```
In [7]: import numpy as np
          import pandas as pd
          import matplotlib.pyplot as plt
          from sklearn.svm import SVC #import Support vector classifier
          from sklearn.model selection import train test split #split train test data
 In [8]: #read the iris data
          df = pd.read_csv('IRIS.csv')
          df.head()
 Out[8]:
             sepal_length sepal width petal_length petal_width
                                                                    species
          0
                      5.1
                                   3.5
                                                1.4
                                                             0.2 Iris-setosa
                      4.9
          1
                                   3.0
                                                1.4
                                                             0.2 Iris-setosa
          2
                      4.7
                                   3.2
                                                1.3
                                                             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
 In [9]: #check for null values
          df.isnull().sum()
 Out[9]: sepal length
                           0
          sepal_width
                           0
          petal_length
                           0
          petal width
                           0
          species
                           0
          dtype: int64
In [12]:
          #Label encoding
          from sklearn.preprocessing import LabelEncoder
          labels = ['species']
          for label in labels:
              label_encoder = LabelEncoder()
              label_encoder.fit(df[label])
          df[label] = label_encoder.transform(df[label])
          df.head()
Out[12]:
             sepal_length sepal_width petal_length petal_width species
          0
                      5.1
                                   3.5
                                                1.4
                                                             0.2
                                                                       0
          1
                      4.9
                                   3.0
                                                1.4
                                                             0.2
                                                                       0
          2
                                   3.2
                                                                       0
                      4.7
                                                1.3
                                                             0.2
          3
                                                1.5
                                                             0.2
                      4.6
                                   3.1
                                                                       0
                                                                       0
                      5.0
                                   3.6
                                                             0.2
          4
                                                1.4
In [13]: print(df['species'])
```

```
0
               0
        2
        3
                0
                0
        145
               2
                2
        146
                2
        147
        148
                2
        149
                2
        Name: species, Length: 150, dtype: int64
In [15]:
          # separate independent and dependent variable
          x = df.iloc[:,0:3] # sepal length-width, petal length-width
          y = df.iloc[:,-1:] #species
In [16]:
         y.head()
Out[16]:
             species
          0
                  0
          1
                  0
          2
                  0
          3
                  0
          4
                  0
In [17]:
          #split the data into train and test set
          from sklearn.model_selection import train_test_split
          X_train, X_test, y_train, y_test = train_test_split(x, y, test_size=0.3, random_sta
Out[17]: X_train hardength sepal_width
                                                 3.5
           60
                        5.0
                                    2.0
          116
                        6.5
                                    3.0
                                                 5.5
          144
                        6.7
                                    3.3
                                                 5.7
          119
                        6.0
                                    2.2
                                                 5.0
          108
                        6.7
                                    2.5
                                                 5.8
In [18]:
          y_train.head()
```

```
Out[18]: species

60 1

116 2

144 2

119 2

108 2
```

```
In [19]: from sklearn.svm import LinearSVC
    clf = LinearSVC() #by default LinearSVC follows one -vs- rest approach.
    #train the alorithm or model
    clf.fit(X_train,y_train)
    #test the model
    yp = clf.predict(X_test)
    print (yp)
    print ("accuracy is ", clf.score(X_test,y_test)*100)

[2 1 0 2 0 2 0 1 1 1 2 1 2 1 2 0 1 2 0 0 2 2 0 0 2 0 0 1 1 0 2 2 0 2 2 1 0
    2 1 1 2 0 1 0 0]
```

```
In [20]: from sklearn import svm
#SVC that follows one-vs-one approch
    clf = svm.SVC(decision_function_shape='ovo')
    clf.fit(X_train, y_train)
    y_pred = clf.predict(X_test)
    print(yp)
    print("accuracy is ", clf.score(X_test,y_pred)*100)
[2 1 0 2 0 2 0 1 1 1 2 1 2 1 2 0 1 2 0 0 2 2 0 0 2 0 0 1 1 0 2 2 0 2 2 1 0
    2 1 1 2 0 1 0 0]
```

```
In [ ]:
```