

# CSC2545 – Topics in Machine Learning: Kernel Methods and Support Vector Machines

- A comprehensive introduction to SVMs and other kernel methods, including theory, algorithms and applications.
- Instructor: Anthony Bonner
- Prerequisites: linear algebra, vector calculus, basic probability, mathematical maturity, programming experience.
- Expected work: 3-4 assignments.
- Lectures: TF 2-3:30pm (sometimes 2-3pm)
- For more information see [www.cs.toronto.edu/~bonner](http://www.cs.toronto.edu/~bonner)

# Course Content

- Some things kernel methods and SVMs can do:
  - non-linear classification
  - non-linear regression
  - non-linear dimensionality reduction
- How and why they work:
  - Hilbert space
  - statistical learning theory
  - regularization
- Possible additional topics:
  - convex optimization
  - implementation issues
  - Bayesian kernel methods

# Data

## Vectors

- Collections of features
  - e.g. height, weight, blood pressure, age, ...
- Can map categorical variables into vectors

## Matrices

- Images, Movies
- Remote sensing and satellite data (multispectral)

## Strings

- Documents
- Gene sequences

## Structured Objects

- XML documents
- Graphs

# Optical Character Recognition

0 9 0 2 6 0 2 5 6 0  
9 0 2 6 0 5 2 1 6 0  
5 4 6 6 4 7 3 6 1 8  
- 1 6 0 9 0 2 6 0 4  
3 5 6 0 5 6 1 8 2 0  
6 0 9 0 2 6 1 8 0 1

# Reuters Database

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Net 2,276,000 vs 1,674,000
Revs 32.6 mln vs 24.4 mln
Year
Shr 90 cts vs 69 cts
Net 4,508,000 vs 3,096,000
Revs 101.0 mln vs 76.9 mln
Avg shrs 5,029,000 vs 4,464,000
NOTE: 1986 fiscal year ended Feb 1, 1986
Reuter
&#3;</BODY></TEXT>
</REUTERS>
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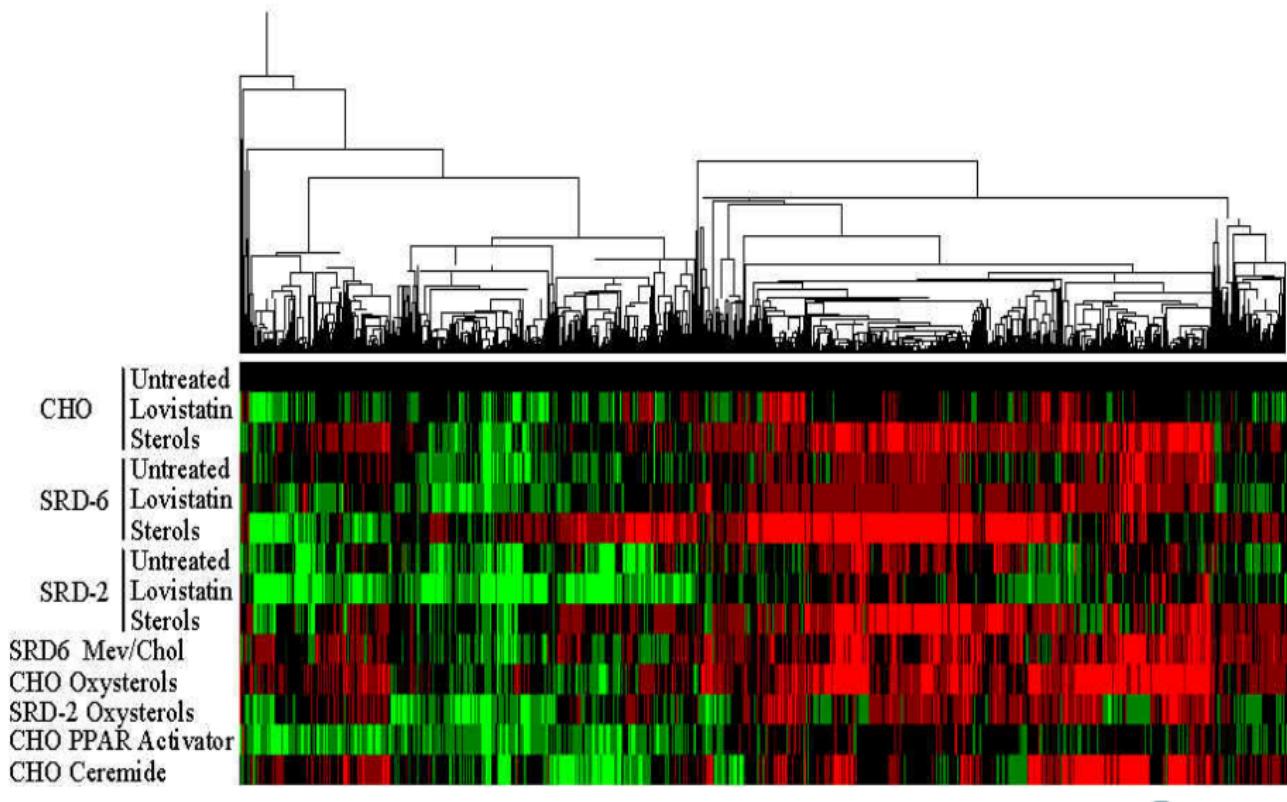
# Faces



