



الجامعة السورية الخاصة
SYRIAN PRIVATE UNIVERSITY

المحاضرة العاشرة

كلية الهندسة المعلوماتية

مقرر تعلم الآلة

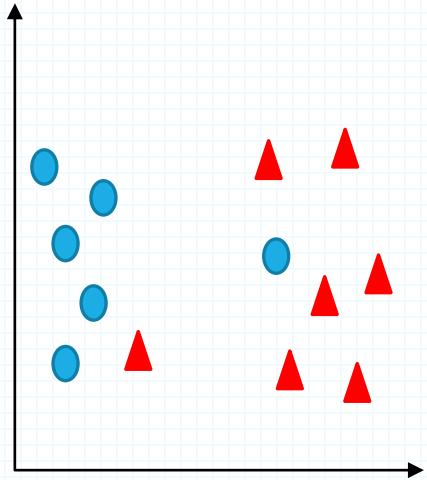
Support Vector Machine (SVM) 3

د. رياض سنبل

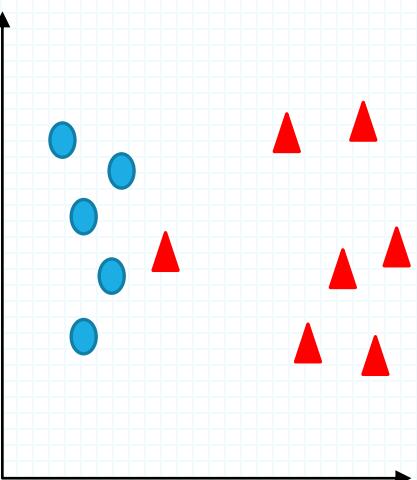
What if?

What are the problems of the current version for SVM?

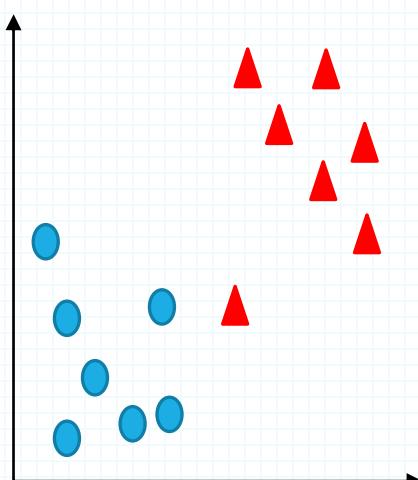
(A)



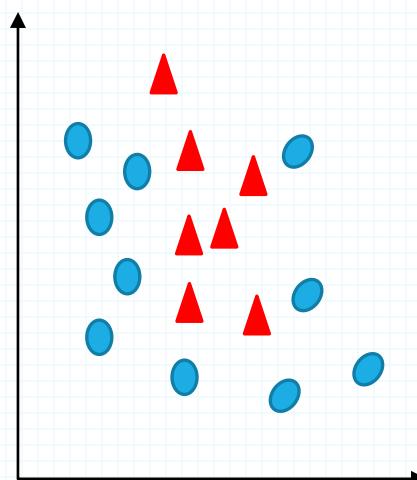
(B)



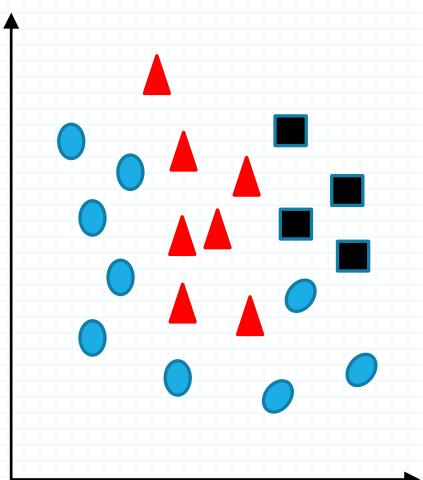
(C)



(D)



(E)



