

Support Vector Machine (SVM)

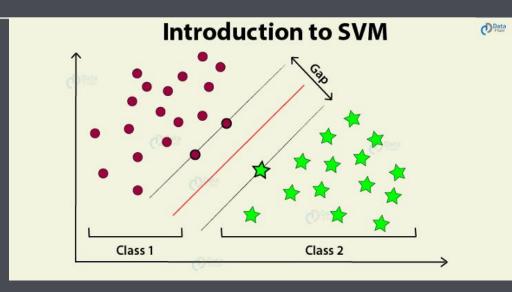
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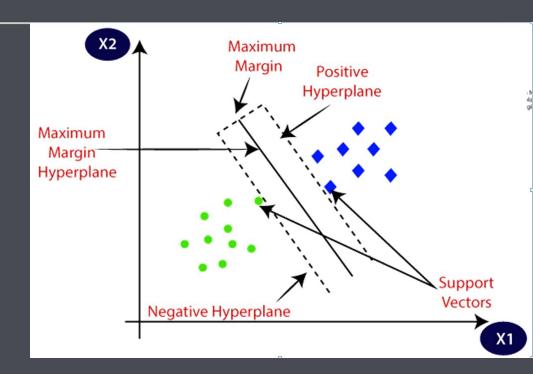
- > Supervised Learning method
- > Classification or Regression
- > Separates the data linearly by finding the best decision boundary
- > The Decision Boundary is called a Hyperplane in SVM
- > Can be linear... and non-linear (!) (we get to this in a sec)



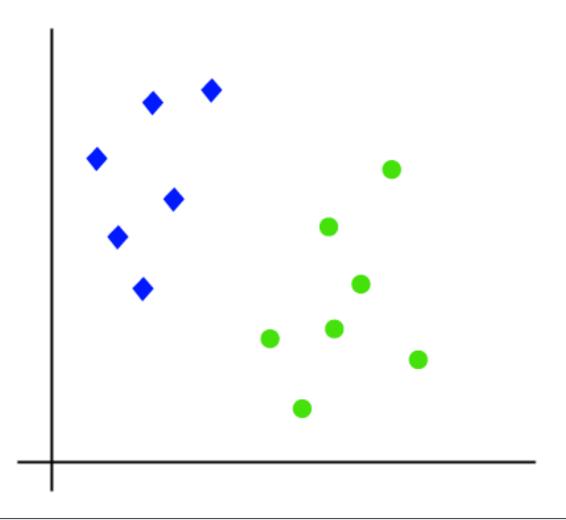


Support Vectors and Margins

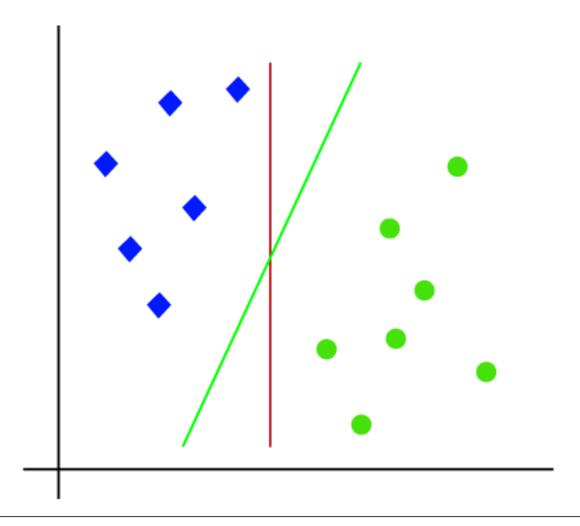
- > The hyperplanes are in the D-1 dimension:
 - > 2 features straight line (1D)
 - > 3 features 2D plane
- > Hyperplanes have a **maximum margin** from the points
- > SVM is looking to **maximize** this margin
- > The nearest points (or **vectors** array of points) to the margins are called the **support vectors**



