

# **UVA CS 4501: Machine Learning**

## **Lecture 17: Support Vector Machine (nonlinear) Kernel Trick and in Practice**

Dr. Yanjun Qi

University of Virginia

Department of  
Computer Science

# Where are we ? ➔

## Five major sections of this course

- ☐ ~~Regression (supervised)~~
- ☐ Classification (supervised)
- ☐ Unsupervised models
- ☐ Learning theory
- ☐ Graphical models

# Today

- ❑ Support Vector Machine (SVM)
  - ✓ History of SVM
  - ✓ Large Margin Linear Classifier
  - ✓ Define Margin ( $M$ ) in terms of model parameter
  - ✓ Optimization to learn model parameters ( $w, b$ )
  - ✓ Non linearly separable case
  - ✓ Optimization with dual form
  - ➔ ✓ Nonlinear decision boundary
  - ✓ Practical Guide

# Dual SVM for linearly separable case – Training / Testing

Our dual target function:  $\max_{\alpha} \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j \mathbf{x}_i^T \mathbf{x}_j$

$$\sum_i \alpha_i y_i = 0$$


$$\alpha_i \geq 0 \quad \forall i$$

Dot product for all training samples

Dot product with (“all” ??) training samples

To evaluate a new sample  $\mathbf{x}_{ts}$   
we need to compute:

$$\mathbf{w}^T \mathbf{x}_{ts} + b = \sum_i \alpha_i y_i \mathbf{x}_i^T \mathbf{x}_{ts} + b$$



$$\hat{y}_{ts} = \text{sign} \left( \sum_{i \in \text{SupportVectors}} \alpha_i y_i \left( \mathbf{x}_i^T \mathbf{x}_{ts} \right) + b \right)$$

$$\max_{\alpha} \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j \mathbf{x}_i^T \mathbf{x}_j$$

$$\sum_i \alpha_i y_i = 0$$

$$C > \alpha_i \geq 0, \forall i$$



$$\max_{\alpha} \sum_i \alpha_i - \sum_{i,j} \alpha_i \alpha_j y_i y_j \Phi(\mathbf{x}_i)^T \Phi(\mathbf{x}_j)$$

$$\sum_i \alpha_i y_i = 0$$

$$C > \alpha_i \geq 0, \forall i$$

# Training

$$\mathbf{w}^T \mathbf{x}_{ts} + b = \sum_i \alpha_i y_i \mathbf{x}_i^T \mathbf{x}_{ts} + b$$

$$\hat{y}_{ts} = \text{sign} \left( \sum_{i \in \text{SupportVectors}} \alpha_i y_i (\mathbf{x}_i^T \mathbf{x}_{ts}) + b \right)$$

Handwritten diagram illustrating the matrix multiplication  $\mathbf{X}^T \mathbf{x}_{ts}$ :

A vertical rectangle represents a matrix of size  $n \times 1$ . The rows are indexed from 1 to  $n$ . The label  $\mathbf{x}_{ts}$  is written above the rectangle. Inside the rectangle, the expression  $\mathbf{x}_i^T \mathbf{x}_{ts}$  is written, indicating the dot product of each support vector  $\mathbf{x}_i$  with the test vector  $\mathbf{x}_{ts}$ .

Handwritten equation showing the summation over support vectors:

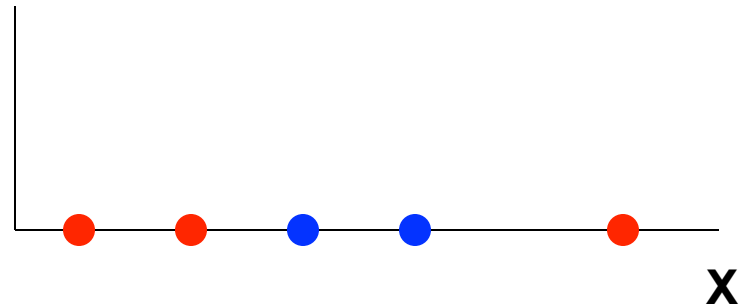
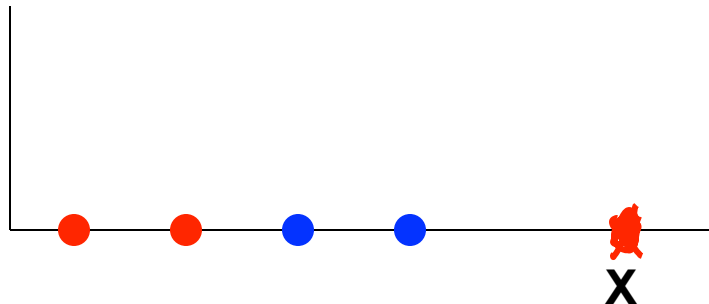
$$\Rightarrow \sum_{SV} \alpha_i y_i \underbrace{\Phi(\mathbf{x}_i) \Phi(\mathbf{x}_{ts})}_{+b}$$

Testing

# Classifying in 1-d

Can an SVM correctly classify this data?

What about this?

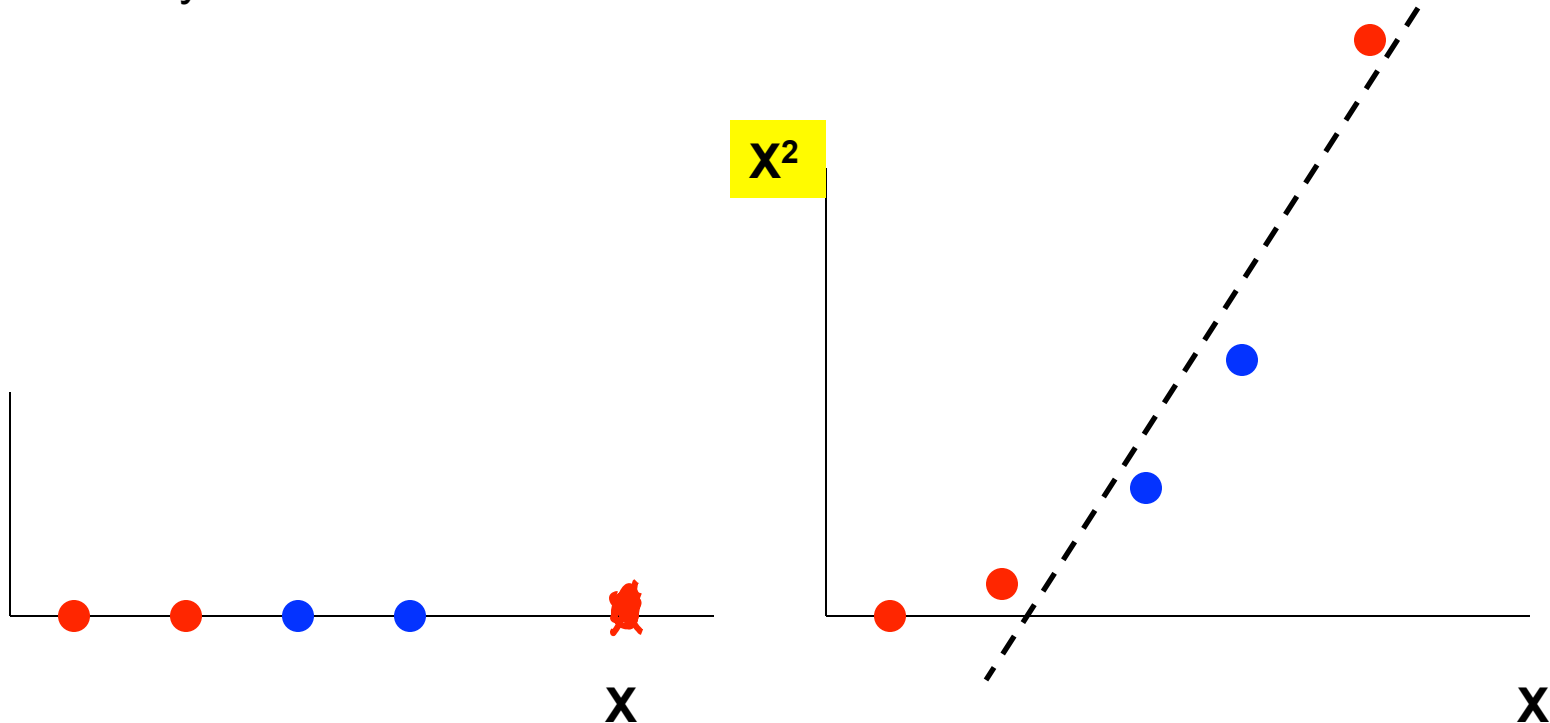


# Classifying in 1-d

$\int \rightarrow$  separable  
 $\lfloor \rightarrow$  nonlinear

Can an SVM correctly classify this data?

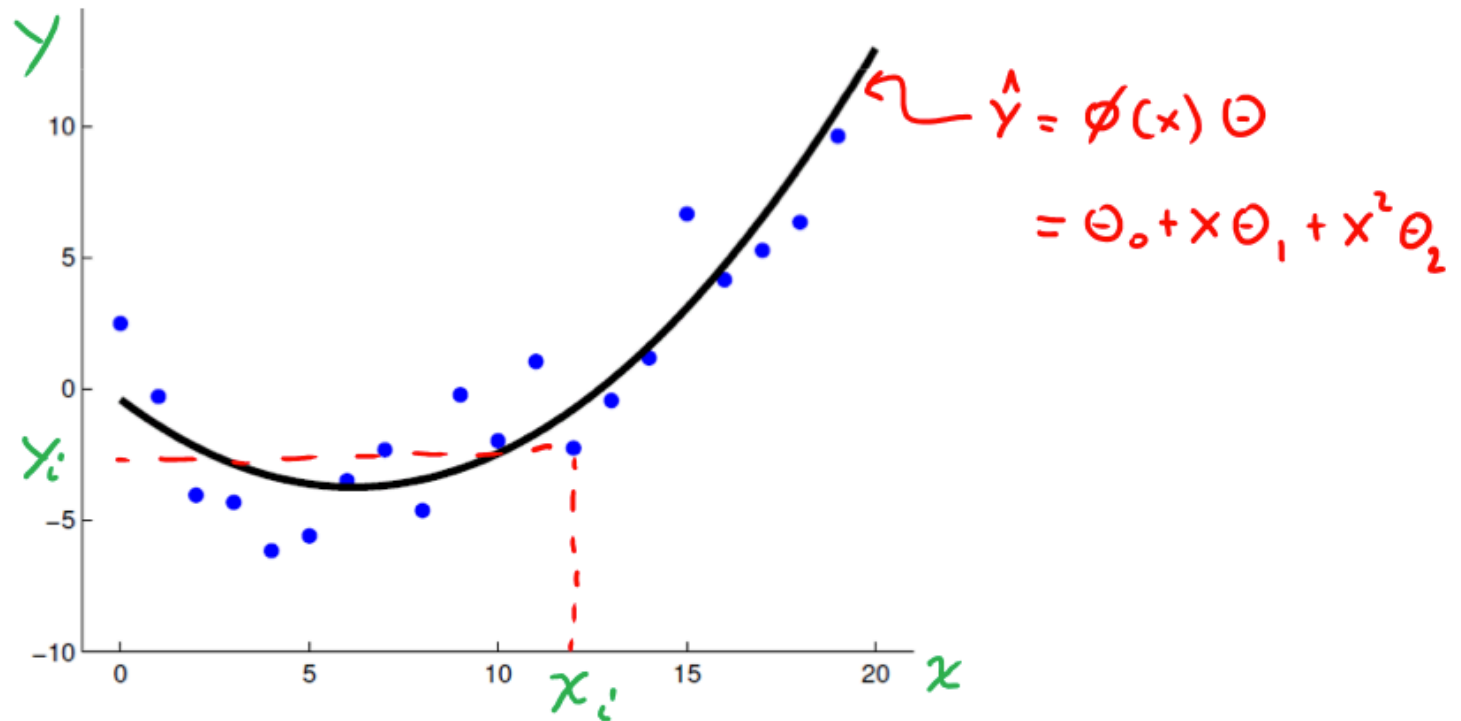
And now? (extend with polynomial basis )





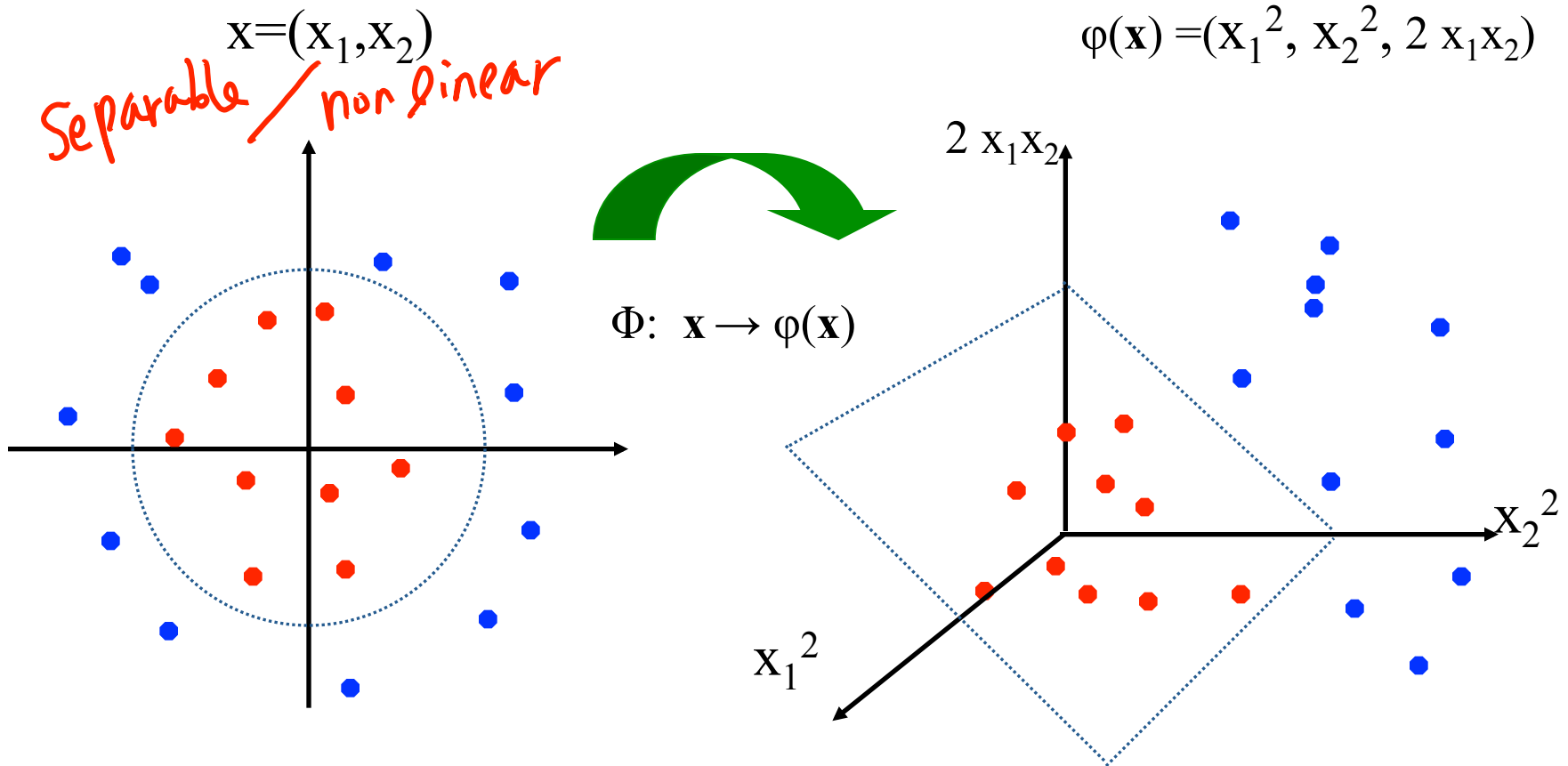
# RECAP: Polynomial regression

For example,  $\phi(x) = [1, x, x^2]$



# Non-linear SVMs: 2D

- The original input space ( $\mathbf{x}$ ) can be mapped to some higher-dimensional feature space ( $\phi(\mathbf{x})$ ) where the training set is separable:

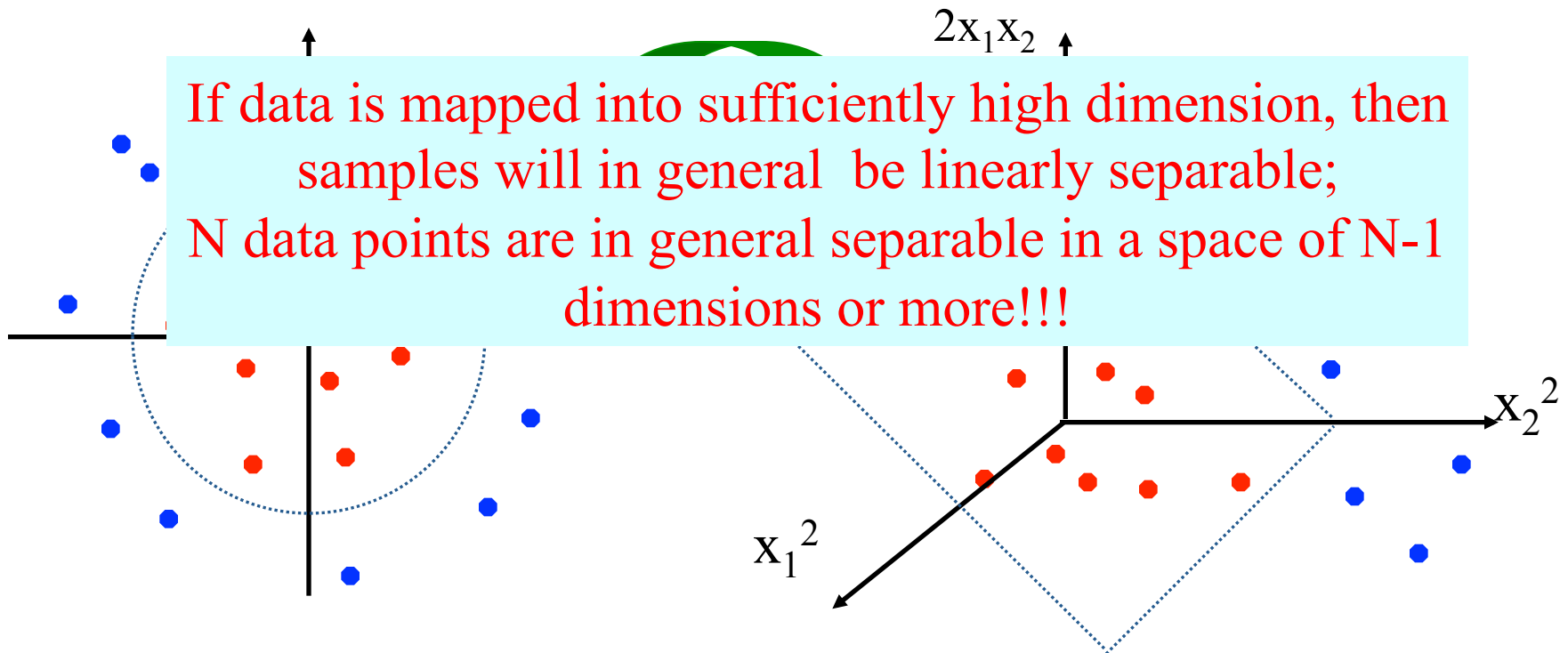


# Non-linear SVMs: 2D

- The original input space ( $\mathbf{x}$ ) can be mapped to some higher-dimensional feature space ( $\phi(\mathbf{x})$ ) where the training set is separable:

$$\mathbf{x} = (x_1, x_2)$$

$$\phi(\mathbf{x}) = (x_1^2, x_2^2, 2x_1x_2)$$

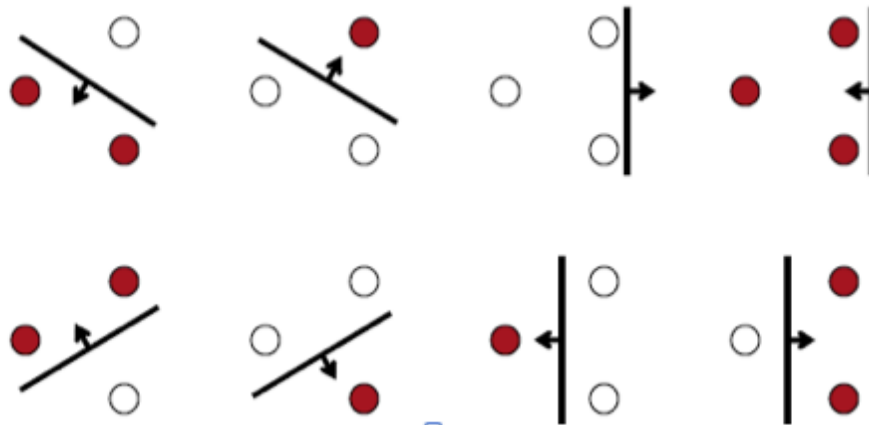


# A little bit theory: Vapnik-Chervonenkis (VC) dimension

If data is mapped into sufficiently high dimension, then samples will in general be linearly separable;

N data points are in general separable in a space of N-1 dimensions or more!!!

- **VC dimension of the set of oriented lines in  $R^2$  is 3**
  - It can be shown that the VC dimension of the family of oriented separating hyperplanes in  $R^N$  is at least N+1



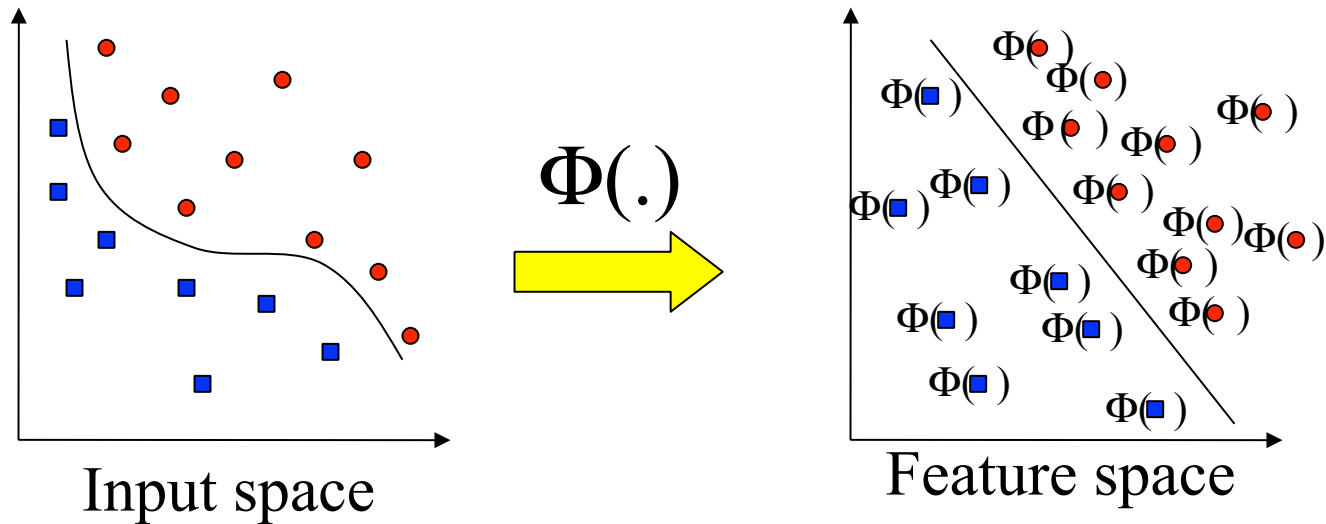
If data is mapped into sufficiently high dimension, then samples will in general be linearly separable;

N data points are in general separable in a space of N-1 dimensions or more!!!

$$X \rightarrow \Phi(X)$$

- Possible problems

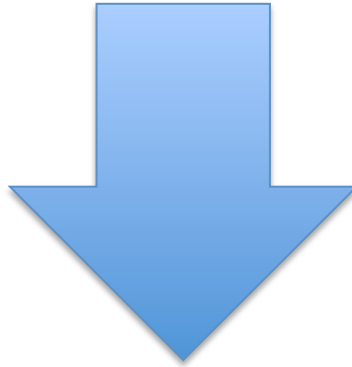
- High computation burden due to high-dimensionality
- Many more parameters to estimate



SVM solves these two issues simultaneously

- “Kernel tricks” for efficient computation
- Dual formulation only assigns parameters to samples, not to features

- SVM solves these two issues simultaneously
  - “Kernel tricks” for efficient computation
  - Dual formulation only assigns parameters to samples, not features



## (1). “Kernel tricks” for efficient computation

~~\$(X)~~

Never represent features explicitly

- ☐ Compute dot products in closed form

Very interesting theory – Reproducing Kernel Hilbert Spaces

- ☐ Not covered in detail here

$K(\mathbf{x}_i, \mathbf{x}_j) \equiv \phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$  is called the kernel function.

- Linear kernel (we've seen it)  $K(\mathbf{x}, \mathbf{z}) = \mathbf{x}^T \mathbf{z}$

$$\begin{cases} \mathbf{x} \in \mathbb{R}^p \\ \mathbf{z} \in \mathbb{R}^p \end{cases}$$

- Polynomial kernel (we will see an example)

$$K(\mathbf{x}, \mathbf{z}) = (1 + \mathbf{x}^T \mathbf{z})^d = \underbrace{\Phi_p(\mathbf{x})^T}_{p \rightarrow d(p)} \underbrace{\Phi_p(\mathbf{z})}_{p \rightarrow d(p)}$$

where  $d = 2, 3, \dots$  To get the feature vectors we concatenate all  $d$ th order polynomial terms of the components of  $\mathbf{x}$  (weighted appropriately)

- Radial basis kernel

$$K(\mathbf{x}, \mathbf{z}) = \exp\left(-r \|\mathbf{x} - \mathbf{z}\|^2\right) = \underbrace{\Phi_r(\mathbf{x})^T}_{p = \infty} \underbrace{\Phi_r(\mathbf{z})}_{p = \infty}$$

In this case.,  $r$  is hyperpara. The feature space of the RBF kernel has an infinite number of dimensions

Never represent features explicitly

☐ Compute dot products with a closed form

Very interesting theory – Reproducing Kernel Hilbert Spaces

☐ Not covered in detail here



# Example: Quadratic kernels

$$K(\mathbf{x}, \mathbf{z}) = (1 + \mathbf{x}^T \mathbf{z})^d \quad \Rightarrow \quad (1 + \mathbf{x}^T \mathbf{z})^2$$

$$K(\mathbf{x}, \mathbf{z}) := \Phi(\mathbf{x})^T \Phi(\mathbf{z})$$

- Consider all quadratic terms for  $x_1, x_2 \dots x_p$

$$\max_{\alpha} \sum_i \alpha_i - \sum_{i,j} \alpha_i \alpha_j y_i y_j \Phi(\mathbf{x}_i)^T \Phi(\mathbf{x}_j)$$

$$\sum_i \alpha_i y_i = 0$$

$$\alpha_i \geq 0 \quad \forall i$$

$$\Phi(\mathbf{x}) =$$

$$\begin{bmatrix} 1 \\ \sqrt{2}x_1 \\ \vdots \\ \sqrt{2}x_p \\ x_1^2 \\ \vdots \\ x_p^2 \\ \sqrt{2}x_1x_2 \\ \vdots \\ \sqrt{2}x_{p-1}x_p \end{bmatrix}$$

$$K(\mathbf{x}, \mathbf{z}) = (1 + \mathbf{x}^T \mathbf{z})^2, \quad [d=2], \quad [p=2] \quad \begin{cases} \mathbf{x} = (x_1, x_2) \\ \mathbf{z} = (z_1, z_2) \end{cases}$$


$$k(x, z) = (1 + x_1 z_1 + x_2 z_2)^2 \Rightarrow \mathcal{O}(p)$$

$$\mathcal{O}(p^2) \left( \begin{array}{l} = (1, \sqrt{2}x_1, \sqrt{2}x_2, x_1^2, x_2^2, \sqrt{2}x_1x_2)^T \\ (1, \sqrt{2}z_1, \sqrt{2}z_2, z_1^2, z_2^2, \sqrt{2}z_1z_2) \end{array} \right)$$


$$= \Phi(\mathbf{x})^T \Phi(\mathbf{z})$$

# The kernel trick

$p^d n^2$  operations if using the basis function representations in building a poly-kernel matrix



So, if we define the **kernel function** as follows, there is no need to carry out basis function explicitly

$$K(\mathbf{x}, \mathbf{z}) = (\mathbf{x}^T \mathbf{z} + 1)^{d=2}$$


$p n^2$  operations in building a poly-kernel matrix for training

$$\max_{\alpha} \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j K(\mathbf{x}_i, \mathbf{x}_j)$$

$$\sum_i \alpha_i y_i = 0$$

$$C > \alpha_i \geq 0, \forall i \in \text{train}$$

# Summary:

## Modification Due to Kernel Trick

- Change all inner products to kernel functions
- For training,

Original  
Linear

$$\max_{\alpha} \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j \mathbf{x}_i^T \mathbf{x}_j$$

$$\sum_i \alpha_i y_i = 0$$

$$C > \alpha_i \geq 0, \forall i \in \text{train}$$

With kernel  
function -  
nonlinear

$$\max_{\alpha} \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j K(\mathbf{x}_i, \mathbf{x}_j)$$

$$\sum_i \alpha_i y_i = 0$$

$$C > \alpha_i \geq 0, \forall i \in \text{train}$$

# Summary:

## Modification Due to Kernel Function

- For testing, the new data  $\mathbf{x}_{ts}$

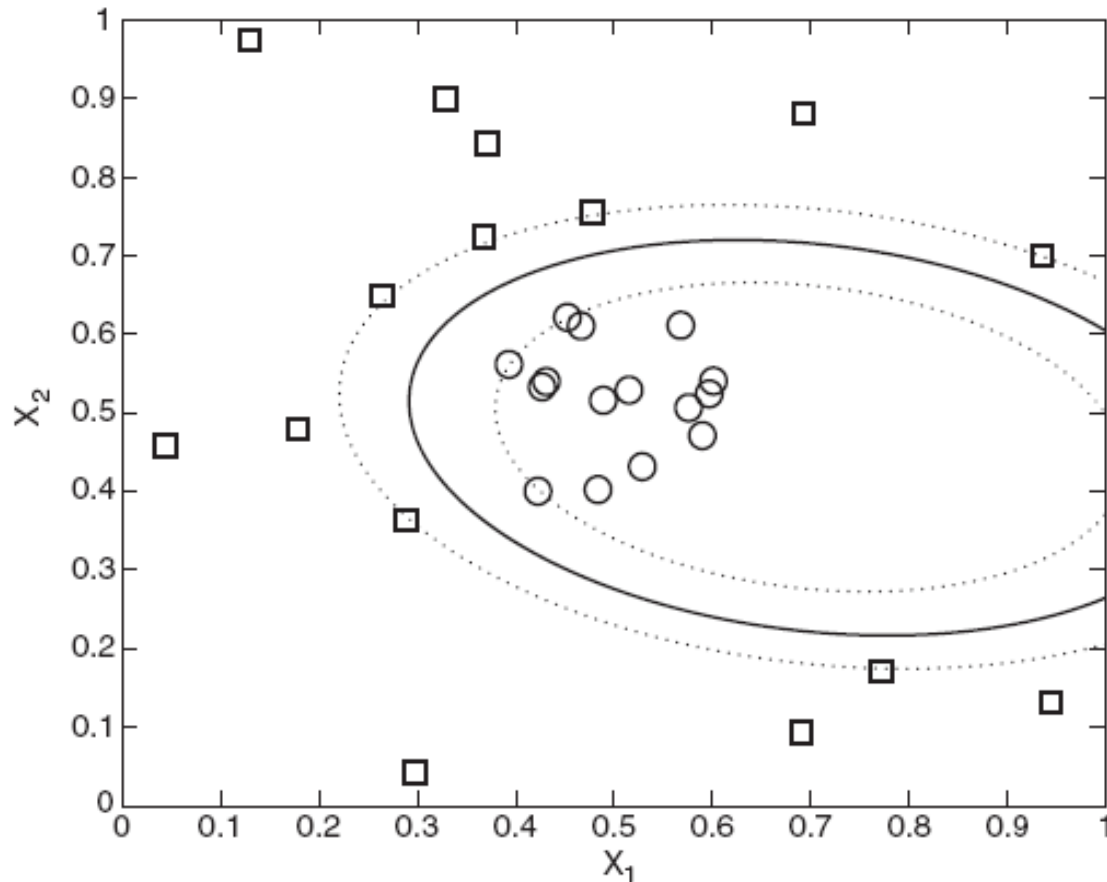
Original  
Linear

$$\hat{y}_{ts} = \text{sign} \left( \sum_{i \in \text{train}} \alpha_i y_i \mathbf{x}_i^T \mathbf{x}_{ts} + b \right)$$

With kernel  
function -  
nonlinear

$$\hat{y}_{ts} = \text{sign} \left( \sum_{i \in \text{supportVectors}} \alpha_i y_i K(\mathbf{x}_i, \mathbf{x}_{ts}) + b \right)$$

# An example: Support vector machines with polynomial kernel



**Figure 5.29.** Decision boundary produced by a nonlinear SVM with polynomial kernel.

# Kernel Trick: Implicit Basis Representation

- For some kernels (e.g. RBF ) the implicit transform basis form  $\phi(\mathbf{x})$  is infinite-dimensional!
  - But calculations with kernel are done in original space, so computational burden and curse of dimensionality aren't a problem.

$$K(\mathbf{x}, \mathbf{z}) = \exp\left(-r \|\mathbf{x} - \mathbf{z}\|^2\right)$$

$p \cdot n^2$  operations in building a RBF-kernel matrix for training

➔ Gaussian RBF Kernel corresponds to an infinite-dimensional vector space.

YouTube video of Caltech: Abu-Mostafa explaining this in more detail

<https://www.youtube.com/watch?v=XUj5JbQihIU&t=25m53s>

# Kernel Functions (Extra)

- In practical use of SVM, only the kernel function (and not basis function) is specified
- Kernel function can be thought of as a similarity measure between the input objects
- Not all similarity measure can be used as kernel function, however Mercer's condition states that any positive semi-definite kernel  $K(x, y)$ , i.e.

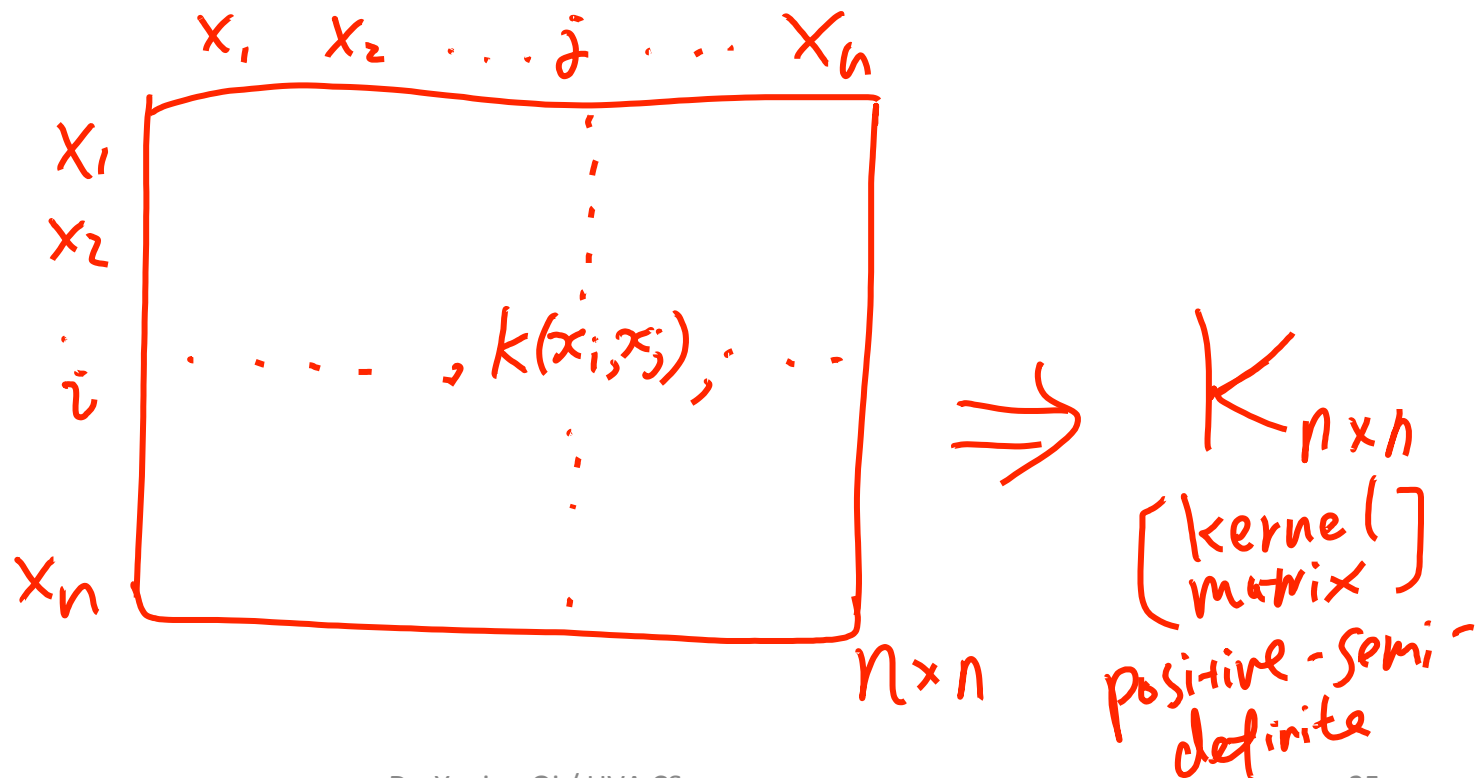
$$\sum_{i,j} K(x_i, x_j) c_i c_j \geq 0$$

can be expressed as a dot product in a high dimensional space.



# Kernel Matrix

- Kernel function creates the kernel matrix, which summarize all the data



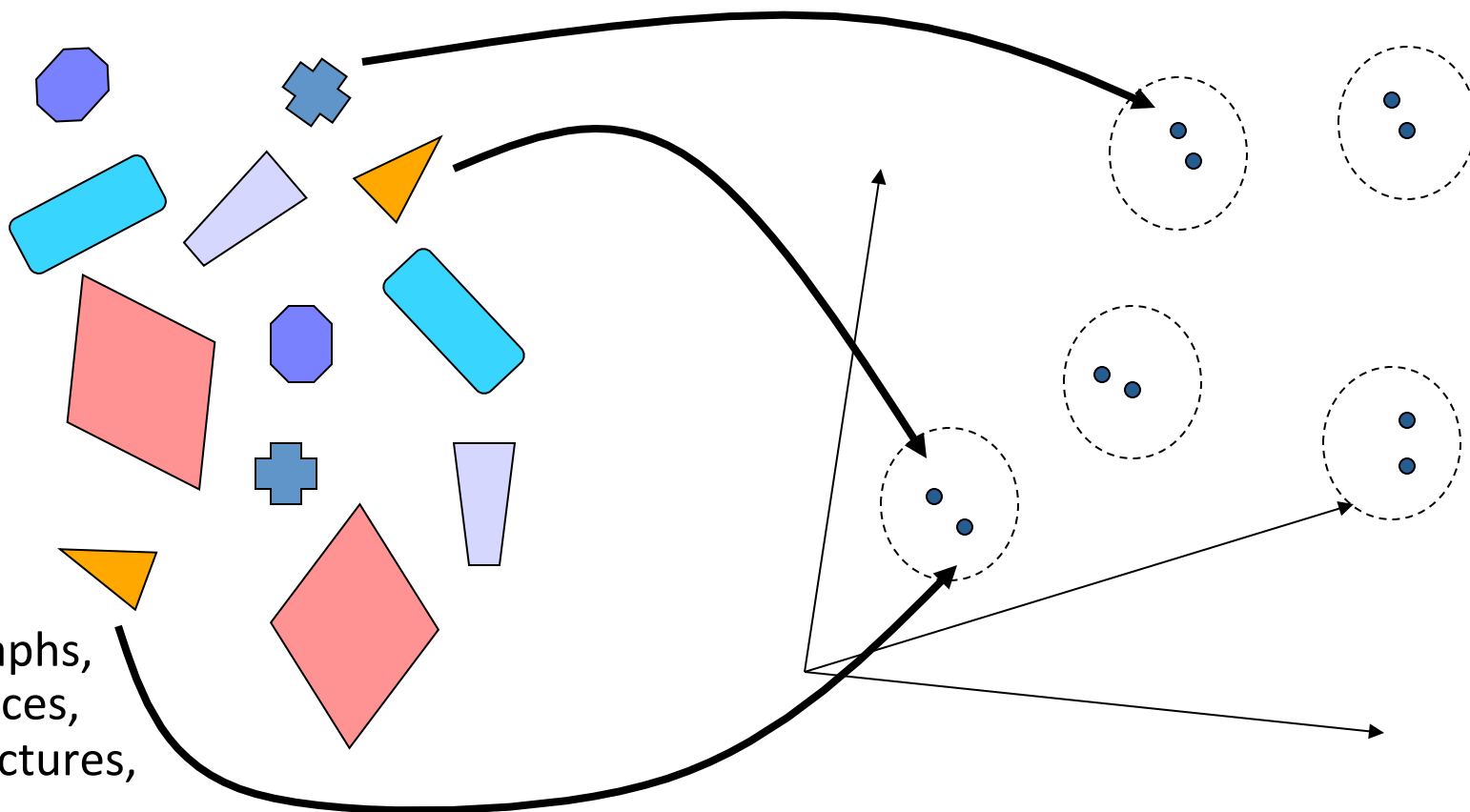
# Choosing the Kernel Function

- Probably the most tricky part of using SVM.
- The kernel function is important because it creates the kernel matrix, which summarize all the data
- Many principles have been proposed (diffusion kernel, Fisher kernel, string kernel, tree kernel, graph kernel, ...)
  - Kernel trick has helped Non-traditional data like strings and trees able to be used as input to SVM, instead of feature vectors
- In practice, a low degree polynomial kernel or RBF kernel with a reasonable width is a good initial try for most applications.

Kernel trick has helped Non-traditional data like strings and trees able to be used as input to SVM, instead of feature vectors

$$K(x, z)$$

Vector vs. Relational data



e.g. Graphs,  
Sequences,  
3D structures,

# Mercer Kernel vs. Smoothing Kernel

- The Kernels used in Support Vector Machines are different from the Kernels used in LocalWeighted /Kernel Regression.
- We can think
  - Support Vector Machines' kernels as **Mercer Kernels**
  - Local Weighted / Kernel Regression's kernels as **Smoothing Kernels**

# Why do SVMs work?

- ❑ If we are using huge features spaces (e.g., with kernels), how come we are not overfitting the data?
  - ✓ Number of parameters remains the same (and most are set to 0)
  - ✓ While we have a lot of input values, at the end we only care about the support vectors and these are usually a small group of samples
  - ✓ The minimization (or the maximizing of the margin) function acts as a sort of regularization term leading to reduced overfitting

# Why SVM Works? (Extra)

- Vapnik argues that the fundamental problem is not the number of parameters to be estimated. Rather, the problem is about the flexibility of a classifier
- Vapnik argues that the flexibility of a classifier should not be characterized by the number of parameters, but by the capacity of a classifier
  - This is formalized by the “VC-dimension” of a classifier
- The SVM objective can also be justified by structural risk minimization: the empirical risk (training error), plus a term related to the generalization ability of the classifier, is minimized
- Another view: the SVM loss function is analogous to ridge regression. The term  $\frac{1}{2} ||w||^2$  “shrinks” the parameters towards zero to avoid overfitting

# Today

- ❑ Support Vector Machine (SVM)
  - ✓ History of SVM
  - ✓ Large Margin Linear Classifier
  - ✓ Define Margin ( $M$ ) in terms of model parameter
  - ✓ Optimization to learn model parameters ( $w, b$ )
  - ✓ Non linearly separable case
  - ✓ Optimization with dual form
  - ✓ Nonlinear decision boundary
  - ✓ Practical Guide

# Software

- A list of SVM implementation can be found at
  - <http://www.kernel-machines.org/software.html>
- Some implementation (such as LIBSVM) can handle multi-class classification
- SVMLight is among one of the earliest implementation of SVM
- Several Matlab toolboxes for SVM are also available



# Summary: Steps for Using SVM in HW

- Prepare the feature-data matrix
- Select the kernel function to use
- Select the parameter of the kernel function and the value of  $C$ 
  - You can use the values suggested by the SVM software, or you can set apart a validation set to determine the values of the parameter
- Execute the training algorithm and obtain the  $\alpha_i$
- Unseen data can be classified using the  $\alpha_i$  and the support vectors

# Practical Guide to SVM

- From authors of as LIBSVM:
  - A Practical Guide to Support Vector Classification  
Chih-Wei Hsu, Chih-Chung Chang, and Chih-Jen Lin, 2003-2010
  - <http://www.csie.ntu.edu.tw/~cjlin/papers/guide/guide.pdf>

# LIBSVM

- <http://www.csie.ntu.edu.tw/~cjlin/libsvm/>
  - ✓ Developed by Chih-Jen Lin etc.
  - ✓ Tools for Support Vector classification
  - ✓ Also support multi-class classification
  - ✓ C++/Java/Python/Matlab/Perl wrappers
  - ✓ Linux/UNIX/Windows
  - ✓ SMO implementation, fast!!!

A Practical Guide to Support Vector  
Classification

# (a) Data file formats for LIBSVM

- Training.dat

+1 1:0.708333 2:1 3:1 4:-0.320755

-1 1:0.583333 2:-1 4:-0.603774 5:1

+1 1:0.166667 2:1 3:-0.333333 4:-0.433962

-1 1:0.458333 2:1 3:1 4:-0.358491 5:0.374429

...

- Testing.dat

## (b) Feature Preprocessing

- (1) Categorical Feature
  - Recommend using  $m$  numbers to represent an  $m$ -category attribute.
  - Only one of the  $m$  numbers is one, and others are zero.
  - For example, a three-category attribute such as {red, green, blue} can be represented as (0,0,1), (0,1,0), and (1,0,0)

# Feature Preprocessing

- (2) Scaling before applying SVM is very important
  - to avoid attributes in greater numeric ranges dominating those in smaller numeric ranges.
  - to avoid numerical difficulties during the calculation
  - Recommend linearly scaling each attribute to the range  $[-1, +1]$  or  $[0, 1]$ .

① Normalization  $\rightarrow \begin{cases} \text{mean} & 0 \\ \text{std} & 1 \end{cases}$

② Scaling  $\rightarrow \text{linear} \Rightarrow [ax+b]$

e.g.  $\left[ \frac{X - X_{\min}}{\max - X_{\min}} \right]$

For  $i$ -th feature  $\Rightarrow$   $\left[ \begin{array}{c} \text{Column operation} \\ \text{on } \sum_{n \times p} \end{array} \right]$

- Centering :  $X_i - \bar{X}_i \Rightarrow E(X_i) = 0$
- Scaling :  $aX_i + b \Rightarrow \text{e.g. } \frac{X_i - \min(X_i)}{\max(X_i) - \min(X_i)}$
- Normalization :  $\Rightarrow \begin{cases} E(X_i) = 0 \\ \text{Var}(X_i) = 1 \end{cases}$

Of course we have to use the same method to scale both training and testing data. For example, suppose that we scaled the first attribute of training data from  $[-10, +10]$  to  $[-1, +1]$ . If the first attribute of testing data lies in the range  $[-11, +8]$ , we must scale the testing data to  $[-1.1, +0.8]$ . See Appendix B for some real examples.

If training and testing sets are separately scaled to  $[0, 1]$ , the resulting accuracy is lower than 70%.

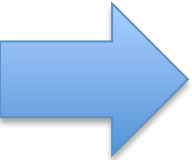
```
$ ../svm-scale -l 0 svmguide4 > svmguide4.scale
$ ../svm-scale -l 0 svmguide4.t > svmguide4.t.scale
$ python easy.py svmguide4.scale svmguide4.t.scale
Accuracy = 69.2308% (216/312) (classification)
```

Using the same scaling factors for training and testing sets, we obtain much better accuracy.

```
$ ../svm-scale -l 0 -s range4 svmguide4 > svmguide4.scale
$ ../svm-scale -r range4 svmguide4.t > svmguide4.t.scale
$ python easy.py svmguide4.scale svmguide4.t.scale
Accuracy = 89.4231% (279/312) (classification)
```

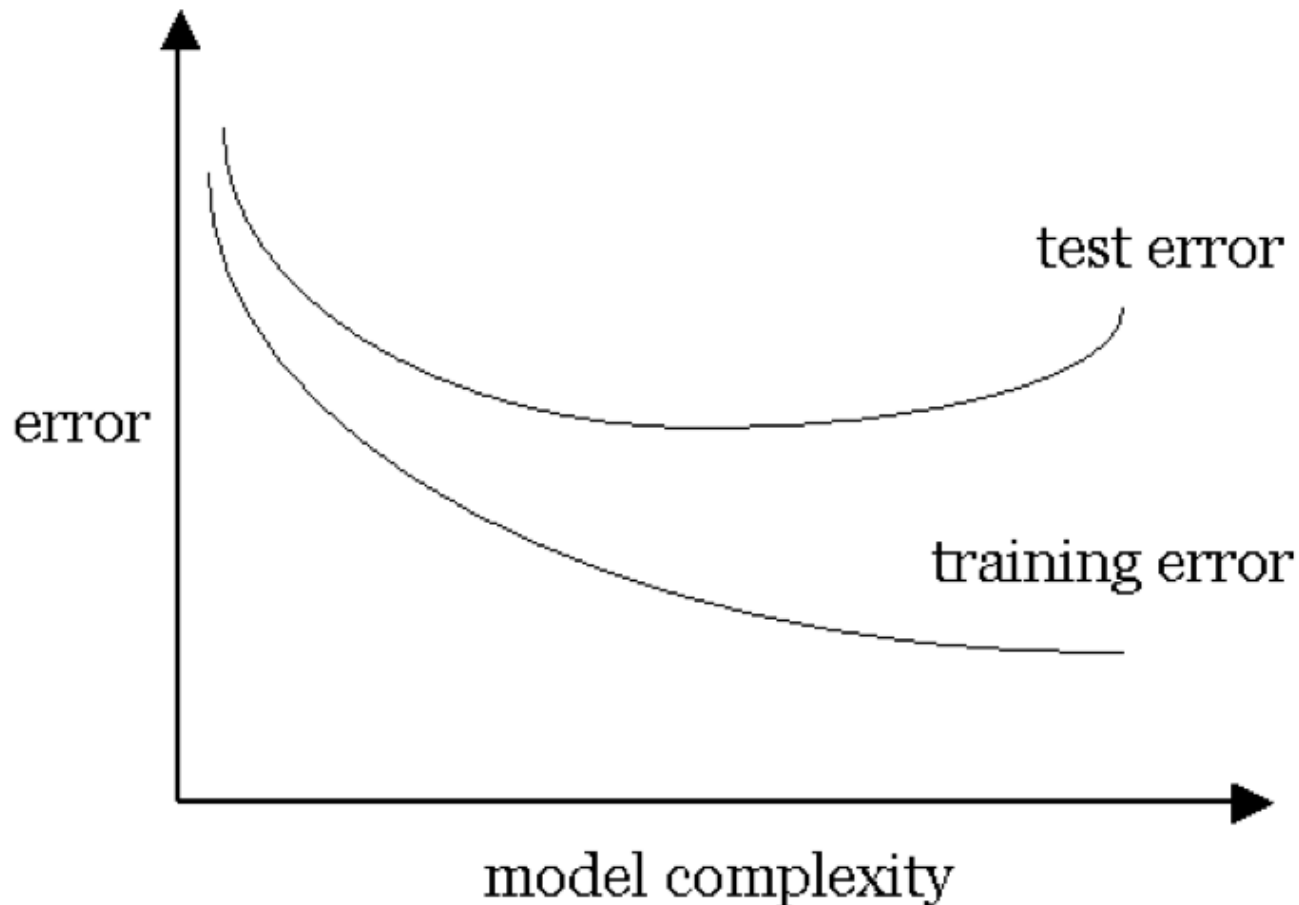


# Feature Preprocessing

- (3) missing value
  - Very very tricky !
  -  – **Easy way:** to substitute the missing values by the mean value of the variable
  - A little bit harder way: imputation using nearest neighbors
  - Even more complex: e.g. EM based (beyond the scope)

# (c) Model Selection

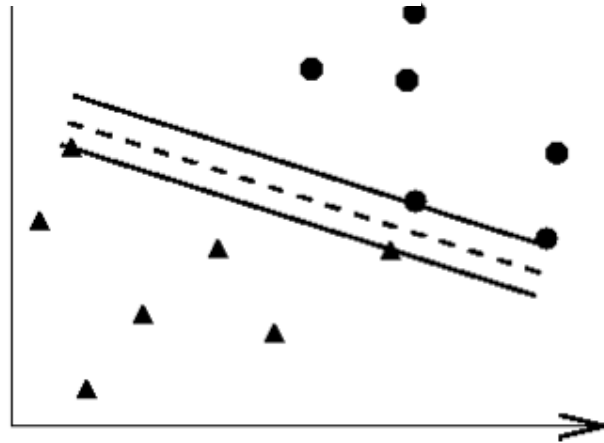
Our goal: find the model  $M$  which minimizes the test error:



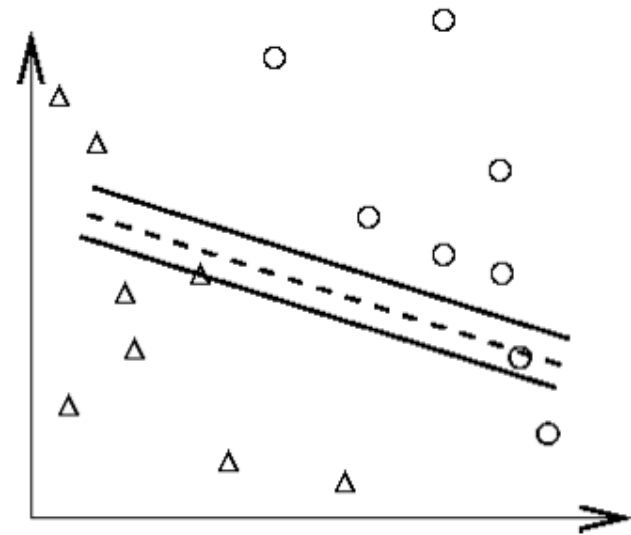
# (c) Model Selection (e.g. for linear kernel)

- linear:  $K(\mathbf{x}_i, \mathbf{x}_j) = \mathbf{x}_i^T \mathbf{x}_j$ .

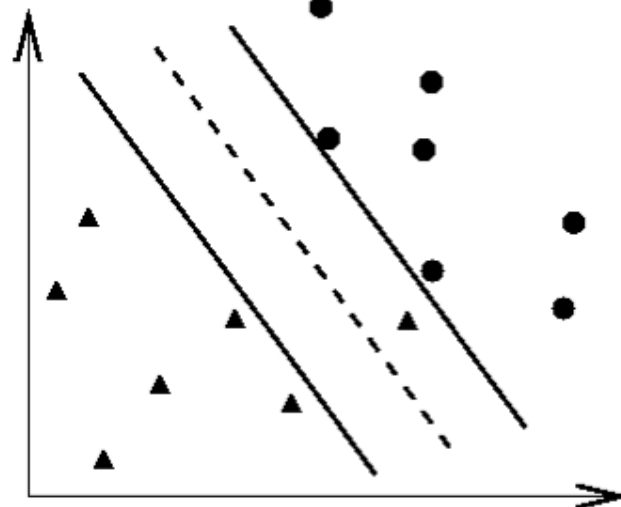
Select the  
right  
penalty  
parameter  
 $C$



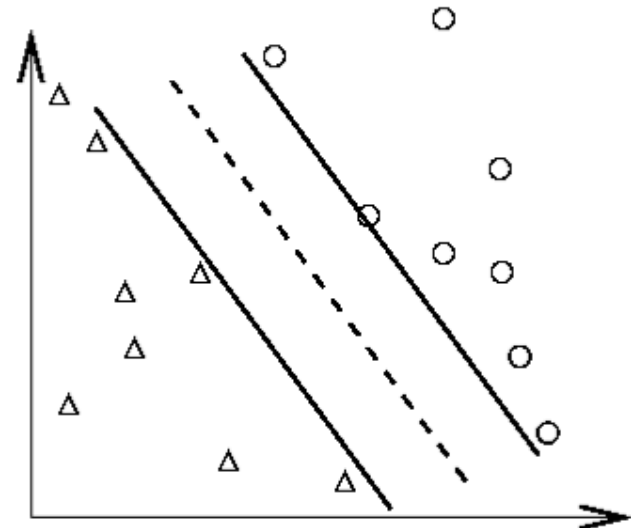
(a) Training data and an overfitting classifier



(b) Applying an overfitting classifier on testing data



(c) Training data and a better classifier



(d) Applying a better classifier on testing data

## (c) Model Selection

- radial basis function (RBF):  $K(\mathbf{x}_i, \mathbf{x}_j) = \exp(-\gamma \|\mathbf{x}_i - \mathbf{x}_j\|^2)$ ,  $\gamma > 0$ .

two parameters for an RBF kernel:  $C$  and  $\gamma$

- polynomial:  $K(\mathbf{x}_i, \mathbf{x}_j) = (\gamma \mathbf{x}_i^T \mathbf{x}_j + r)^d$ ,  $\gamma > 0$ .

Three parameters for a polynomial kernel

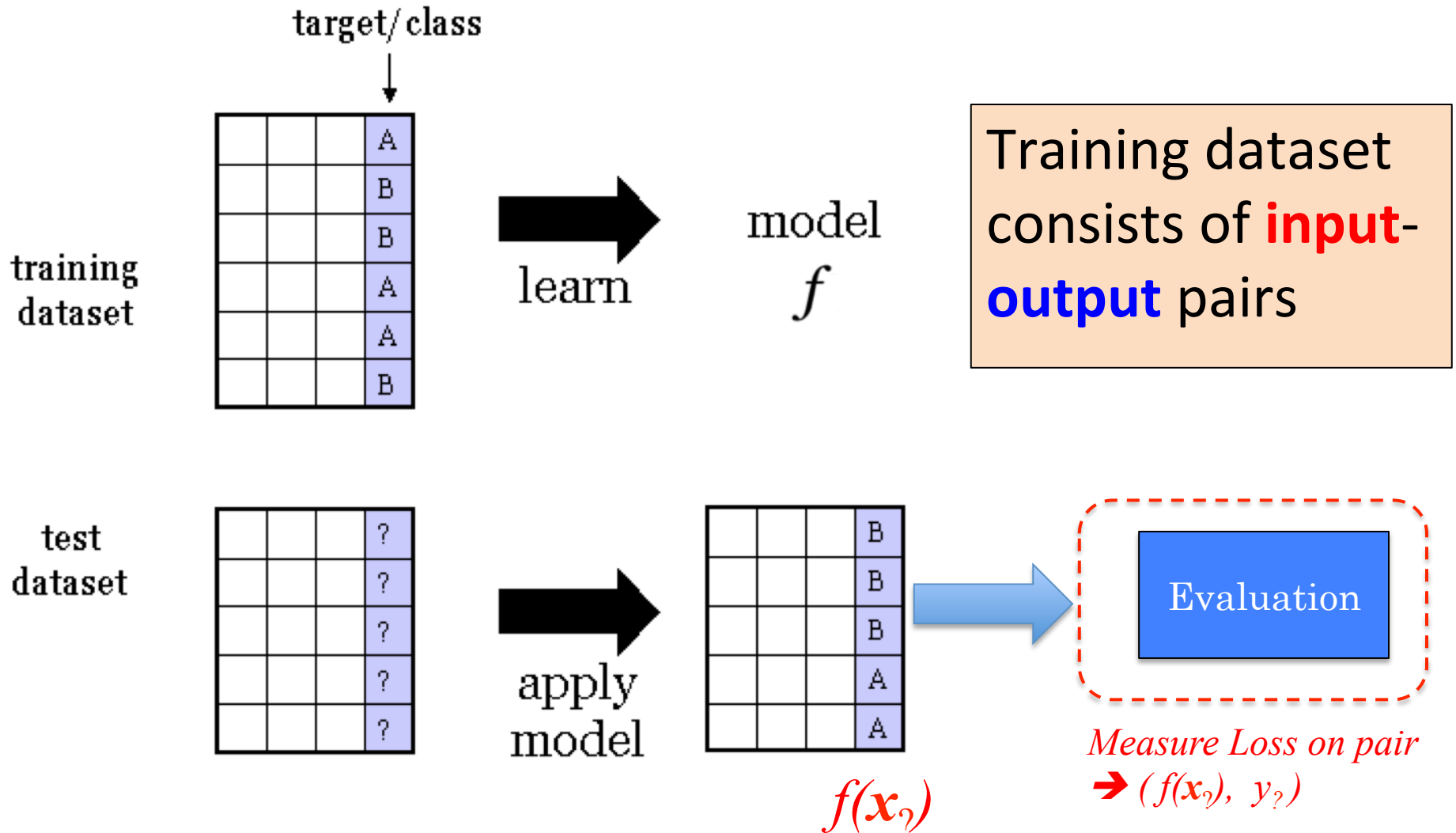
## (d) Pipeline Procedures



- (1) train / test
- (2) k-folds cross validation
- (3) k-CV on train to choose hyperparameter / then test

# Evaluation Choice-I:

## Train and Test



# Evaluation Choice-II:

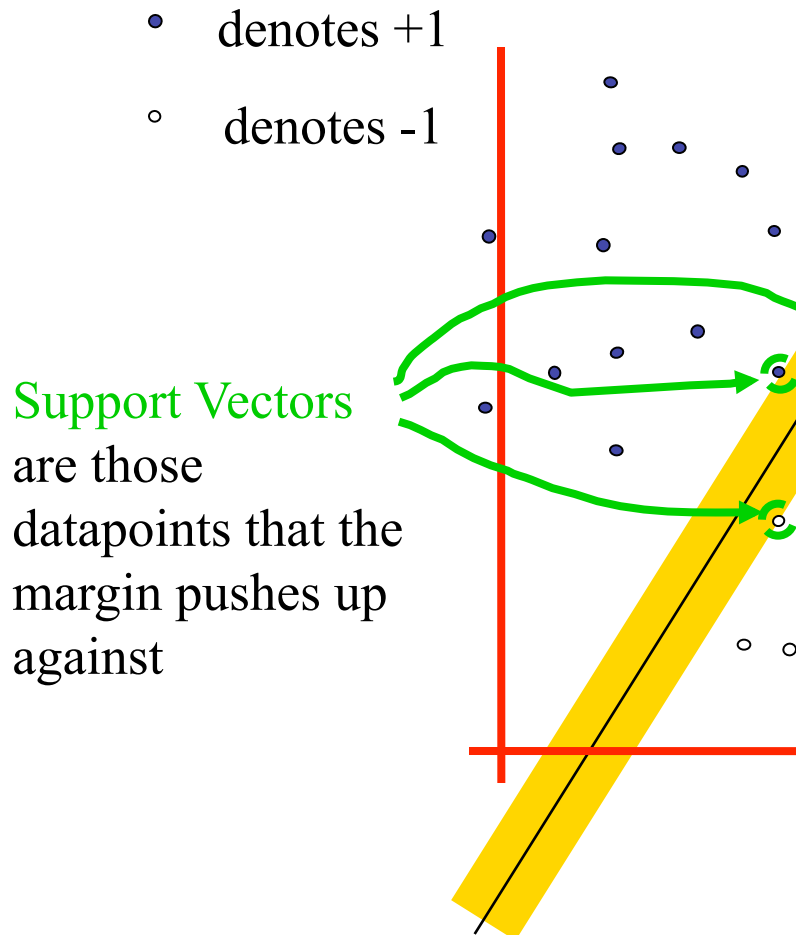
## Cross Validation

---

- Problem: don't have enough data to set aside a test set
- Solution: Each data point is used both as train and test
- Common types:
  - K-fold cross-validation (e.g.  $K=5$ ,  $K=10$ )
  - 2-fold cross-validation
  - Leave-one-out cross-validation (LOOCV)

A good practice is : to random shuffle all training sample before splitting

# Why Maximum Margin for SVM ?



1. Intuitively this feels safest.
2. If we've made a small error in the location of the boundary (it's been jolted in its perpendicular direction) this gives us least chance of causing a misclassification.
3. **LOOCV is easy since the model is immune to removal of any non-support-vector datapoints.**
4. There's some theory (using VC dimension) that is related to (but not the same as) the proposition that this is a good thing.
5. Empirically it works very very well.



# Evaluation Choice-III:

Many beginners use the following procedure now:

- Transform data to the format of an SVM package
- Randomly try a few kernels and parameters
- Test

---

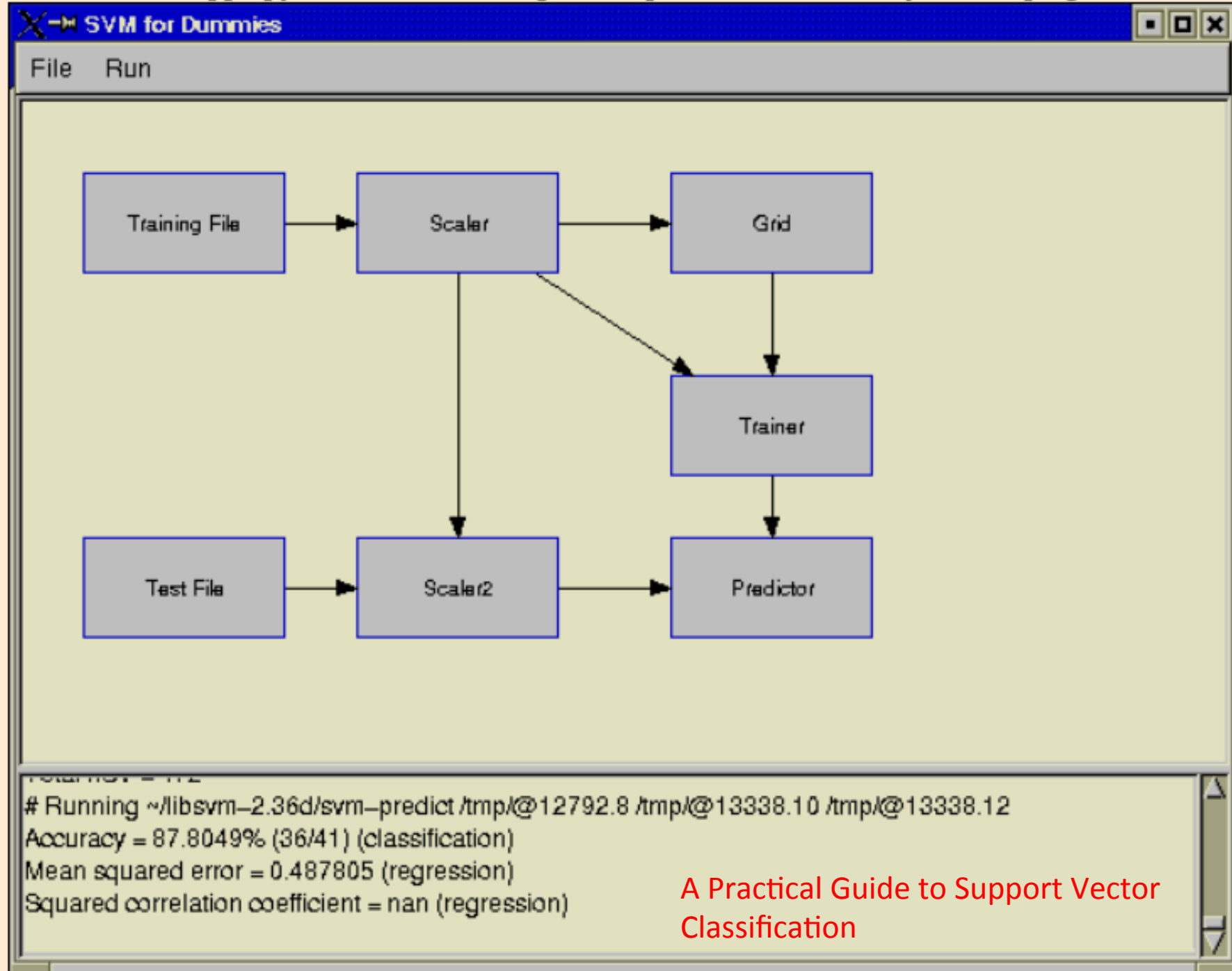
We propose that beginners try the following procedure first:

- Transform data to the format of an SVM package
- Conduct simple scaling on the data
- Consider the RBF kernel  $K(\mathbf{x}, \mathbf{y}) = e^{-\gamma \|\mathbf{x} - \mathbf{y}\|^2}$
- Use cross-validation to find the best parameter  $C$  and  $\gamma$
- Use the best parameter  $C$  and  $\gamma$  to train the whole training set<sup>5</sup>
- Test



For HW2-Q2

Run the file with `python.py`. You then have to give the path of libsvm binary files in `python/svm/svm_`

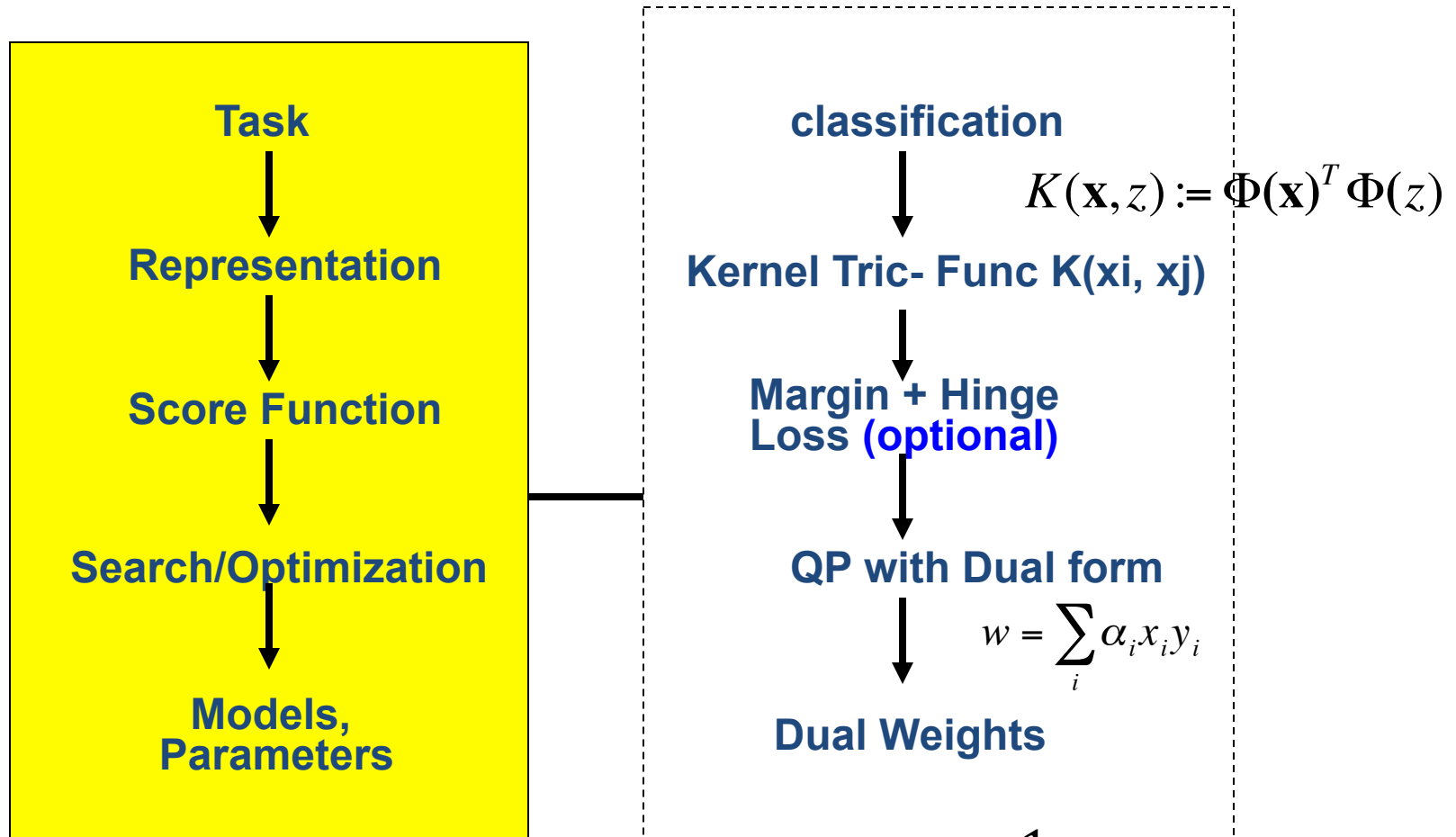


# Today: Review & Practical Guide

## □ Support Vector Machine (SVM)

- ✓ History of SVM
- ✓ Large Margin Linear Classifier
- ✓ Define Margin ( $M$ ) in terms of model parameter
- ✓ Optimization to learn model parameters ( $w, b$ )
- ✓ Non linearly separable case
- ✓ Optimization with dual form
- ✓ Nonlinear decision boundary
- ✓ Practical Guide
  - ✓ File format / LIBSVM
  - ✓ Feature preprocsssing
  - ✓ Model selection
  - ✓ Pipeline procedure

# Support Vector Machine



$$\operatorname{argmin}_{\mathbf{w}, b} \sum_{i=1}^p w_i^2 + C \sum_{i=1}^n \varepsilon_i$$

$$\text{subject to } \forall \mathbf{x}_i \in D_{\text{train}} : y_i (\mathbf{x}_i \cdot \mathbf{w} + b) \geq 1 - \varepsilon_i$$

$$\max_{\alpha} \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j \mathbf{x}_i^T \mathbf{x}_j$$

$$\sum_i \alpha_i y_i = 0, \quad \alpha_i \geq 0 \quad \forall i$$

# References

- Big thanks to Prof. Ziv Bar-Joseph and Prof. Eric Xing @ CMU for allowing me to reuse some of his slides
- Elements of Statistical Learning, by Hastie, Tibshirani and Friedman
- Prof. Andrew Moore @ CMU's slides
- Tutorial slides from Dr. Tie-Yan Liu, MSR Asia
- A Practical Guide to Support Vector Classification  
Chih-Wei Hsu, Chih-Chung Chang, and Chih-Jen Lin, 2003-2010
- Tutorial slides from Stanford "Convex Optimization I — Boyd & Vandenberghe