1st Improvement

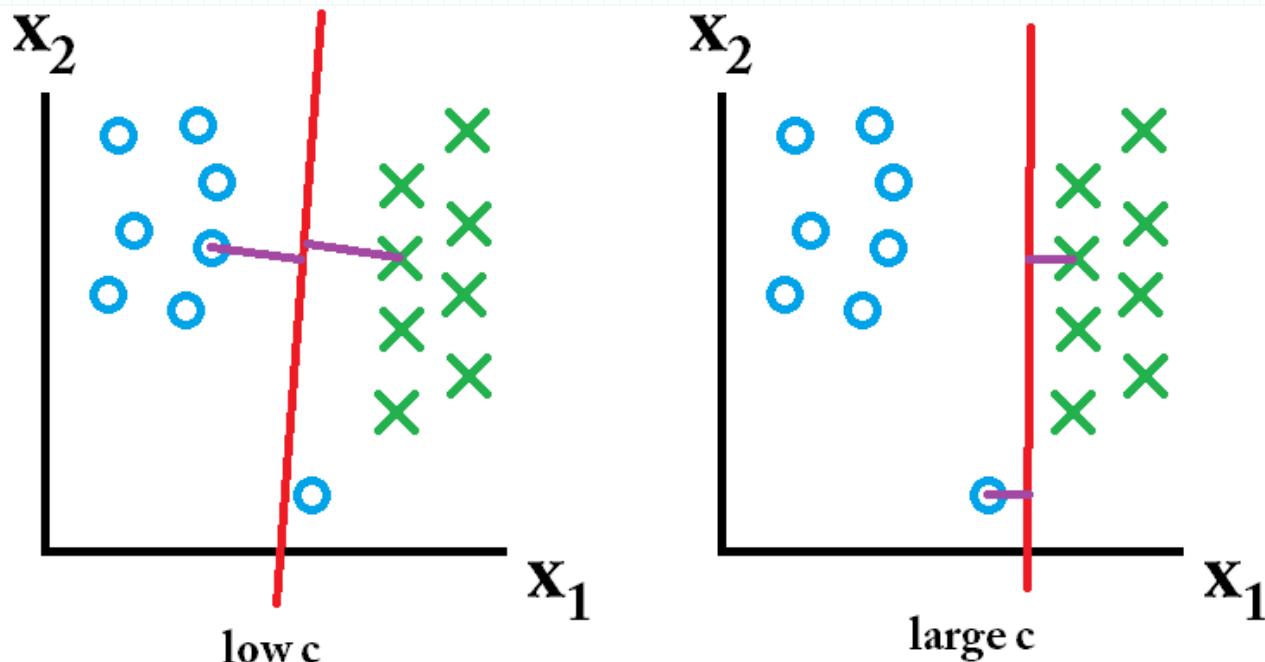
Soft Margin SVM

(allows few misclassifications)

C Hyper-parameter

- When C is high it will classify all the data points correctly, also there is a chance to overfit.
- SVM Error = Margin Error + Classification Error**

$$\operatorname{argmin}(w^*, b^*) \frac{\|w\|}{2} + c \sum_{i=1}^n \zeta_i$$

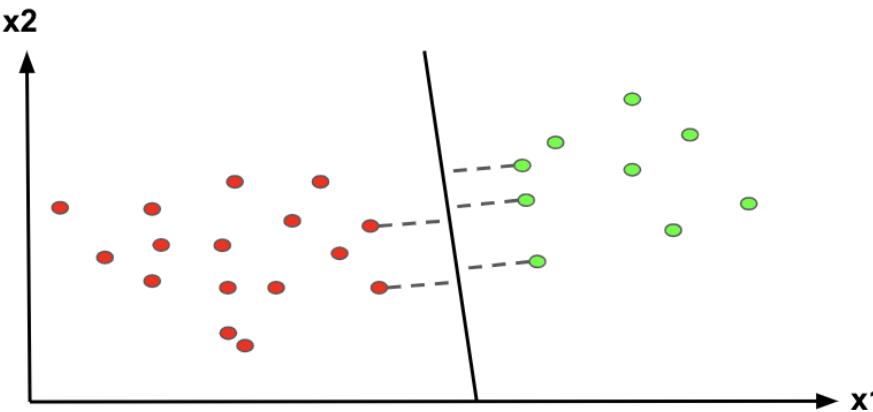


2nd Improvement

Consider more points to get the decision boundary.