# More Faces



# Microarray Data



# Biological Sequences

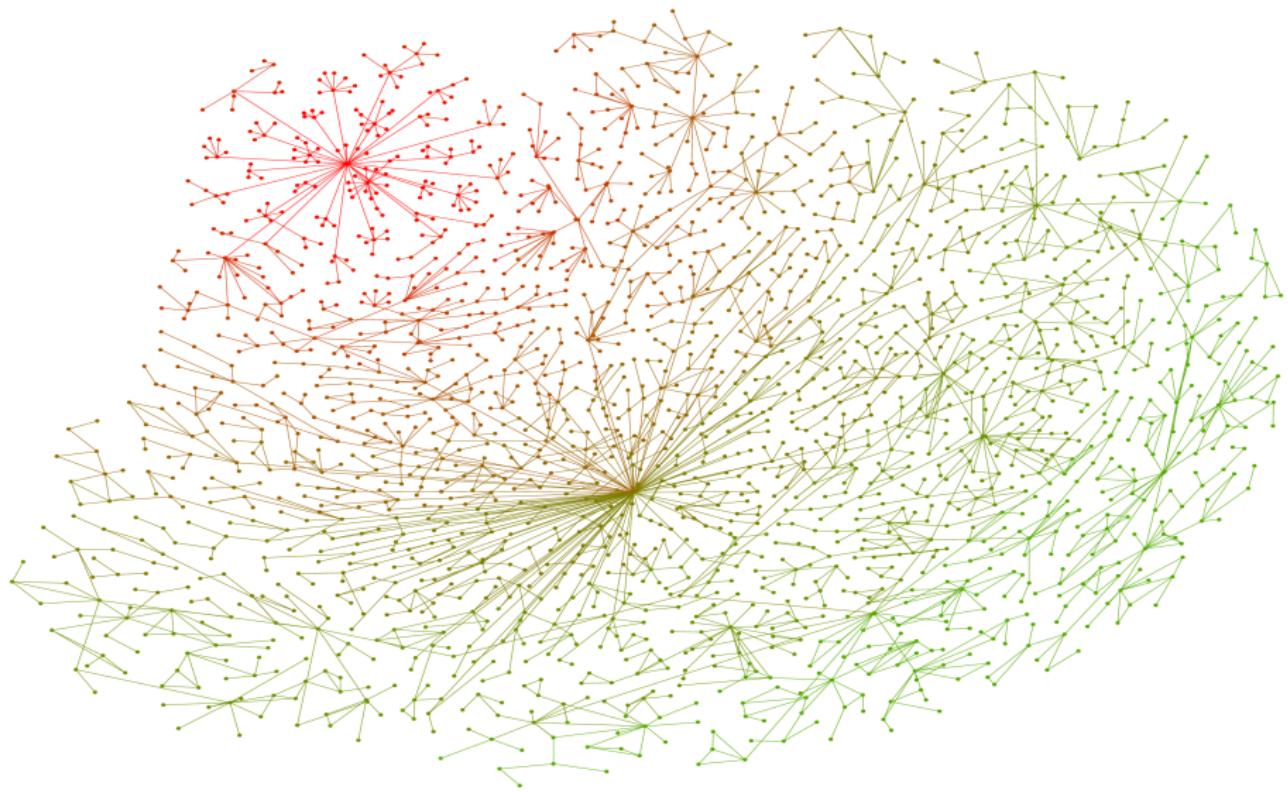
## Goal

Estimate function of protein based on sequence information.

