0	2 4.9 3.0 1.4 0.2 Iris-setosa
2 3 4	3 4.7 3.2 1.3 0.2 Iris-setosa 4 4.6 3.1 1.5 0.2 Iris-setosa 5 5.0 3.6 1.4 0.2 Iris-setosa .drop('Id', axis=1, inplace = True)
df	.drop('Id', axis=1, inplace = True) .head() SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm Species 5.1 3.5 1.4 0.2 Iris-setosa 4.9 3.0 1.4 0.2 Iris-setosa
3 4	4.7 3.2 1.3 0.2 Iris-setosa 4.6 3.1 1.5 0.2 Iris-setosa 5.0 3.6 1.4 0.2 Iris-setosa
(15	.shape 0, 5) .dtypes
Sep Pet Pet Spe dty	alLengthCm float64 alWidthCm float64 alLengthCm float64 alLengthCm float64 alLengthCm float64 alWidthCm float64 cies object pe: object
Fal	.isnull().values.any() se .describe()
cou mea s	td 0.828066 0.433594 1.764420 0.763161
25 50 75 ma	% 5.100000 2.800000 1.600000 0.300000 % 5.800000 3.00000 4.350000 1.300000 % 6.40000 3.30000 5.100000 1.800000
Sep	SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm
Pet Pe	palWidthCm
Iri Iri Nam	s-setosa 50 s-versicolor 50 s-virginica 50 e: Species, dtype: int64 s.pairplot(df.iloc[:,1:])
4 4 E	aborn.axisgrid.PairGrid at 0x1a09424c100>
PetalLengthCm	
0	.hist()
	ay([[<axessubplot:title={'center':'sepallengthcm'}>,</axessubplot:title={'center':'sepallengthcm'}>
20 - 10 - 0 -	PetalLengthcm 8 2 PetalWidthCm ⁴
30 - 20 - 10 - 0 -	
ax ax co sp ct	<pre>g = plt.figure(figsize=(9, 6))</pre>
pl	t.title('Percentage of different species in the Dataset') t.show() Percentage of different species in the Dataset
	Iris-setosa Iris-virginica
	33.33%
	33.33%
Sep	.skew() alLengthCm 0.314911
Pet Pet dty	alWidthCm
0 1 2 3 4 145 146	False
147 148 149 Len	False False
<fi 4 4 E</fi 	aborn.axisgrid.PairGrid at 0x1a094b3a880> gure size 720x360 with 0 Axes>
SepalWidthCr	
PetalLengthCm	
WidthCm 1	
	SepalWidthCm PetalLengthCm PetalWidthCm
	.boxplot(column='PetalLengthCm') esSubplot:>
5 - 4 - 3 -	
2 1 df	PetalLengthCm .quantile(0.75)-df.quantile(0.25)
Sep Pet Pet dty	alLengthCm 1.3 alWidthCm 0.5 alLengthCm 3.5 alWidthCm 1.5 pe: float64
0	SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm Species 5.1 3.5 1.4 0.2 Iris-setosa 4.9 3.0 1.4 0.2 Iris-setosa
	4.7 3.2 1.3 0.2 Iris-setosa 4.6 3.1 1.5 0.2 Iris-setosa 5.0 3.6 1.4 0.2 Iris-setosa df.drop('Species', axis=1)
fr X_	<pre>df['Species'] om sklearn.model_selection import train_test_split train, X_test, y_train, y_test = train_test_split(x, y, test_size = 0.30, random_state=1)</pre>
dT dT	<pre>om sklearn.tree import DecisionTreeClassifier ree = DecisionTreeClassifier(criterion = 'entropy', max_depth=5) ree.fit(X_train, y_train) isionTreeClassifier(criterion='entropy', max_depth=5)</pre>
Acc	<pre>int("Accuracy:", dTree.score(X_test,y_test) * 100) uracy: 95.555555555556 pred = dTree.predict(X_test) int(y_pred)</pre>
['I 'I 'I 'I 'I	ris-setosa' 'Iris-versicolor' 'Iris-versicolor' 'Iris-setosa' ris-virginica' 'Iris-versicolor' 'Iris-virginica' 'Iris-setosa' ris-setosa' 'Iris-virginica' 'Iris-versicolor' 'Iris-setosa' ris-virginica' 'Iris-versicolor' 'Iris-setosa' ris-virginica' 'Iris-versicolor' 'Iris-setosa' ris-versicolor' 'Iris-versicolor' 'Iris-setosa' ris-versicolor' 'Iris-versicolor' 'Iris-setosa' ris-versicolor' 'Iris-versicolor' 'Iris-setosa' 'Iris-setosa' ris-virginica' 'Iris-versicolor' 'Iris-setosa' 'Iris-setosa'
'I 'I 'I 'I	ris-versicolor' 'Iris-virginica' 'Iris-versicolor' 'Iris-virginica' ris-versicolor' 'Iris-virginica' 'Iris-virginica' 'Iris-setosa' ris-versicolor' 'Iris-setosa' 'Iris-versicolor' 'Iris-virginica' ris-versicolor' 'Iris-setosa' 'Iris-versicolor' 'Iris-virginica' ris-virginica' 'Iris-setosa' 'Iris-versicolor' 'Iris-virginica' ris-versicolor'] atures = df.columns[:-1]
fr ou	<pre>asses = df['Species'].unique().tolist() om sklearn.tree import plot_tree tput = plot_tree(dTree, feature_names=features, class_names=classes, filled=True) r o in output: arrow = o.arrow_patch if arrow is not None: arrow.set_edgecolor('black')</pre>
	arrow.set_linewidth(1) Pedallangstr.cm == 2.8
er su valu class	PetaliangtiCm <= 9.0 minary = 0.611 arriphs = 25 value = [5, 51, 51] class = Intervityion region = 0.0 surphs = 0.0 whitesy = 0.811 samplas = 4 value = [0, 1, 2] class = Intervityion class = Interv
	ree.fit(X_train, y_train) isionTreeClassifier(criterion='entropy', max_depth=5)
ou	t.figure(figsize=(20, 15)) tput = plot_tree(dTree, feature_names=features, class_names=classes, filled= True) r o in output: arrow = o.arrow_patch if arrow is not None: arrow.set_edgecolor('black')
	PetalLengthCm <= 2.6 entropy = 1.582 samples = 105
	value = [36, 32, 37] class = Iris-virginica PetalWidthCm <= 1.65
	entropy = 0.0 samples = 36 value = [36, 0, 0] class = Iris-setosa retalWidthCH <= 1.03 entropy = 0.996 samples = 69 value = [0, 32, 37] class = Iris-virginica
	PetalLengthCm <= 5.0 entropy = 0.431 samples = 34 value = [0, 31, 3] class = Iris versiseler
	entropy = 0.0 SepalLengthCm <= 6.05 entropy = 0.811 SepalWidthCm <= 3.1 entropy = 0.811 SepalWidthCm <= 3.1 entropy = 0.811 SepalWidthCm <= 3.1 entropy = 0.811
	samples = 30 value = [0, 30, 0] class = Iris-versicolor samples = 4 value = [0, 1, 3] class = Iris-virginica samples = 4 value = [0, 1, 3] class = Iris-virginica samples = 31 value = [0, 0, 31] class = Iris-virginica
	entropy = 0.0 samples = 1 value = [0, 1, 0] class = Iris-versicolor entropy = 0.0 samples = 3 value = [0, 0, 3] class = Iris-virginica entropy = 0.0 samples = 3 value = [0, 0, 3] class = Iris-virginica entropy = 0.0 samples = 3 value = [0, 0, 3] class = Iris-virginica class = Iris-virginica