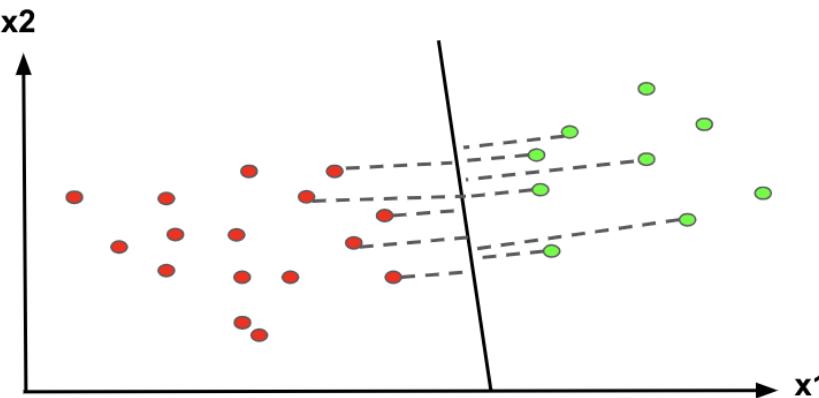
Gamma Hyper-Parameters

- It defines how far influences the calculation of plausible line of separation.
- when gamma is higher, nearby points will have high influence; low gamma means far away points also be considered to get the decision boundary.



High Gamma

- only near points are considered.



Low Gamma

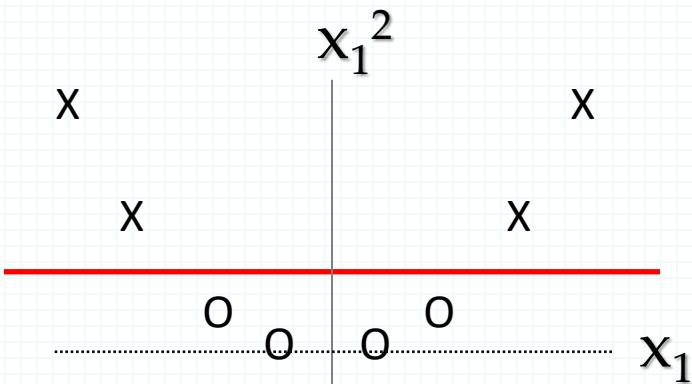
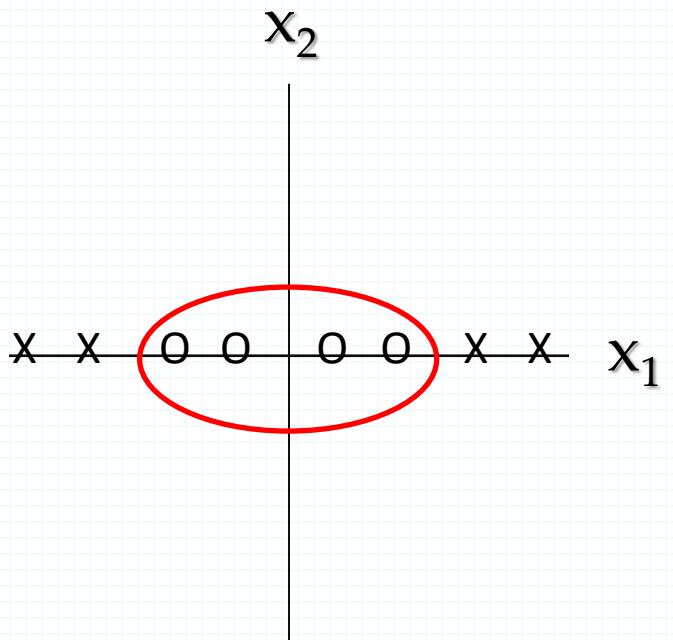
- far away points are also considered

3rd Improvement

How to make a plane curved?

