Experiment 6: Support Vector Machines

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Class: TE EXTC

```
import pandas as pd
import numpy as np
import matplotlib as mpl
import matplotlib.pyplot as plt
from sklearn.metrics import confusion matrix
%matplotlib inline
# We'll define a function to draw a nice plot of an SVM
def plot_svc(svc, X, y, h=0.02, pad=0.25):
    x_{min}, x_{max} = X[:, 0].min()-pad, <math>X[:, 0].max()+pad
    y_{min}, y_{max} = X[:, 1].min()-pad, <math>X[:, 1].max()+pad
    xx, yy = np.meshgrid(np.arange(x_min, x_max, h), np.arange(y_min, y_max, h))
    Z = svc.predict(np.c_[xx.ravel(), yy.ravel()])
    Z = Z.reshape(xx.shape)
    plt.contourf(xx, yy, Z, cmap=plt.cm.Paired, alpha=0.2)
    plt.scatter(X[:,0], X[:,1], s=70, c=y, cmap=mpl.cm.Paired)
    # Support vectors indicated in plot by vertical lines
    sv = svc.support_vectors_
    plt.scatter(sv[:,0], sv[:,1], c='k', marker='x', s=100, linewidths='1')
    plt.xlim(x min, x max)
    plt.ylim(y_min, y_max)
    plt.xlabel('X1')
    plt.ylabel('X2')
    plt.show()
    print('Number of support vectors: ', svc.support .size)
```

Above is a function to make a plot of the support vector machines

```
from sklearn.svm import SVC

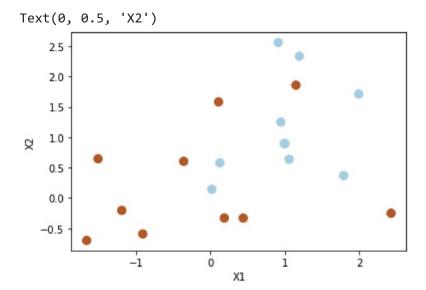
# Generating random data: 20 observations of 2 features and divide into two classes.
np.random.seed(5)

X = np.random.randn(20,2)
y = np.repeat([1,-1], 10)
```

$$X[y == -1] = X[y == -1]+1$$

Generating 20 Observations randomly and seperating them into classes.

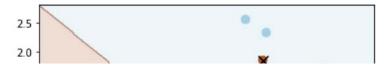
```
plt.scatter(X[:,0], X[:,1], s=70, c=y, cmap=mpl.cm.Paired)
plt.xlabel('X1')
plt.ylabel('X2')
```



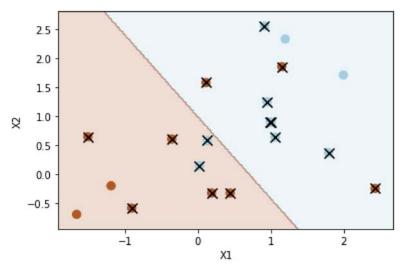
Plotting the randomly generated points

```
svc = SVC(C=1, kernel='linear')
svc.fit(X, y)
SVC(C=1, kernel='linear')
```

