- using gradient boosting algorithms for modern data science problems. It falls under the

category of Ensemble Learning in ML, where we train and predict using many models to produce one superior output.

What is XGBoost?

XGBoost is a new Machine Learning algorithm designed with speed and performance in mind. XGBoost stands for eXtreme Gradient Boosting and is based on decision trees. In this project, we will import the XGBClassifier from the xgboost library; this is an implementation of the scikit-learn API for XGBoost classification.

```
from xgboost import XGBClassifier
model = XGBClassifier()
model.fit(x_train, y_train,)
print(model)
```

OUTPUT:

```
XGBClassifier(base_score=0.5, booster='gbtree', colsample_bylevel=1,
              colsample_bynode=1, colsample_bytree=1, gamma=0, gpu_id=-1,
              importance_type='gain', interaction_constraints='',
              learning_rate=0.300000012, max_delta_step=0, max_depth=6,
              min_child_weight=1, missing=nan, monotone_constraints='()',
              n_estimators=100, n_jobs=4, num_parallel_tree=1, random_state=0,
              reg_alpha=0, reg_lambda=1, scale_pos_weight=1, subsample=1,
              tree_method='exact', validate_parameters=1, verbosity=None)
```

To standardize and fit the Data and transfer the data in similar range.

```
#Data Standardization
scaler=StandardScaler()
scaler.fit(x_train)
x_train=scaler.transform(x_train)
x_test=scaler.transform(x_test)
print(x_train)
```

OUTPUT:

```
[ [ 0.63239631 -0.02731081 -0.87985049 ... -0.97586547 -0.55160318
    0.07769494]
  [-1.05512719 -0.83337041 -0.9284778 ... 0.3981808 -0.61014073
    0.39291782]
  [ 0.02996187 -0.29531068 -1.12211107 ... -0.43937044 -0.62849605
    -0.50948408]
  ...
  [-0.9096785 -0.6637302 -0.160638 ... 1.22001022 -0.47404629
    -0.2159482 ]
  [-0.35977689 0.19731822 -0.79063679 ... -0.17896029 -0.47272835
    0.28181221]
  [ 1.01957066 0.19922317 -0.61914972 ... -0.716232 1.23632066
    -0.05829386]]
```

Support Vector Machine Model(SVM) and Accuracy Score

```
#Support Vector Machine Model(SVM)
model=svm.SVC(kernel='linear')

#Training the svm model with training data
model.fit(x_train,y_train)

#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 testing data
x_test_prediction=model.predict(x_test)
test_data_accuracy=accuracy_score(y_test,x_test_prediction)
print("Accuracy score of test data: ",test_data_accuracy)
```

OUTPUT:

```
[20:40:20] WARNING: C:/Users/Administrator/workspace/xgboost-win64_release_1.4.0
/src/learner.cc:1095: Starting in XGBoost 1.3.0, the default evaluation metric u
sed with the objective 'binary:logistic' was changed from 'error' to 'logloss'.
Explicitly set eval_metric if you'd like to restore the old behavior.
Accuracy score of training data: 0.8846153846153846
Accuracy score of test data: 0.8717948717948718
```

Predicting the output that the person is either having Parkinson's disease or not.

```
#Predicting
input_data=(119.992,157.302,74.997,0.00784,0.00007,0.0037,0.00554,0.01109,0.04374,0.426,0.02182,
#changing input data to a numpy array
input_data_as_numpy_array= np.asarray(input_data)
#reshape the numpy array
input_data_reshaped= input_data_as_numpy_array.reshape(1,-1)
#standarize the data
std_data= scaler.transform(input_data_reshaped)

prediction= model.predict(std_data)
print(prediction)

if(prediction[0]==0):
    print("Person is not having Parkinson Disease")
else:
    print("Person is having Parkinson Disease")
```

OUTPUT:

```
[20:42:07] WARNING: C:/Users/Administrator/workspace/xgboost-win64_release_1.4.0
/src/learner.cc:1095: Starting in XGBoost 1.3.0, the default evaluation metric u
sed with the objective 'binary:logistic' was changed from 'error' to 'logloss'.
Explicitly set eval_metric if you'd like to restore the old behavior.
[1]
Person is having Parkinson Disease

[0]
Person is not having Parkinson Disease
```

CONCLUSION:

Parkinson's disease affects the CNS of the brain and has yet no treatment unless it's detected early. Late detection leads to no treatment and loss of life. Thus its early detection is significant. For early detection of the disease, we utilized machine learning algorithms such as XGBoost . We checked our Parkinson disease data and find out XGBoost is the best Algorithm to predict the onset of the disease which will enable early treatment and save a life. Using support vector and Standard scaler we conclude whether the person has got Parkinson's disease or not.