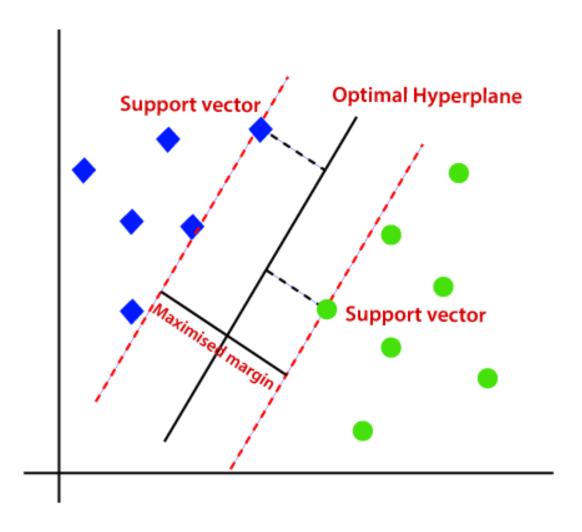
Example 1



Example 1



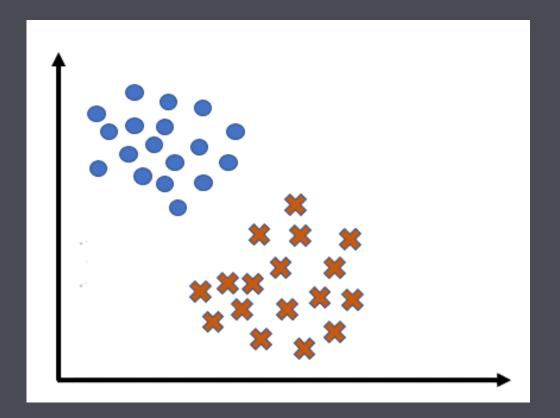
Example 1





SVM Margins

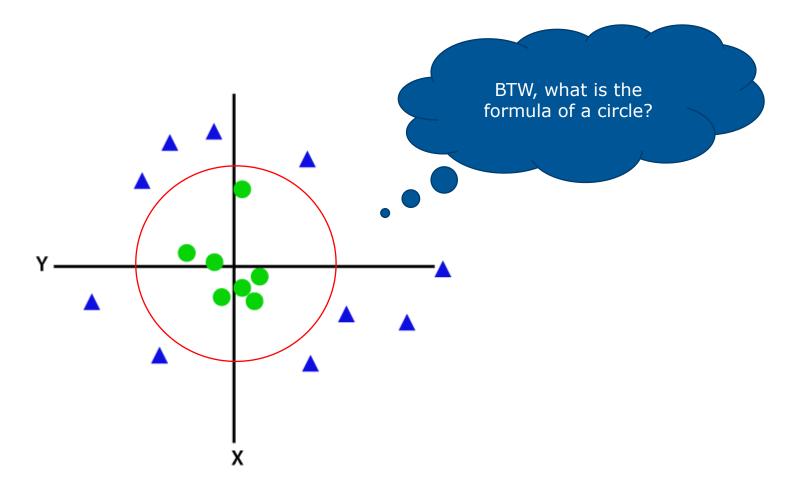
- > Bigger is better!
- > Notice: the objective is different: Instead of minimizing distances, we maximize the margins



Big deal when the data is linear...

Linear regression does the same, so why do we need this?!

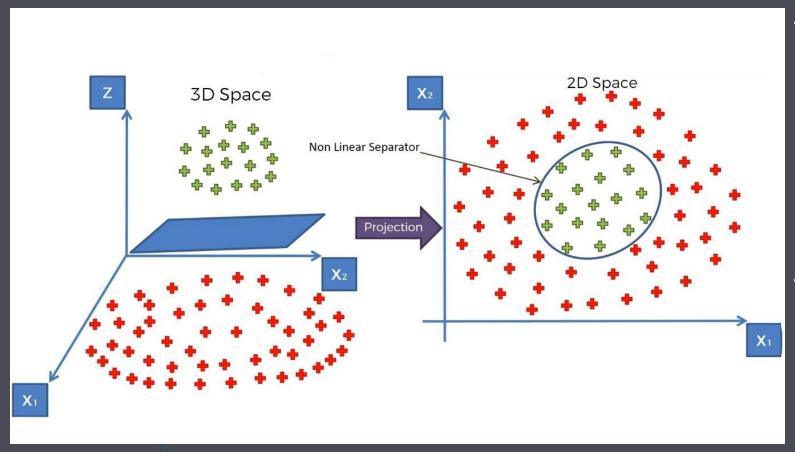
Non-Linearly Separable data



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We can add a dimension!