Making the SVC of the generated points with C = 1 and the kernel as linear



We can see that when C = 1 No. of support vectrs is 13



Number of support vectors: 16

For C = 0.1 no. of support vectors is 16

We wan to see which C parameter is the best to make our support vector machine so we pass tuned parameters which have multiple C values. After we obtained the results we will see which C value is the best

```
clf.cv_results_
     {'mean fit time': array([0.00146143, 0.00114441, 0.00138247, 0.0006315 , 0.00043905,
             0.00045528, 0.00056968]),
      'mean score time': array([0.00058742, 0.00046158, 0.00035613, 0.00028646, 0.00025413,
             0.00025156, 0.00025833]),
      'mean_test_score': array([0.8 , 0.8 , 0.8 , 0.75, 0.75, 0.75, 0.75]),
      'param C': masked array(data=[0.001, 0.01, 0.1, 1, 5, 10, 100],
                   mask=[False, False, False, False, False, False],
             fill value='?',
                  dtype=object),
      'params': [{'C': 0.001},
       {'C': 0.01},
       {'C': 0.1},
       {'C': 1},
       {'C': 5},
       {'C': 10},
       {'C': 100}],
      'rank test score': array([1, 1, 1, 4, 4, 4, 4], dtype=int32),
      'split0 test score': array([0.5, 0.5, 0.5, 0.5, 0.5, 0.5, 0.5]),
      'split1 test score': array([0.5, 0.5, 0.5, 0. , 0. , 0. , 0. ]),
      'split2 test score': array([0.5, 0.5, 0.5, 0.5, 0.5, 0.5, 0.5]),
      'split3_test_score': array([1., 1., 1., 1., 1., 1., 1.]),
      'split4_test_score': array([1., 1., 1., 1., 1., 1., 1.]),
      'split5_test_score': array([1., 1., 1., 1., 1., 1., 1.]),
      'split6_test_score': array([1., 1., 1., 1., 1., 1., 1.]),
      'split7_test_score': array([1., 1., 1., 1., 1., 1., 1.]),
      'split8_test_score': array([0.5, 0.5, 0.5, 0.5, 0.5, 0.5]),
      'split9 test score': array([1., 1., 1., 1., 1., 1., 1.]),
      'std_fit_time': array([1.53381734e-03, 1.64096783e-03, 2.21696624e-03, 4.63546288e-04,
             1.01731003e-05, 2.38197224e-05, 4.70846453e-05]),
      'std score time': array([4.67495911e-04, 3.81544568e-04, 2.79883611e-05, 4.18144660e-0!
             8.33814231e-06, 6.84178244e-06, 1.01926382e-05]),
      'std_test_score': array([0.24494897, 0.24494897, 0.24494897, 0.3354102, 0.3354102,
             0.3354102 , 0.3354102 ])}
clf.best_params_
     {'C': 0.001}
We can see that the best C value is C = 0.001
np.random.seed(1)
X test = np.random.randn(20,2)
y test = np.random.choice([-1,1], 20)
X \text{ test[y test == 1] = } X \text{ test[y test == 1]-1}
```

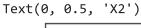
We again generate 20 random numbers

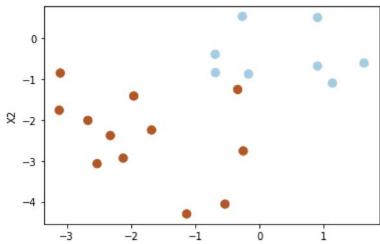
```
svc2 = SVC(C=0.001, kernel='linear')
svc2.fit(X, y)
y_pred = svc2.predict(X_test)
pd.DataFrame(confusion_matrix(y_test, y_pred), index=svc2.classes_, columns=svc2.classes_)
```

```
-1 1
-1 2 6
1 0 12
```

We make SVC using C = 0.01 since it is the best C value.

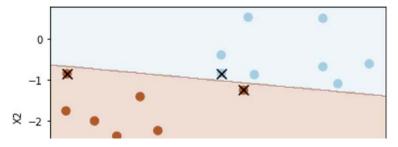
```
X_test[y_test == 1] = X_test[y_test == 1] -1
plt.scatter(X_test[:,0], X_test[:,1], s=70, c=y_test, cmap=mpl.cm.Paired)
plt.xlabel('X1')
plt.ylabel('X2')
```





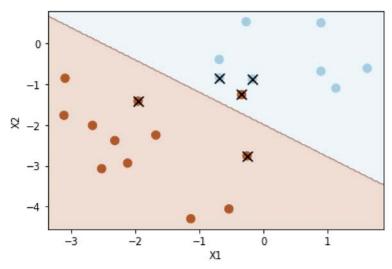
Plotting all the points generated

```
svc3 = SVC(C=1e5, kernel='linear')
svc3.fit(X_test, y_test)
plot_svc(svc3, X_test, y_test)
```



No we make the support vector line and we can see that for C = 1e5 No.of support vectors are 3

```
svc4 = SVC(C=1, kernel='linear')
svc4.fit(X_test, y_test)
plot_svc(svc4, X_test, y_test)
```



Number of support vectors: !

