#Name : Rajshri Kirandas Satpute #Year : 3rd year #Section : B #Roll No : 55 #Date : 08/02/2024 import numpy as np import pandas as pd import seaborn as sns import matplotlib.pyplot as plt import os In [4]: os.chdir("C:\\Users\\fatin\\OneDrive\\Desktop") df = pd.read_csv("Iris.csv") In [6]: df.head() Out[6]: Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm Species 0 1 5.1 3.5 1.4 0.2 Iris-setosa 0.2 Iris-setosa **1** 2 4.9 3.0 1.4 **2** 3 3.2 0.2 Iris-setosa 4.7 1.3 **3** 4 4.6 3.1 1.5 0.2 Iris-setosa **4** 5 5.0 3.6 1.4 0.2 Iris-setosa In [7]: df.describe() Out[7]: Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm count 150.000000 150.000000 150.000000 150.000000 150.000000 75.500000 5.843333 3.054000 3.758667 1.198667 mean 43.445368 0.828066 0.433594 1.764420 0.763161 0.100000 min 1.000000 4.300000 2.000000 1.000000 **25**% 38.250000 5.100000 2.800000 1.600000 0.300000 3.000000 75.500000 5.800000 4.350000 1.300000 1.800000 **75**% 112.750000 6.400000 3.300000 5.100000 4.400000 2.500000 max 150.000000 7.900000 6.900000 In [8]: df.info() <class 'pandas.core.frame.DataFrame'> RangeIndex: 150 entries, 0 to 149 Data columns (total 6 columns): # Column Non-Null Count Dtype -----0 Id 150 non-null int64 SepalLengthCm 150 non-null float64 2 SepalWidthCm 150 non-null float64 3 PetalLengthCm 150 non-null float64 PetalWidthCm 150 non-null 4 float64 5 150 non-null Species object dtypes: float64(4), int64(1), object(1) memory usage: 7.2+ KB df.shape (150, 6)In [10]: df.drop("Id", axis=1, inplace=True) #droping id becuase it is no use to us , Inplace = True means changes will take effect in original dataframe df.head() Out[10]: SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm **Species** 0 5.1 0.2 Iris-setosa 3.5 1.4 4.9 3.0 1.4 1 0.2 Iris-setosa 2 1.3 4.7 3.2 0.2 Iris-setosa 3 4.6 3.1 1.5 0.2 Iris-setosa 4 5.0 3.6 1.4 0.2 Iris-setosa **Data Visualization** In [11]: print(df["Species"].value_counts()) sns.countplot(df["Species"]) 50 Iris-setosa Iris-versicolor 50 Iris-virginica 50 Name: Species, dtype: int64 C:\Users\fatin\anaconda3\lib\site-packages\seaborn_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional a rgument will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation. warnings.warn(<AxesSubplot:xlabel='Species', ylabel='count'> Out[11]: 50 40 30 30 20 10 Iris-versicolor Iris-setosa Iris-virginica In [12]: plt.figure(figsize=(8,4)) sns.heatmap(df.corr(),annot=True,fmt=".0%") #draws heatmap with input as the correlation matrix calculted by(df.corr()) plt.show() -1.0 SepalLengthCm -100% -11% 87% 82% - 0.8 - 0.6 -11% 100% 42% -36% SepalWidthCm 0.4 0.2 PetalLengthCm 87% 42% 100% 96% 0.0 -0.2 PetalWidthCm -36% 100% 82% 96% SepalLengthCm SepalWidthCm PetalLengthCm In [13]: # We'll use seaborn's FacetGrid to color the scatterplot by species sns.FacetGrid(df, hue="Species", height=5).map(plt.scatter, "SepalLengthCm", "SepalWidthCm").add_legend() <seaborn.axisgrid.FacetGrid at 0x1ee1709f760> Out[13]: 4.0 SepalWidthCm 0.8 Species Iris-setosa Iris-versicolor Iris-virginica 2.5 2.0 5.0 5.5 6.0 6.5 7.0 7.5 SepalLengthCm In [14]: #let Create a pair plot of some columns sns.pairplot(df.iloc[:,:], hue='Species') # graph also tell us about the realationship between the two columns <seaborn.axisgrid.PairGrid at 0x1ee17139c10> SepalLengthCm SepalWidthCm 3.0 2.0 Species Iris-setosa Iris-versicolor Iris-virginica 2.5 2.0 0.5 SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm In [15]: # We can quickly make a boxplot with Pandas on each feature split out by species df.boxplot(by="Species", figsize=(15,15)) array([[<AxesSubplot:title={'center':'PetalLengthCm'}, xlabel='[Species]'>, <AxesSubplot:title={'center':'PetalWidthCm'}, xlabel='[Species]'>], [<AxesSubplot:title={'center':'SepalLengthCm'}, xlabel='[Species]'>, <AxesSubplot:title={'center':'SepalWidthCm'}, xlabel='[Species]'>]], dtype=object) Boxplot grouped by Species PetalLengthCm PetalWidthCm SepalLengthCm SepalWidthCm Iris-setosa Iris-versicolor Iris-virginica Iris-setosa Iris-versicolor Iris-virginica # importing all1 the necessary packages to Logistic Regression from sklearn.linear_model import LogisticRegression # for Logistic Regression algorithm from sklearn.model_selection import train_test_split #to split the dataset for training and testing from sklearn import metrics #for checking the model accuracy Splitting the Data into Training and Testing Dataset In [17]: X=df.iloc[:,0:4] Y=df["Species"] X.head() SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm Out[17]: 3.5 1.4 0.2 4.9 3.0 1.4 0.2 2 4.7 3.2 1.3 0.2 3.1 1.5 4.6 0.2 4 5.0 3.6 1.4 0.2 In [18]: X_train, X_test, Y_train, Y_test=train_test_split(X, Y, test_size=0.25, random_state=0)# in this our main data is split into train and test # the attribute test_size=0.3 splits the data into 70% and 30% ratio. train=70% and test=30% print("Train Shape", X_train.shape) print("Test Shape", X_test.shape) Train Shape (112, 4) Test Shape (38, 4) Logistic Regression In [19]: log = LogisticRegression() log.fit(X_train,Y_train) prediction=log.predict(X_test) print('The accuracy of the Logistic Regression is', metrics.accuracy_score(prediction, Y_test)) The accuracy of the Logistic Regression is 0.9736842105263158 In []: