## importing useful libraries In [2]: from sklearn.cluster import KMeans import pandas as pd import seaborn as sns import matplotlib.pyplot as plt plt.style.use('ggplot') import numpy as np importing dataset In [10]: d=pd.read\_csv("Iris.csv") Out[10]: Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm Species 3.5 1.4 5.1 Iris-setosa 1 2 4.9 3.0 1.4 0.2 Iris-setosa 1.3 4.7 3.2 0.2 Iris-setosa 3 4 4.6 3.1 1.5 0.2 Iris-setosa 5 5.0 3.6 1.4 Iris-setosa 6.7 **145** 146 3.0 5.2 2.3 Iris-virginica **146** 147 6.3 2.5 5.0 1.9 Iris-virginica **147** 148 6.5 3.0 5.2 2.0 Iris-virginica **148** 149 6.2 3.4 5.4 2.3 Iris-virginica 5.9 1.8 Iris-virginica **149** 150 3.0 5.1 150 rows × 6 columns In [11]: import pandas\_profiling d.profile\_report() Pandas Profiling Report Overview Variables Interactions Correlations Missing values Sample Overview Overview Warnings 4 Reproduction Dataset statistics Variable types **Number of variables** 5 6 NUM **Number of observations** 150 CAT 1 0 Missing cells Missing cells (%) 0.0% 0 **Duplicate rows Duplicate rows (%)** 0.0% Total size in memory 7.2 KiB Average record size in memory 48.9 B **Variables Distinct** 75.5 150 Mean Id Real number ( $\mathbb{R}_{\geq 0}$ ) **Distinct (%)** 100.0% Minimum 1 Out[11]: **Data Visualization** import numpy as np In [12]: from sklearn import datasets import matplotlib.pyplot as plt from yellowbrick.target.feature\_correlation import feature\_correlation #Load the iris dataset data = datasets.load\_iris() X, y = data['data'], data['target'] features = np.array(data['feature\_names']) visualizer = feature\_correlation(X, y, labels=features) plt.tight\_layout() Features correlation with dependent variable petal width (cm) petal length (cm) sepal width (cm) sepal length (cm) -0.4-0.20.0 0.2 0.4 0.6 0.8 1.0 Pearson Correlation <Figure size 576x396 with 0 Axes> In [21]: iris\_setosa=d.loc[d["Species"]=="Iris-setosa"] iris\_virginica=d.loc[d["Species"]=="Iris-virginica"] iris\_versicolor=d.loc[d["Species"]=="Iris-versicolor"] $\verb|sns.FacetGrid(d,hue="Species",size=3).map(sns.distplot,"PetalLengthCm").add\_legend()|$ sns.FacetGrid(d, hue="Species", size=3).map(sns.distplot, "PetalWidthCm").add\_legend() sns.FacetGrid(d,hue="Species",size=3).map(sns.distplot,"SepalLengthCm").add\_legend() sns.FacetGrid(d, hue="Species", size=3).map(sns.distplot, "SepalWidthCm").add\_legend() plt.show() /home/goutami/anaconda3/lib/python3.7/site-packages/seaborn/axisgrid.py:316: UserWarning: The `size` parameter has been renamed to `height`; please update your code. warnings.warn(msg, UserWarning) /home/goutami/anaconda3/lib/python3.7/site-packages/seaborn/distributions.py:2551: FutureWarn ing: `distplot` is a deprecated function and will be removed in a future version. Please adap t your code to use either `displot` (a figure-level function with similar flexibility) or `hi stplot` (an axes-level function for histograms). warnings.warn(msg, FutureWarning) /home/goutami/anaconda3/lib/python3.7/site-packages/seaborn/distributions.py:2551: FutureWarn ing: `distplot` is a deprecated function and will be removed in a future version. Please adap t your code to use either `displot` (a figure-level function with similar flexibility) or `hi stplot` (an axes-level function for histograms). warnings.warn(msg, FutureWarning) /home/goutami/anaconda3/lib/python3.7/site-packages/seaborn/distributions.py:2551: FutureWarn ing: `distplot` is a deprecated function and will be removed in a future version. 