HELP MANUAL

SVM Classification on Lists

SVM Classification on Lists

- Required packages to implement SVM
 Classification algorithm on Lists
 - import numpy as np
 - import matplotlib.pyplot as plt
 - from sklearn import svm

SVM Classification on Lists Contd...

- Creating Lists and displaying them
 - $\square x=[1, 5, 1.5, 8, 1, 9]$
 - y=[2, 8, 1.8, 8, 0.6, 11]
 - print (x)
 - print (y)
- Plot and display scatter chart of x and y
 - plt.scatter(x,y)
 - plt.show()
- Creating an array X which stores pair (x, y)
 - \square X = np.array([[1,2], [5,8], [1.5,1.8], [8,8], [1,0.6], [9,11]])
- □ Create the target class Y as Y = [0,1,0,1,0,1]

SVM Classification on Lists Contd...

- Classifying list X and comparing with target Y
 - clf = svm.SVC(kernel='linear', C = 1.0) # Linear Kernel
 and SVC method
 - clf.fit(X,Y)
 - print(clf.predict([0.58,0.76]))
 - print(clf.predict([10.58,10.76]))
 - \square w = clf.coef_[0]
 - print("weight",w)
 - a = -w[0] / w[1]
 - print ("Bias",a)

SVM Classification on Lists Contd...

Visualizing Results

- $\square xx = np.linspace(0,12)$
- print ("xx",xx)
- \square yy = a * xx clf.intercept_[0] / w[1]
- print ("yy",yy)
- h0 = plt.plot(xx, yy, 'k-', label="SVM Linear Classifier Chart")
- \square plt.scatter(X[:, 0], X[:, 1], c = y)
- plt.legend()
- plt.show()

SVM Classification on Dataset

SVM Classification on Dataset

- Required packages to implement SVM
 Classification algorithm on datasets
 - from sklearn import datasets
 - □import numpy as np
 - import matplotlib.pyplot as plt
 - from sklearn import svm

Loading dataset

- iris_dataset = datasets.load_iris()
- print("Iris data set Description :: ",
 iris_dataset['DESCR'])
- print ("Iris feature data :: ",
 iris_dataset['data'])
- print ("Iris target :: ", iris_dataset['target'])

Visualizing dataset for sepal attribute

- def visuvalize_sepal_data():
 - iris = datasets.load_iris()
 - X = iris.data[:,:2] # we only take the first two features
 - y = iris.target
 - plt.scatter(X[:, 0], X[:, 1], c=y, cmap=plt.cm.coolwarm)
 - plt.xlabel('Sepal length')
 - plt.ylabel('Sepal width')
 - plt.title('Sepal Width & Length')
 - plt.show()

Visualizing dataset for petal attribute

- def visuvalize_petal_data():
 - iris = datasets.load_iris()
 - X = iris.data[:, 2:] # we only take the last two features
 - y = iris.target
 - plt.scatter(X[:, 0], X[:, 1], c=y, cmap=plt.cm.coolwarm)
 - plt.xlabel('Petal length')
 - plt.ylabel('Petal width')
 - plt.title('Petal Width & Length')
 - plt.show()

Classification based on Sepal attribute

- iris = datasets.load_iris()
- X = iris.data[:, :2] # we only take the Sepal two features
- y = iris.target
- □ C = 1.0 # SVM regularization parameter
- Linear Kernel
 - svc = svm.SVC(kernel='linear', C=C).fit(X, y) #SVC function
 - lin_svc = svm.LinearSVC(C=C).fit(X, y) # Linear SVC
- rbf/poly Kernel
 - rbf_svc = svm.SVC(kernel='rbf', gamma=0.7, C=C).fit(X, y)
 - poly_svc = svm.SVC(kernel='poly', degree=3, C=C).fit(X, y)

Creating Meshgrid

- \blacksquare h = .02 #step size in the mesh
- # create a mesh to plot in
- \blacksquare x_min, x_max = X[:, 0].min() 1, X[:, 0].max() + 1
- \square y_min, y_max = X[:, 1].min() 1, X[:, 1].max() + 1
- xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h))

Visualizing Results

- for i, clf in enumerate((svc, lin_svc, rbf_svc, poly_svc)):
- # Plot the decision boundary assigning a color to each
- # point in the mesh [x_min, x_max]x[y_min, y_max]
- \square plt.subplot(2, 2, i + 1)
- plt.subplots_adjust(wspace=0.4, hspace=0.4)
- $Z = clf.predict(np.c_[xx.ravel(), yy.ravel()])$
- # Put the result into a color plot
- \Box Z = Z.reshape(xx.shape)
- plt.contourf(xx, yy, Z, cmap=plt.cm.coolwarm, alpha=0.8)