

# hw1-yuquan-cai-q2

March 7, 2025

## 1 Linear SVM

```
[18]: %matplotlib inline
import matplotlib
import matplotlib.pyplot as plt

from sklearn.svm import SVC
from sklearn import datasets

# Dataset
iris = datasets.load_iris()
X = iris["data"][:, (2, 3)] # petal length, petal width
y = iris["target"]

setosa_or_versicolor = (y == 0) | (y == 1)
X = X[setosa_or_versicolor]
y = y[setosa_or_versicolor]

# SVM Classifier model
svm_clf = SVC(kernel="linear", C=float("inf"))
svm_clf.fit(X, y)

# Usage Example:
# SVM classifiers do not output a probability like logistic regression
svm_clf.predict([[2.4, 3.1]])
```

```
[18]: array([1])
```

```
[19]: def plot_svc_decision_boundary(svm_clf, xmin, xmax):
    w = svm_clf.coef_[0]
    b = svm_clf.intercept_[0]
    x0 = np.linspace(xmin, xmax, 200)

    #####
    ## WRITE YOUR CODE HERE (10 Points) ##
    #####
```

```

# Hint: at the decision boundary,  $w_0x_0 + w_1x_1 + b = 0$ ,
# write down the formula of  $x_1$  as decision_boundary here

decision_boundary = -(w[0] * x0 + b) / w[1] #
margin = 1 / np.linalg.norm(w) #

gutter_up = decision_boundary + margin
gutter_down = decision_boundary - margin

svs = svm_clf.support_vectors_
plt.scatter(svs[:, 0], svs[:, 1], s=180, facecolors='#FFAAAA')
plt.plot(x0, decision_boundary, "k-", linewidth=2)
plt.plot(x0, gutter_up, "k--", linewidth=2)
plt.plot(x0, gutter_down, "k--", linewidth=2)

```

```

[20]: #Plot the decision boundaries
import numpy as np
from sklearn.preprocessing import StandardScaler

plt.figure(figsize=(12,3.2))
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
svm_clf.fit(X_scaled, y)

plt.plot(X_scaled[:, 0][y==1], X_scaled[:, 1][y==1], "bo")
plt.plot(X_scaled[:, 0][y==0], X_scaled[:, 1][y==0], "ms")
plot_svc_decision_boundary(svm_clf, -2, 2)

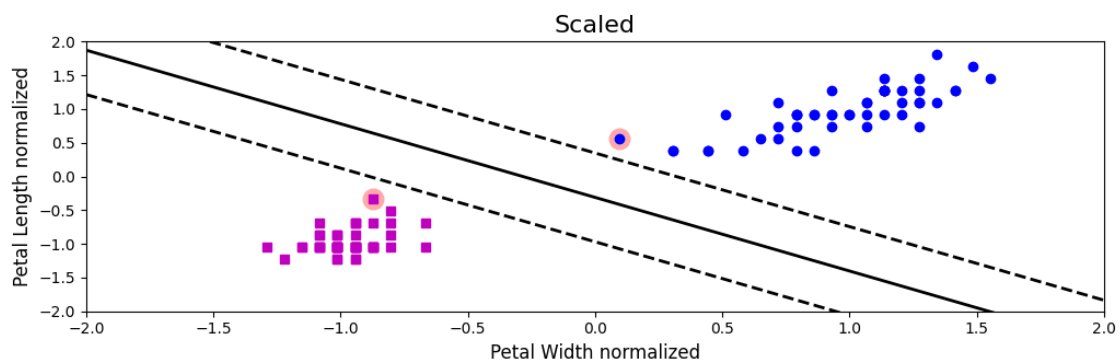
plt.xlabel("Petal Width normalized", fontsize=12)
plt.ylabel("Petal Length normalized", fontsize=12)
plt.title("Scaled", fontsize=16)
plt.axis([-2, 2, -2, 2])

```

```

[20]: (np.float64(-2.0), np.float64(2.0), np.float64(-2.0), np.float64(2.0))