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Please adap t your code to use either `displot` (a figure-level function with similar flexibility) or `hi stplot` (an axes-level function for histograms). warnings.warn(msg, FutureWarning) 3.0 25 2.0 Species 15 lris-setosa lris-versicolor 1.0 lris-virginica 0.5 0.0 6 PetalLengthCm 10 8 Species 6 lris-setosa lris-versicolor lris-virginica 2 0 2 PetalWidthCm 1.50 1.25 1.00 Species 0.75 lris-setosa lris-versicolor 0.50 lris-virginica 0.25 0.00 8 SepalLengthCm1.50 1.25 1.00 Species 0.75 lris-setosa lris-versicolor 0.50 lris-virginica 0.25 0.00 SepalWidthCm In [22]: sns.boxplot(x="Species", y="PetalLengthCm", data=d) plt.show() PetalLengthCm Iris-versicolor Iris-setosa Iris-virginica Species In [23]: sns.violinplot(x="Species", y="PetalLengthCm", data=d) plt.show() Iris-setosa Iris-versicolor Iris-virginica Species sns.set\_style("whitegrid") sns.pairplot(d, hue="Species", size=3); plt.show() /home/goutami/anaconda3/lib/python3.7/site-packages/seaborn/axisgrid.py:1912: UserWarning: Th e `size` parameter has been renamed to `height`; please update your code. warnings.warn(msg, UserWarning) 8.0 7.5 2.5 2.0 2.0 PetalWidthCm PetalLengthCm **Observations** In [31]: #if 0≤petallength≤2 and 0≤petalwidth≤0.7then setosa #if 2≤petallenght≤5.2 and 1≤petallength≤1.7 then versicolor and #else virginica split independent & target variable inputs = d.drop(['Id', 'Species'], axis='columns')#independent variable target = d.drop(['Id', 'SepalLengthCm', 'SepalWidthCm', 'PetalLengthCm', 'PetalWidthCm'], axis='c olumns') In [33]: inputs Out[33]: SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm 0.2 5.1 3.5 1.4 1 4.9 3.0 1.4 0.2 0.2 4.7 3.2 1.3 3 4.6 3.1 1.5 0.2 5.0 3.6 1.4 0.2 2.3 145 6.7 3.0 5.2 146 6.3 2.5 5.0 1.9 147 6.5 5.2 2.0 3.0 148 6.2 3.4 5.4 2.3 5.9 149 3.0 5.1 1.8 150 rows × 4 columns In [34]: target Out[34]: **Species** Iris-setosa Iris-setosa Iris-setosa Iris-setosa Iris-setosa 145 Iris-virginica 146 Iris-virginica 147 Iris-virginica 148 Iris-virginica 149 Iris-virginica 150 rows × 1 columns In [35]: #encode categorical feature from sklearn.preprocessing import LabelEncoder In [36]: le\_Species = LabelEncoder() In [37]: | target['Species'] = le\_Species.fit\_transform(target['Species']) target.head(70) Out[37]: Species 0 1 0 0 3 0 0 ... 65 66 67

Out[42]: array([0]) In [43]: model.predict([[6.7,3.0,5.2,2.3]])#iris virginica Out[43]: array([2])

In [44]: model.predict([[6.4,2.9,4.3,1.3]])#iris versicolor

**Decision Tree model implementation** 

Out[44]: array([1]) In [34]: model.predict([[6.6,3.9,5.3,1.3]]) Out[34]: array([2])

model accuracy

In [41]: model.fit(inputs, target) Out[41]: DecisionTreeClassifier(class\_weight=None, criterion='gini', max\_depth=None, max\_features=None, 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, presort=False, random\_state=None, splitter='best') value prediction In [42]: model.predict([[5.1,3.5,1.4,0.2]])#iris setosa

68 69

70 rows × 1 columns

In [39]: **from sklearn import** tree

In [40]: | model = tree.DecisionTreeClassifier()

**Submitted By Goutami Dey** 

score

Out[45]: 1.0 Thank you

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In [45]: score=model.score(inputs, target)