We make the support vector line and we can see that for C = 1 No.of support vectors are 5

Using cost = 1, we misclassify a training observation, but we also obtain a much wider margin and make use of five support vectors. It seems likely that this model will perform better on test data than the model with cost = 1e5.

```
from sklearn.model_selection import train_test_split

np.random.seed(8)

X = np.random.randn(200,2)

X[:100] = X[:100] +2

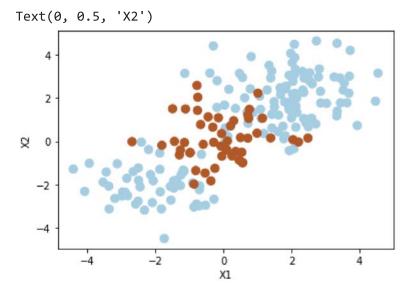
X[101:150] = X[101:150] -2

y = np.concatenate([np.repeat(-1, 150), np.repeat(1,50)])

X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.5, random_state=2)

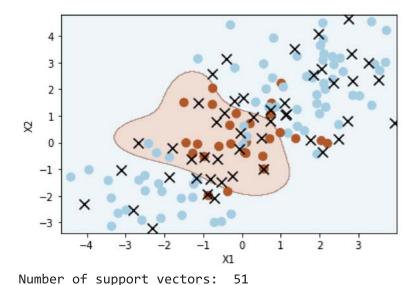
plt.scatter(X[:,0], X[:,1], s=70, c=y, cmap=mpl.cm.Paired)
```

```
plt.xlabel('X1')
plt.ylabel('X2')
```



Random data points are generated which belong to 2 seperate classes that are linearly not seperable. Since one class is stuck between another class we will have to use the radial kernel method of SVM with gamma = 1.

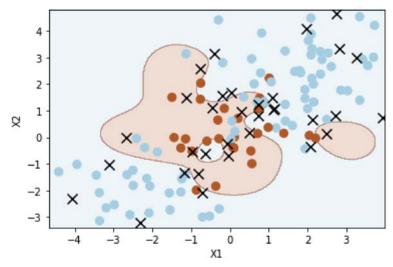
```
svm = SVC(C=1.0, kernel='rbf', gamma=1)
svm.fit(X_train, y_train)
plot_svc(svm, X_test, y_test)
```



We cash see that there are alot of errors and the two classes are not seperated. It can be improved by increasing the cost from 1 to 100. The number of support vectors is 51

Increasing C parameter, allowing more flexibility

```
svm2 = SVC(C=100, kernel='rbf', gamma=1.0)
svm2.fit(X_train, y_train)
plot_svc(svm2, X_test, y_test)
```

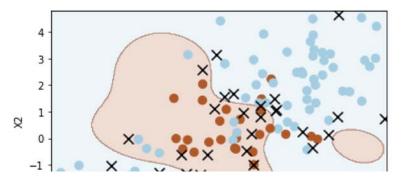


Number of support vectors: 36

The number of support vectors is 36. The cost is increased but because of that the boundary is very irregular.