```



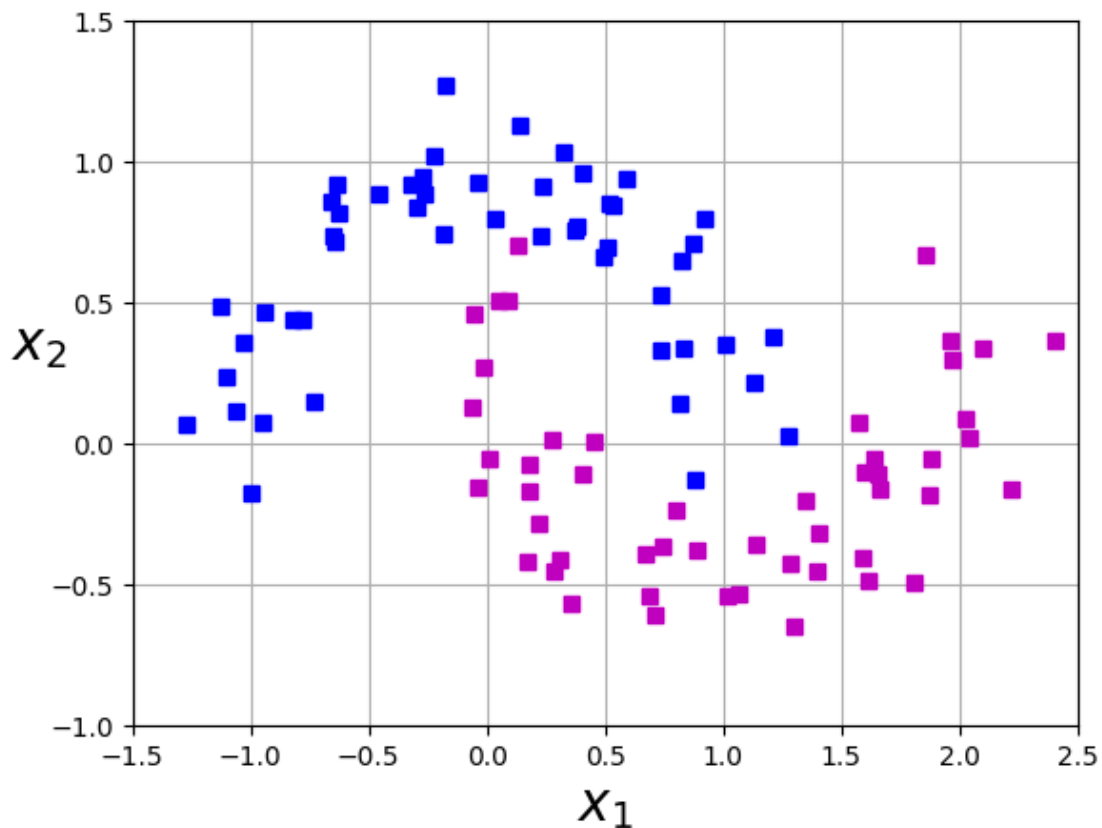
## 2 Non Linear SVM

```
[21]: from sklearn.datasets import make_moons
from sklearn.pipeline import Pipeline
from sklearn.preprocessing import PolynomialFeatures
from sklearn.svm import LinearSVC
```

```
[22]: # Construct some test data
from sklearn.datasets import make_moons
X, y = make_moons(n_samples=100, noise=0.15, random_state=42)

# Define a function to plot the dataset
def plot_dataset(X, y, axes):
    plt.plot(X[:, 0][y==0], X[:, 1][y==0], "bs")
    plt.plot(X[:, 0][y==1], X[:, 1][y==1], "ms")
    plt.axis(axes)
    plt.grid(True, which='both')
    plt.xlabel(r"$x_1$", fontsize=20)
    plt.ylabel(r"$x_2$", fontsize=20, rotation=0)

# Let's have a look at the data we have generated
plot_dataset(X, y, [-1.5, 2.5, -1, 1.5])
plt.show()
```



```
[23]: # Define a function plot the decision boundaries
```

```
def plot_predictions(clf, axes):  
    # create data in continous linear space  
    x0s = np.linspace(axes[0], axes[1], 100)  
    x1s = np.linspace(axes[2], axes[3], 100)  
    x0, x1 = np.meshgrid(x0s, x1s)  
    X = np.c_[x0.ravel(), x1.ravel()]  
    y_pred = clf.predict(X).reshape(x0.shape)  
    y_decision = clf.decision_function(X).reshape(x0.shape)  
    plt.contourf(x0, x1, y_pred, cmap=plt.cm.brg, alpha=0.2)  
    plt.contourf(x0, x1, y_decision, cmap=plt.cm.brg, alpha=0.1)
```

```
[24]: # Build the model and set hyperparameters  
# Create a pipeline to create features, scale data and fit the model
```

```
#####  
## WRITE YOUR CODE HERE (15 Points) ##  
#####  
  
polynomial_svm_clf = Pipeline([("poly_features", PolynomialFeatures(degree=3)),  
    ↪ ("scaler", StandardScaler()), ("svm_clf", LinearSVC(C=1, loss="hinge"))]) #  
    ↪ finish the pipeline  
  
# Hint:  
# Pipeline allows you to chain multiple preprocessing and modeling steps↪  
    ↪ together.  
# Step1 begin by applying polynomial feature transformation to the input data,↪  
    ↪ using a degree of 3 to capture non-linear relationships.  
# Step2: scale the data with a standard scaler to ensure that the features are↪  
    ↪ standardized  
# Step3: use a linear SVM classifier with hinge loss and appropriate↪  
    ↪ hyperparameter.  
  
# Once the pipeline is constructed, you can fit the model on the training data  
polynomial_svm_clf.fit(X,y)
```

```
[24]: Pipeline(steps=[('poly_features', PolynomialFeatures(degree=3)),  
    ('scaler', StandardScaler()),  
    ('svm_clf', LinearSVC(C=1, loss='hinge'))])
```

```
[25]: #plot the decision boundaries  
plt.figure(figsize=(11, 4))
```

```

#plot the decision boundaries
plot_predictions(polynomial_svm_clf, [-1.5, 2.5, -1, 1.5])

#plot the dataset
plot_dataset(X, y, [-1.5, 2.5, -1, 1.5])

plt.title(r"$d=3, C=10$", fontsize=18)
plt.show()

```