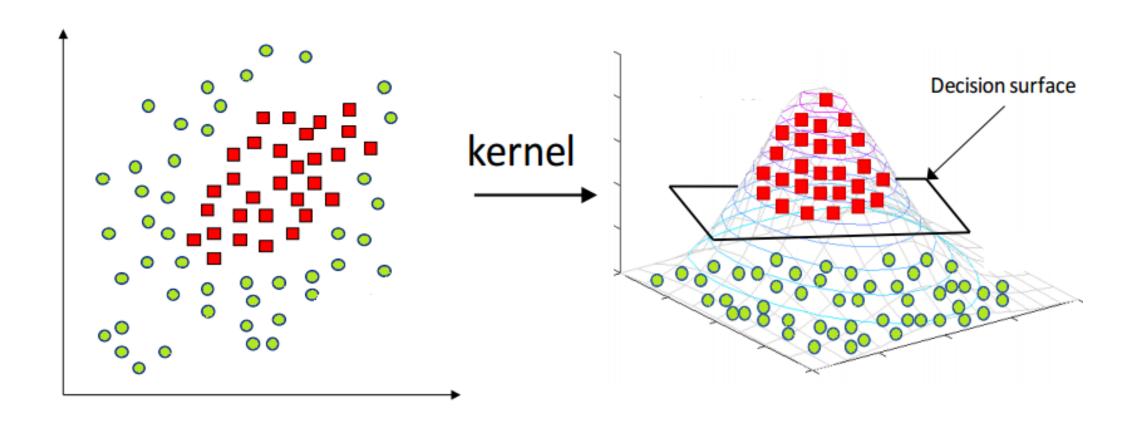


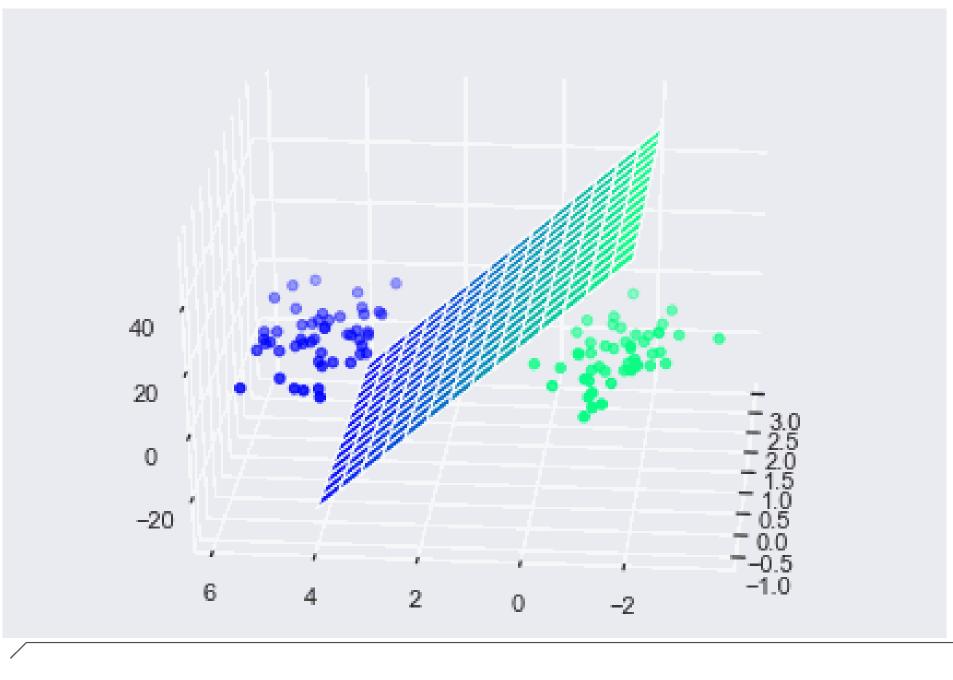
The new dimension – z can be created with some formula, e.g.:

$$z = x_1^2 + x_2^2$$

This formula is called a **kernel**

A kernel is a transformation function.





