Train the Iris data and Test the Model

• Iris is perhaps the best known database to be found in the pattern recognition literature.

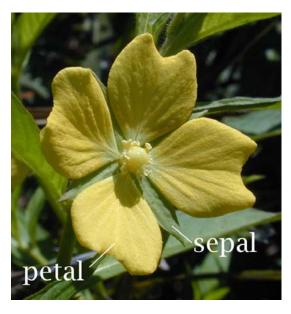


Figure 1: Flower showing petal and sepal [1].

- The data set contains 3 classes of 50 instances each, where each class refers to a type of iris plant.
- One class is linearly separable from the other 2; the latter are NOT linearly separable from each other.
 - Number of Instances: 150 (50 in each of three classes)
 - Number of Attributes/features: 4 numeric, predictive attributes and the class
 - Attribute Information:
 - 1. sepal length in cm
 - 2. sepal width in cm
 - 3. petal length in cm
 - 4. petal width in cm
 - 5. class:
 - o Iris Setosa
 - o Iris Versicolour
 - Iris Virginica

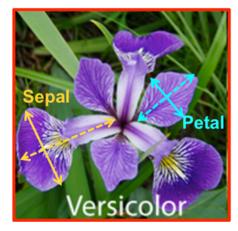






Figure 2: Three categories of Iris flowers [2].

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In [ ]: # Load pandas library. We want to use its DataFrame which supports tabular form.
import pandas as pd

In [ ]: # Load the dataset from the current directory into a DataFrame
iris=pd.read_csv("iris.arff")
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In []: # See the content of housing
 iris