## Example Data

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>0_d1vcaa2 2.1.1.4.1 (1-90) N-terminal domain of vascular cell adhesion molecule-1 (VCAM-1) [human (Homo sapiens)]
FKIETTPESRYLAQIGDSVSLTCSTTGCESPFFSWRTQIDSPLNGKVTNEGTTSTLMNPVSFGNEHSYL
CTATCESRKLEKGIQVEIYS
>0_d1zxq_2 2.1.1.4.2 (1-86) N-terminal domain of intracellular adhesion molecule-2, ICAM-2 [human (Homo sapiens)]
KVFEVHVRPKKLAVERPKGSLEVNCSTTCNQPEVGGLETSLNKILLDEQAQWKHYLVSNIHDTVLCQCHFT
CSGKQESMNSNSVSYQ
>0_d1tlk_ 2.1.1.4.3 Telokin [turkey (Meleagris gallopavo)]
VAEEKPHVKPYFTKTILDMDVVEGSAARFDCKVEGYPDPEVMWFKDNPVKESRHFQIDYDEEGNCSLTI
SEVCGGDDDAKYTCKAVNSLGEATCTAELLVETM
>0_d2ncm_ 2.1.1.4.4 N-terminal domain of neural cell adhesion molecule (NCAM) [human (Homo sapiens)]
RVLQVDIVPSQGEISVGESKFFLCQVAGDAKDNDISWFSPNGEKLSPNQRISVVWNDDDSSTLTIYNAN
IDDAGIYKCVVTAEDGTQSEATNVKIFQ
>0_d1tnm_ 2.1.1.4.5 Titin [Human (Homo sapiens), module M5]
RILTKPRSMTVYEGESARFSCDTDGEPVPTVTWLRKGQVLSTSARHQVTTKYKSTFEISSVQASDEGNY
SVVVENSEGKQEAFTLTIQK
>0_d1wiu_ 2.1.1.4.6 Twitchin [Nematode (Caenorhabditis elegans)]