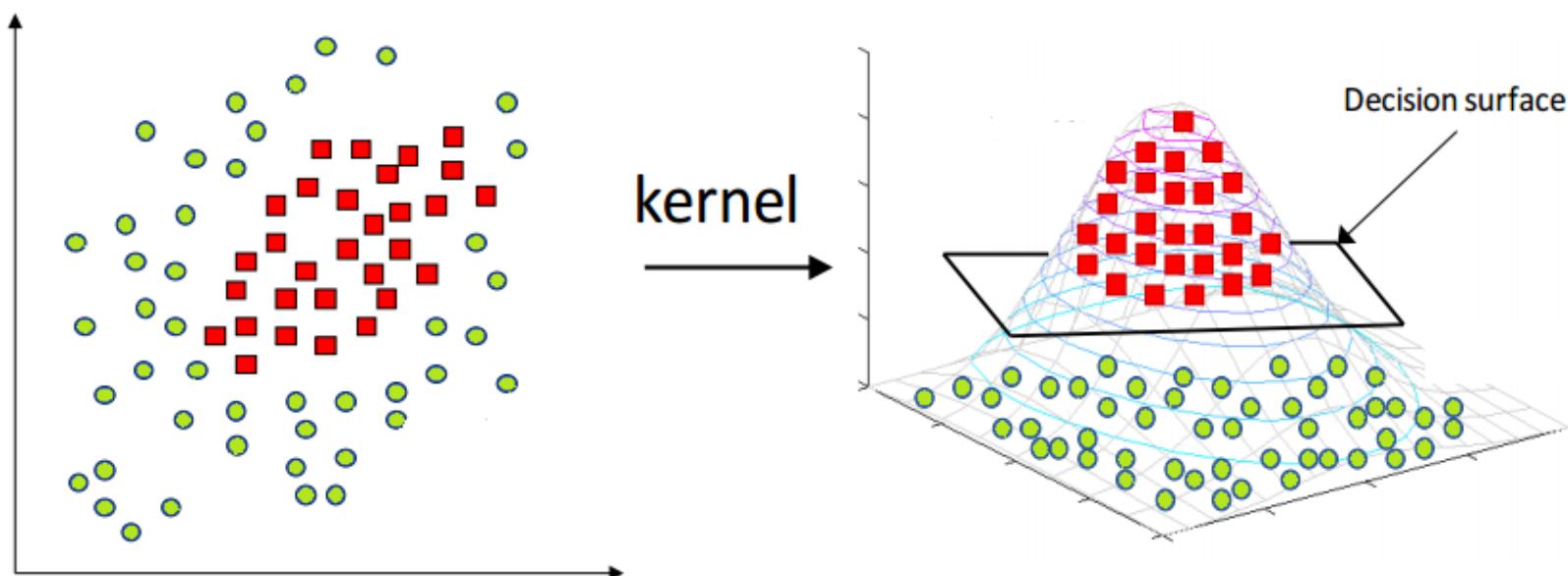
When Linear Separators Fail

- The most interesting feature of SVM is that it can even work with a non-linear dataset.
- We use “Kernel Trick” which makes it easier to classify the points.



The Kernel Trick

- try converting this lower dimension space to a higher dimension space using some quadratic functions which will allow us to find a decision boundary that clearly divides the data points.
- These functions which help us do this are called Kernels and which kernel to use is purely determined by hyperparameter tuning.



Kernel functions: Polynomial kernel

- Following is the formula for the polynomial kernel:

$$f(X_1, X_2) = (X_1^T \cdot X_2 + 1)^d$$

- Here d is the degree of the polynomial, which we need to specify manually.
- Suppose we have two features X_1 and X_2 and output variable as Y, so using polynomial kernel we can write it as:

$$\begin{aligned} X_1^T \cdot X_2 &= \begin{bmatrix} X_1 \\ X_2 \end{bmatrix} \cdot [X_1 \quad X_2] \\ &= \begin{bmatrix} X_1^2 & X_1 \cdot X_2 \\ X_1 \cdot X_2 & X_2^2 \end{bmatrix} \end{aligned}$$

- So we basically need to find X_1^2 , X_2^2 and $X_1 \cdot X_2$, and now we can see that 2 dimensions got converted into 5 dimensions.