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In [ ]: | # Want to see a few rows (actually 4, but 5 including the header)
        iris.head()
In [ ]: # Want to see a few last rows
        iris.tail()
In [ ]: # Information about the dataset
        iris.info()
In [ ]: # Some Statistical info. of the dataset
        iris.describe()
In [ ]: | # Want to see the column
        iris.columns
In [ ]: # I am intertested to see the unique values in the class column because I want to replace
        the text with the numeric values
        iris['class '].unique()
In [ ]: | # Visualizing pairwise relationships
        import seaborn as sns # for more on seaborn, see https://seaborn.pydata.org/
In [ ]: | sns.pairplot(iris); \#';' avoid outputing the internal location info here
In [ ]:
        # Want to see the histogram of the numerical columns using malplotlib
        import matplotlib.pyplot as plt
        %matplotlib inline
        iris.hist()
        plt.show()
In [ ]: | # I want to replace 'Iris-setosa' with 0, 'Iris-versicolor' with 1, 'Iris-virginica' with
        iris.replace("Iris-setosa",0)
In [ ]: | # But the above table is a view - and the replacement will not be a permanent change [we n
        eed to use option: inplace=True]
        iris
In [ ]: # We also do not want to change the original dataset, so we make a copy
        iriscp=iris.copy()
        iriscp
In [ ]: iriscp.replace(to_replace="Iris-setosa", value=0, inplace=True)
        #iriscp.replace("Iris-setosa",0,inplace=True) # This will work as well
In [ ]: iriscp
In [ ]: # Instead of replace them one-by-one I want to replace them all at once
        # So I make a dictionary (dict) first
        myreplacementlist= {"Iris-setosa":0, "Iris-versicolor":1,"Iris-virginica":2}
In [ ]: myreplacementlist
        # Note: I want the replacement to work only for column 'class'
In [ ]: # Testing the dict
        myreplacementlist['Iris-versicolor']
In [ ]: | iriscp.replace({'class ': myreplacementlist}, inplace=True)
In [ ]: iriscp
In [ ]: | # Now, all columns are numerical column - I want to run the pairplot again
        sns.pairplot(iriscp);
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sns.relplot(x='petal_length',y='petal_width',data=iriscp, hue='class ', style='class ');
In [ ]: # I want to save this table into a file
        iriscp.to_csv('myiriscp.csv')
In [ ]: | # read the file to check whether it is saved or not
        !cat 'myiriscp.csv'
In [ ]: # I can also use window's type command
        !type myiriscp.csv
In [ ]: # Now, I see index of each row is saved as well, which is a new item and it is now the 1st
        column of the table.
        # I do not want the index column to be saved - so, I use a modified command below:
        iriscp.to_csv('myiriscp_nonewcolumn.csv', index=False) # So, I made the index=False and it
        worked (see below)
In [ ]: # I want to read from this file - which I might need to do in future.
        # I am reading in, say, 'irisnewcp' DataFrame
        irisnewcp = pd.read_csv('myiriscp_nonewcolumn.csv')
        irisnewcp
In [ ]: # I see the index column above but not in the file - so it is created on the fly. Check th
        e columns:
        irisnewcp.columns
In [ ]: # I see the dataset is originally sorted (class-wise), which is not a good idea for machin
        e learning - let us unsort it
        from sklearn.utils import shuffle # NOTE: sklearn (Scikit-learn) will be our main Machine
        Learning python library
In [ ]: | irisnewcp_sh=shuffle(irisnewcp, random_state=345) # 'random_state' is used for initializin
        g the internal random number generator
In [ ]: | irisnewcp_sh
In [ ]: X=irisnewcp_sh.iloc[:,0:4] # 'iloc' is integer index based, so you have to specify rows an
        d columns by their integer value of the index
In [ ]: y=irisnewcp_sh.iloc[:,4:5]
In [ ]: y
In [ ]: | # Let us use kNN, with k=5, from sklearn
        from sklearn.neighbors import KNeighborsClassifier
In [ ]: # create an instance of KNeighborsClassifier along with necessary parameters
        knn = KNeighborsClassifier(n_neighbors=5)
In [ ]: # print the instance variable to see the parameters of knn
        print(knn)
In [ ]: # Train the classifier with the dataset
        knn.fit(X,y)
In [ ]: | # Since index column (& header) is a problem now, I need to drop the index column (& heade
        r) from both X and y
        # Also, sklearn expects X, y in array
        X=X.values.tolist() # 'values' are the content without the header and index of the DataFra
        me. toList converts into array
        Χ
```

In []: | # Want to examine how features help separate classes

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In []: # flatten() will remove the header and will convert y in a 1d array.
    #You can also use .ravel(). .ravel() returns a view and .flatten() return a copy
    y=y.values.flatten()
    Y

In []: # Now, try to train again
    knn.fit(X,y)

In []: # create sample test dataset => expected answers are: 0, 2, 1
    X_test = [4.8, 2.9, 1.54, 0.15], [5.9, 2.5, 5.5, 1.2], [5.9, 3.0, 4.6, 1.4]

In []: # predict the class to which the sample falls into
    knn.predict(X_test)
```

Save and Load the Model

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In [ ]: # Python pickle module is used for serializing and de-serializing a Python object structur
        import pickle
        # Note: you can also use joblib
        # joblib is optimized to be fast and robust on large data in particular
        # to write use 'joblib.dump' & to read use 'joblib.load'
In [ ]: # Save the model
        f1=open('iris_saved_knn_model','wb') # wb => write binary
        pickle.dump(knn, f1)
In []: # better close (or flush) a file when done.
        f1.close()
In [ ]: # Load the model & Test
        f2=open('iris_saved_knn_model', 'rb')
        loaded_model = pickle.load(f2)
In []: X_test = [4.8, 2.9, 1.54, 0.15], [5.9, 2.5, 5.5, 1.2], [5.9, 3.0, 4.6, 1.4]
In [ ]: loaded_model.predict(X_test)
In [ ]: # If you know the test answers and want to compute the accuracy then do the following
        Y \text{ test}=[0,2,1]
        accuracy = loaded_model.score(X_test, Y_test)
In [ ]: print(accuracy)
In [ ]: f2.close()
```

References:

[1] https://en.wikipedia.org/wiki/Sepal (https://en.wikipedia.org/wiki/Sepal)

[2] http://suruchifialoke.com/2016-10-13-machine-learning-tutorial-iris-classification/ (http://suruchifialoke.com/2016-10-13-machine-learning-tutorial-iris-classification/)