LKPKILTASRKIKIKAGFTHNLEVDFIGAPDPTATWTVGDSGAALAPELLVDAKSSTTSIFFPSAKRADS
GNYKLKVKNELGEDEAIFEVIVQ
>0_d1koa_1 2.1.1.4.6 (351-447) Twitchin [Nematode (Caenorhabditis elegans)]
QPRFIVKPYGTEVGEGQSANFYCRVIASSPPVVTWHKDDRELKQSVKYMKRYNGNDYGLTINRVKGDDKG
EYTVRAKNSYGTKEEIVFLNVTRHSEP
```

# Graphs



# What to do with data

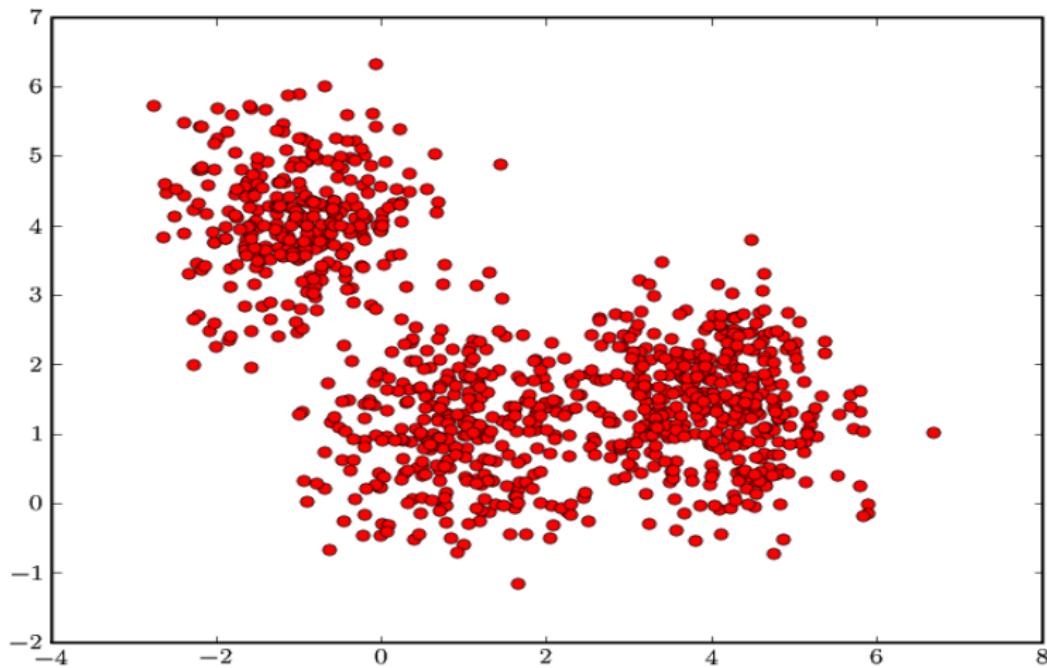
## Unsupervised Learning

- Find clusters of the data
- Find low-dimensional representation of the data  
(e.g. unroll a swiss roll, find structure)
- Find interesting directions in data
- Interesting coordinates and correlations
- Find novel observations / database cleaning

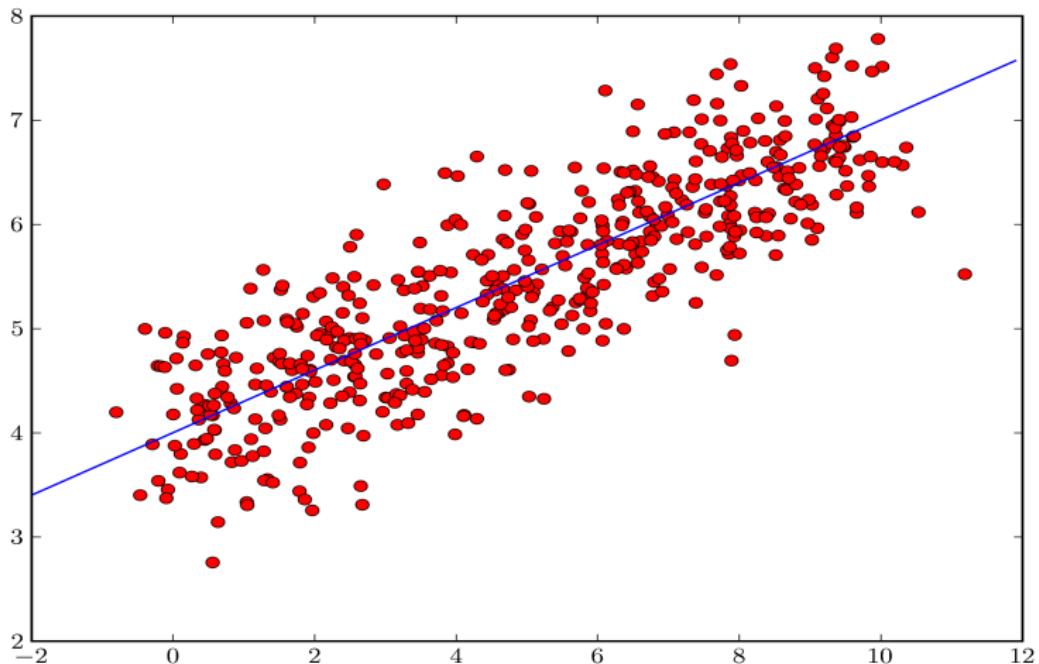
## Supervised Learning

- Classification (distinguish apples from oranges)
- Speech recognition
- Regression (tomorrow's stock value)
- Predict time series
- Annotate strings

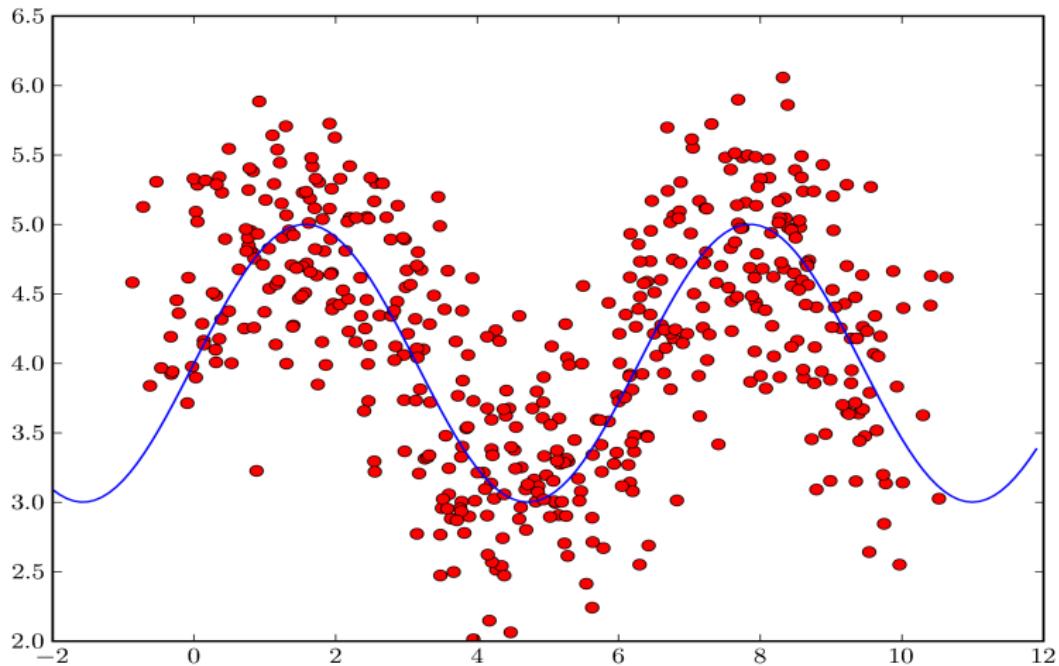
# Clustering



# Principal Components



# Linear Subspace



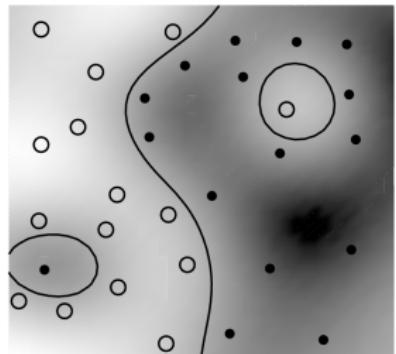
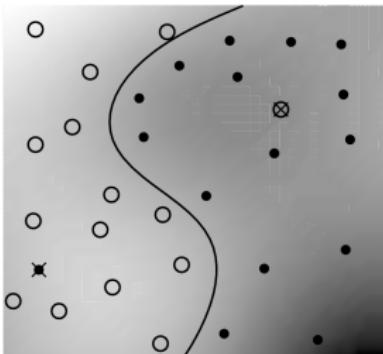