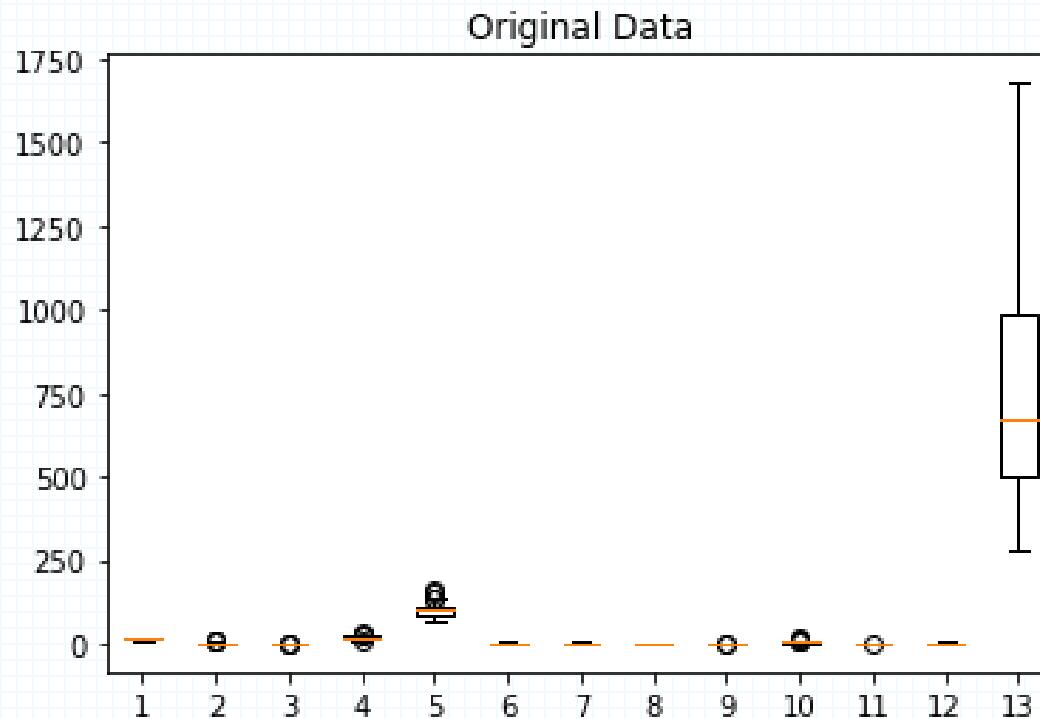
Other commonly used kernels

- linear: $\langle x, x' \rangle$.
- polynomial: $(\gamma \langle x, x' \rangle + r)^d$, where d is specified by parameter `degree`, r by `coef0`.
- rbf: $\exp(-\gamma \|x - x'\|^2)$, where γ is specified by parameter `gamma`, must be greater than 0.
- sigmoid $\tanh(\gamma \langle x, x' \rangle + r)$, where r is specified by `coef0`.

4th Improvement Feature Scaling

Why feature scaling?

- Why?
- Any suggestion?



Feature scaling

- **Feature scaling** is mapping the feature values of a dataset into the same range.
- The two most widely adopted approaches for feature scaling are normalization and standardization.
- Normalization maps the values into the $[0, 1]$ interval:

$$z = \frac{x - \min(x)}{\max(x) - \min(x)}$$

- Standardization shifts the feature values to have a mean of zero, then maps them into a range such that they have a standard deviation of 1:

$$z = \frac{x - \mu}{\sigma}$$

- It centers the data, and it's more flexible to new values that are not yet seen in the dataset. That's why we prefer standardization in general.

Feature scaling