With the help of gridsearchCV() we can find the best combination for the value of C and Gamma. We find out that the best values are C = 10 and Gamma = 0.5

```
plot_svc(clf.best_estimator_, X_test, y_test)
print(confusion_matrix(y_test, clf.best_estimator_.predict(X_test)))
print(clf.best_estimator_.score(X_test, y_test))
```



On applying the best parameters to our radial kernel SVM i.e. C = 10 and Gamma = 0.5 we get the above plot. The number of support vectors is 32 in this case.

```
from sklearn.metrics import auc
from sklearn.metrics import roc_curve
```

We now are plotting the ROC curve so we import auc and roc_curve

```
# More constrained model
svm3 = SVC(C=1, kernel='rbf', gamma=1)
svm3.fit(X_train, y_train)
SVC(C=1, gamma=1)
```

We fit the more constrained model with C = 1 and Gamma = 1

```
# More flexible model
svm4 = SVC(C=1, kernel='rbf', gamma=50)
svm4.fit(X_train, y_train)
SVC(C=1, gamma=50)
```

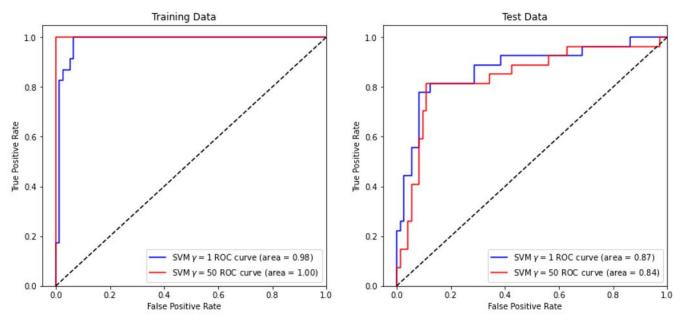
We try to fit the more flexible model cost = 1 and gamma = 50

```
y_train_score3 = svm3.decision_function(X_train)
y_train_score4 = svm4.decision_function(X_train)

y_train_score3 = svm3.decision_function(X_train)
y_train_score4 = svm4.decision_function(X_train)

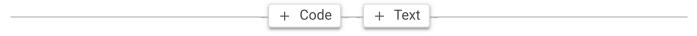
false_pos_rate3, true_pos_rate3, _ = roc_curve(y_train, y_train_score3)
roc_auc3 = auc(false_pos_rate4, _ = roc_curve(y_train, y_train_score4)
roc_auc4 = auc(false_pos_rate4, true_pos_rate4)
```

```
fig, (ax1,ax2) = plt.subplots(1, 2, figsize=(14,6))
ax1.plot(false_pos_rate3, true_pos_rate3, label='SVM $\gamma = 1$ ROC curve (area = %0.2f)' %
ax1.plot(false pos rate4, true pos rate4, label='SVM $\gamma = 50$ ROC curve (area = %0.2f)'
ax1.set_title('Training Data')
y test score3 = svm3.decision function(X test)
y_test_score4 = svm4.decision_function(X_test)
false_pos_rate3, true_pos_rate3, _ = roc_curve(y_test, y_test_score3)
roc auc3 = auc(false pos rate3, true pos rate3)
false_pos_rate4, true_pos_rate4, _ = roc_curve(y_test, y_test_score4)
roc auc4 = auc(false pos rate4, true pos rate4)
ax2.plot(false pos rate3, true pos rate3, label='SVM $\gamma = 1$ ROC curve (area = %0.2f)' %
ax2.plot(false pos rate4, true pos_rate4, label='SVM $\gamma = 50$ ROC curve (area = %0.2f)'
ax2.set title('Test Data')
for ax in fig.axes:
    ax.plot([0, 1], [0, 1], 'k--')
    ax.set_xlim([-0.05, 1.0])
    ax.set_ylim([0.0, 1.05])
    ax.set xlabel('False Positive Rate')
    ax.set_ylabel('True Positive Rate')
    ax.legend(loc="lower right")
```



Here we can see that we have recieved the ROC curves. For the training data we can see that case with gamma = 50 gives better result (area = 1.00 compared to 0.98 for the other case) while for

testing data gamma = 1 gives a better result as compared to gamma = 50 (area = 0.87 as compared to 0.84 in other case



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