

Parkinson_Prediction

```
import numpy as np
import pandas as pd
from sklearn import svm
from sklearn.metrics import accuracy_score
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split
import os

# Define the directory path and list all files in the directory
directory_path = r'C:\Users\HP\Desktop\jupyter projects'
files = os.listdir(directory_path)
print(files)

# Define the file path and check if the file exists
file_path = r'C:\Users\HP\Desktop\jupyter
projects\parkinsons.data'
if os.path.isfile(file_path):
    # Load the .data file into a DataFrame
    df = pd.read_csv(file_path, delimiter=',')

    # Display the first few rows of the DataFrame
    print(df.head())
```

else:

```
print("File not found. Please check the file path.")
```

```
# Display basic information about the DataFrame
```

```
print(df.info())
```

```
# Display statistical summary of the numeric columns
```

```
print(df.describe())
```

```
# Display the shape of the DataFrame (number of rows and columns)
```

```
print(df.shape)
```

```
# Check for missing values in each column
```

```
print(df.isnull().sum())
```

```
# Display the distribution of the target variable
```

```
print(df['status'].value_counts())
```

```
# Convert all columns to numeric, coercing errors to NaN
```

```
df = df.apply(pd.to_numeric, errors='coerce')
```

```
# Ensure 'status' is treated as a categorical variable
```

```
df['status'] = df['status'].astype('category')
```

```
# Select only numeric columns for aggregation
```

```
numeric_df = df.select_dtypes(include='number')
```

```
# Perform groupby operation and calculate the mean for each  
'status'
```

```
result = numeric_df.groupby(df['status']).mean()
```

```
print(result)
```

```
# Separate features (X) and target (Y) variables
```

```
X = df.drop(columns=['name', 'status'], axis=1)
```

```
Y = df['status']
```

```
print(X)
```

```
print(Y)
```

```
# Split the dataset into training and testing sets
```

```
X_train, X_test, Y_train, Y_test = train_test_split(X, Y,  
test_size=0.2, random_state=2)
```

```
# Display the shape of the training and testing data
```

```
print(X.shape, X_train.shape, X_test.shape)
```

```
# Standardize features by removing the mean and scaling to  
unit variance
```

```
ss = StandardScaler()
```

```
X_train = ss.fit_transform(X_train) # Fit and transform  
training data
```

```
X_test = ss.transform(X_test)      # Transform testing data
```

```
# Create and train the SVM model with a linear kernel
```

```
model = svm.SVC(kernel="linear")
```

```
model.fit(X_train, Y_train)
```

```
# Model evaluation: Accuracy on training and testing data
```

```
X_train_pred = model.predict(X_train)
```

```
train_data_acc = accuracy_score(Y_train, X_train_pred)
```

```
print("Accuracy of training data:", train_data_acc)
```

```
X_test_pred = model.predict(X_test)
```

```
test_data_acc = accuracy_score(Y_test, X_test_pred)
```

```
print('Accuracy of testing data:', test_data_acc)
```

```
# Define input data with the same number of features as the  
training data
```

```
input_data = [1.0] * X_train.shape[1] # Replace with actual  
feature values
```

```
input_data_np = np.asarray(input_data).reshape(1, -1)
```

```
# Standardize the input data
```

```
s_data = ss.transform(input_data_np)
```

```
# Make a prediction with the trained SVM model
```

```
pred = model.predict(s_data)
```

```
print(pred)
```

```
# Interpret the prediction
```

```
if pred[0] == 0:
```

```
    print("Negative")
```

```
else:
```

```
    print("Positive")
```