In 2D it finds a line.

In 3D it finds a hyperplane.



How can we possibly add a dimension?!

- > Dimensions aka **z-space**
- > Kernels
 - > A hyperparameter of the classifier

Some Kernel Examples:

> Linear Kernel: k(x,y) = sum(x,y)

> Polynomial Kernel: $k(x,y) = (x,y+1)^d$

> Gaussian Radial Basis Function (RBF): $k(x,y) = \exp(-\gamma * ||x-y||^2)$

> Sigmoid Kernel: $k(x,y) = \tanh(\gamma x^T y + r)$



Python code example

```
from sklearn.svm import SVC # Support Vector Classifier
classifier = SVC(kernel='linear', random_state=0)
classifier.fit(x_train, y_train)
```

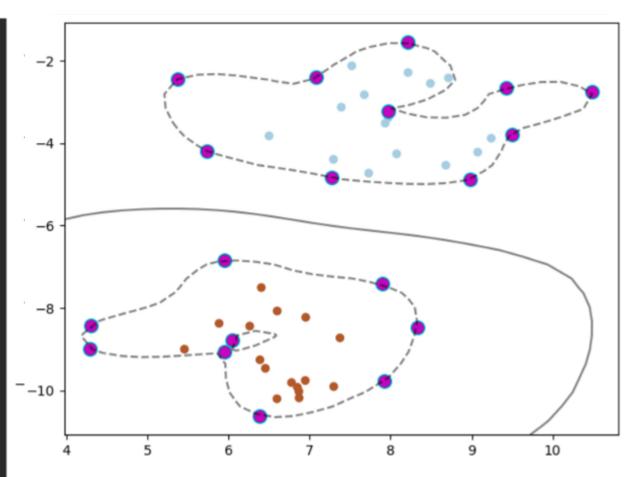
Other kernel options are:
 "poly"
 "rbf"
 "sigmoid"
 ...



Elaborated Code Example

```
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```

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn import svm
from sklearn.datasets import make_blobs
# we create 40 separable points
X, y = make_blobs(n_samples=50, centers=2, random_state=6)
# fit the model, don't regularize for illustration purposes
clf = svm.SVC(kernel='rbf'.
                            C=1000)
clf.fit(X, y)
plt.scatter(X[:, 0], X[:, 1], c=y, s=30, cmap=plt.cm.Paired)
# plot the decision function
ax = plt.gca()
xlim = ax.get_xlim()
ylim = ax.get_ylim()
# create grid to evaluate model
xx = np.linspace(xlim[0], xlim[1], 30)
yy = np.linspace(ylim[0], ylim[1], 30)
YY, XX = np.meshgrid(yy, xx)
xy = np.vstack([XX.ravel(), YY.ravel()]).T
Z = clf.decision_function(xy).reshape(XX.shape)
# plot decision boundary and margins
# plot support vectors
ax.scatter(clf.support_vectors_[:, 0], clf.support_vectors_[:, 1], s=100,
          linewidth=1, facecolors='m', edgecolors='c')
plt.show()
```





SVM - Summary

- > Up until NN this was the most efficient classifier (!)
- > It can solve many real-world problems
- > Works well with high dimensionality
- > Memory Efficient: It only uses the data points which are close to the decision boundary.
- > Can give an *intuition* about the probability or *certainty* of data points (how close they are to the boundary vector)
- > Bonus: It can be used in an *unsupervised* manner for anomaly classification (e.g., analyzing log-texts) called "one-class SVM"