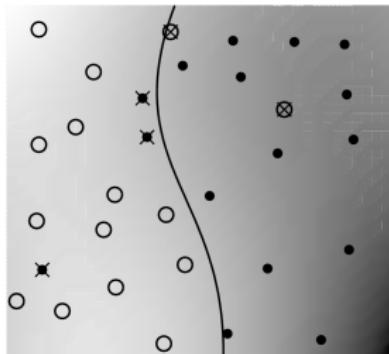
# Classification

## Data

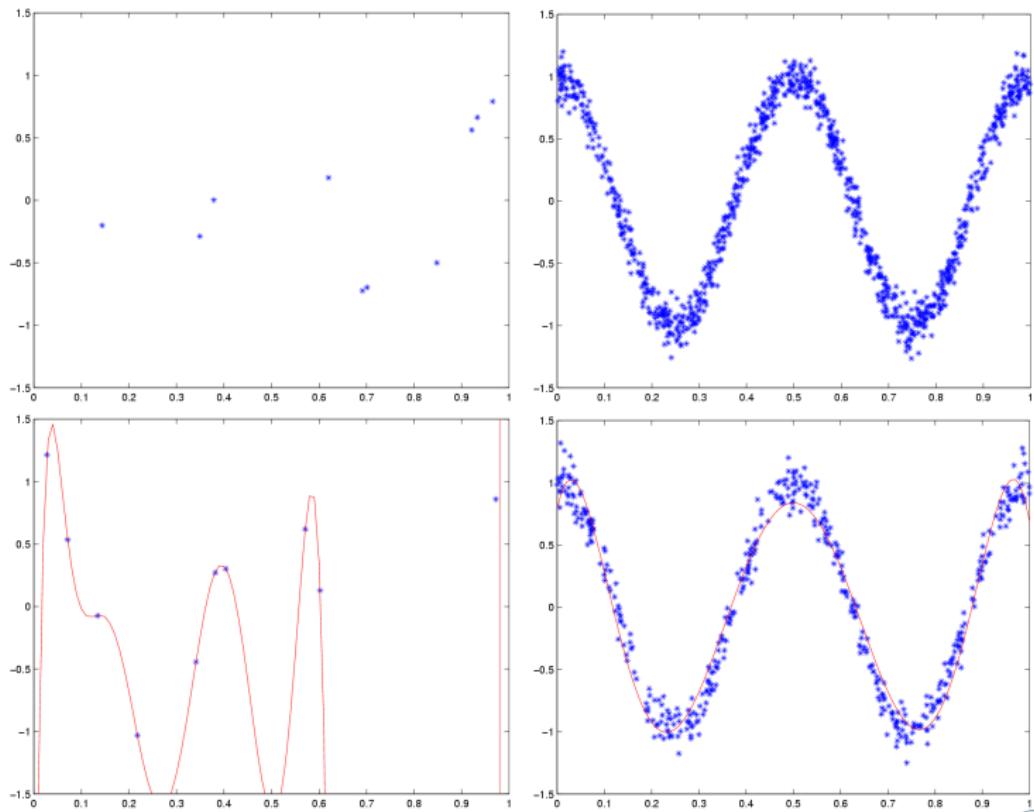
Pairs of observations  $(x_i, y_i)$  drawn from distribution  
e.g., (blood status, cancer), (credit transactions, fraud),  
(sound profile of jet engine, defect)

## Goal

**Estimate**  $y \in \{\pm 1\}$  **given**  $x$  at a new location. Or find a function  $f(x)$  that does the trick.



# Regression



# Regression

## Data

Pairs of observations  $(x_i, y_i)$  generated from some joint distribution  $\Pr(x, y)$ , e.g.,

- market index, SP100
- fab parameters, yield
- user profile, price

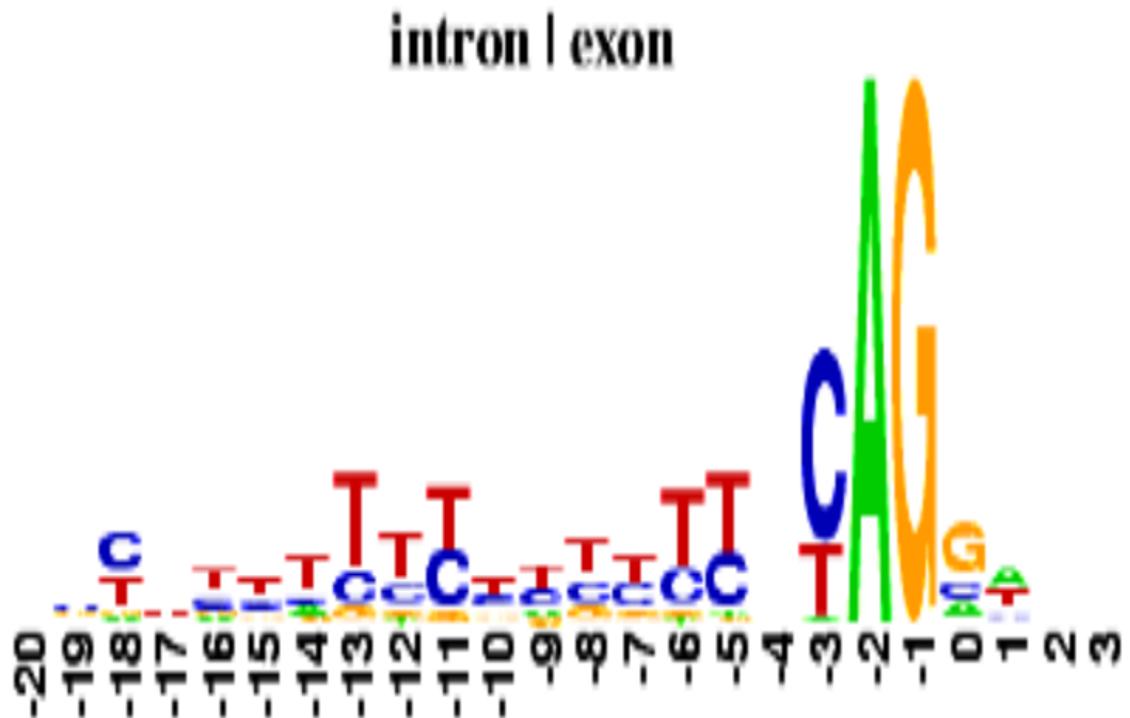
## Task

Estimate  $y$ , given  $x$ , such that some loss  $c(x, y, f(x))$  is minimized.

## Examples

- Quadratic error between  $y$  and  $f(x)$ , i.e.  
 $c(x, y, f(x)) = \frac{1}{2}(y - f(x))^2$ .
- Absolute value, i.e.,  $c(x, y, f(x)) = |y - f(x)|$ .

# Annotating Strings



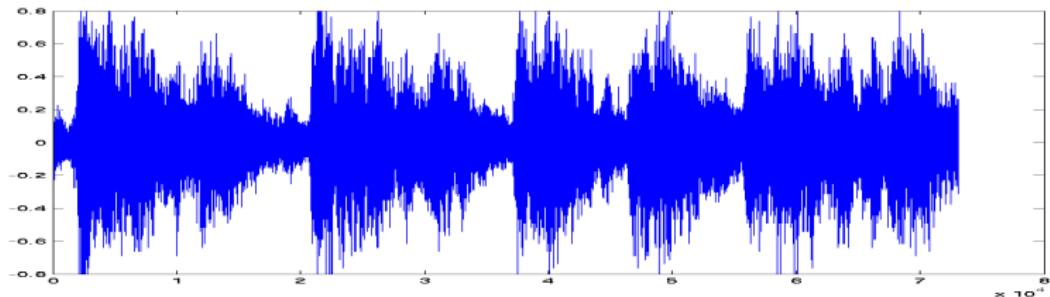
# Annotating Audio

## Goal

- Possible meaning of an audio sequence
- Give confidence measure

## Example (from Australian Prime Minister's speech)

- a stray alien
- Australian



# Novelty Detection

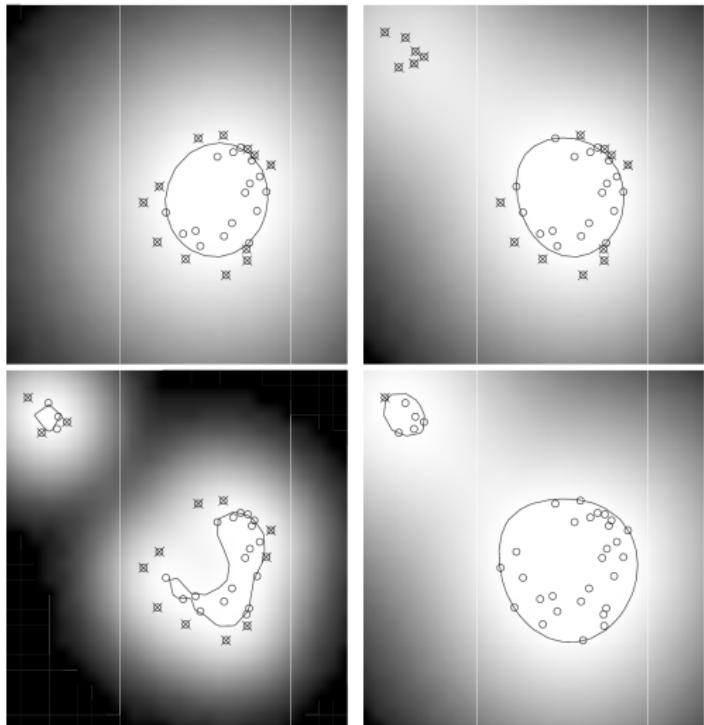
## Data

Observations  $x_i$  from some  $P(x)$ , e.g.,

- network usage patterns
- handwritten digits
- alarm sensors
- factory status

## Task

Find unusual events, clean database, distinguish typical examples.



# Learning and Similarity: some Informal Thoughts

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- input/output sets  $\mathcal{X}, \mathcal{Y}$
- training set  $(x_1, y_1), \dots, (x_m, y_m) \in \mathcal{X} \times \mathcal{Y}$
- “generalization”: given a previously unseen  $x \in \mathcal{X}$ , find a suitable  $y \in \mathcal{Y}$
- $(x, y)$  should be “similar” to  $(x_1, y_1), \dots, (x_m, y_m)$
- how to measure similarity?
  - for outputs: *loss function* (e.g., for  $\mathcal{Y} = \{\pm 1\}$ , zero-one loss)
  - for inputs: *kernel*

## Similarity of Inputs

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- symmetric function

$$\begin{aligned} k : \mathcal{X} \times \mathcal{X} &\rightarrow \mathbb{R} \\ (x, x') &\mapsto k(x, x') \end{aligned}$$

- for example, if  $\mathcal{X} = \mathbb{R}^N$ : canonical dot product

$$k(x, x') = \sum_{i=1}^N [x]_i [x']_i$$

- if  $\mathcal{X}$  is not a dot product space: assume that  $k$  has a **representation** as a dot product in a linear space  $\mathcal{H}$ , i.e., there exists a map  $\Phi : \mathcal{X} \rightarrow \mathcal{H}$  such that

$$k(x, x') = \langle \Phi(x), \Phi(x') \rangle.$$

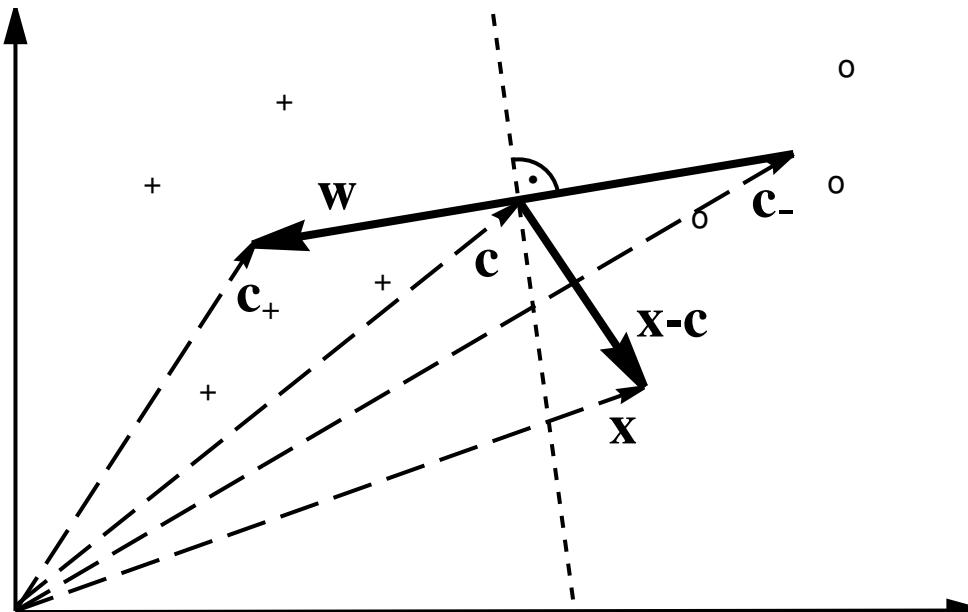
- in that case, we can think of the patterns as  $\Phi(x)$ ,  $\Phi(x')$ , and carry out geometric algorithms in the dot product space (“**feature space**”)  $\mathcal{H}$ .

## An Example of a Kernel Algorithm

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Idea: classify points  $\mathbf{x} := \Phi(x)$  in feature space according to which of the two **class means** is closer.

$$\mathbf{c}_+ := \frac{1}{m_+} \sum_{y_i=1} \Phi(x_i), \quad \mathbf{c}_- := \frac{1}{m_-} \sum_{y_i=-1} \Phi(x_i)$$



Compute the sign of the dot product between  $\mathbf{w} := \mathbf{c}_+ - \mathbf{c}_-$  and  $\mathbf{x} - \mathbf{c}$ .

## An Example of a Kernel Algorithm, ctd. [25]

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$$\begin{aligned} f(x) &= \operatorname{sgn} \left( \frac{1}{m_+} \sum_{\{i:y_i=+1\}} \langle \Phi(x), \Phi(x_i) \rangle - \frac{1}{m_-} \sum_{\{i:y_i=-1\}} \langle \Phi(x), \Phi(x_i) \rangle + b \right) \\ &= \operatorname{sgn} \left( \frac{1}{m_+} \sum_{\{i:y_i=+1\}} k(x, x_i) - \frac{1}{m_-} \sum_{\{i:y_i=-1\}} k(x, x_i) + b \right) \end{aligned}$$

where

$$b = \frac{1}{2} \left( \frac{1}{m_-^2} \sum_{\{(i,j):y_i=y_j=-1\}} k(x_i, x_j) - \frac{1}{m_+^2} \sum_{\{(i,j):y_i=y_j=+1\}} k(x_i, x_j) \right).$$

- provides a geometric interpretation of Parzen windows

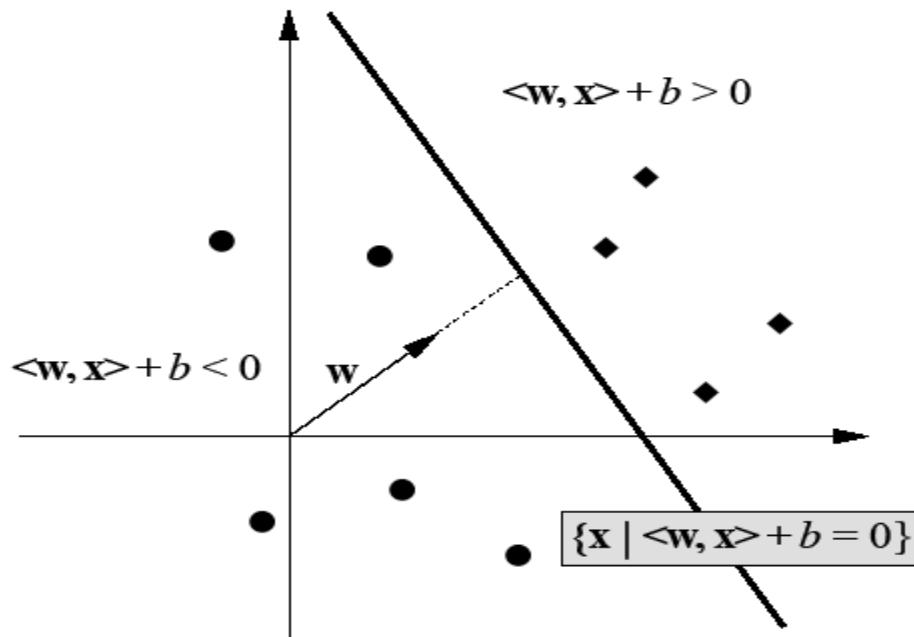
# Non-linear Classification via Support Vector Machines

- Based on several ideas:
  - linear classification
  - *increasing* the dimensionality
  - the kernel trick
  - maximizing the margin
  - duality

# Linear Classification

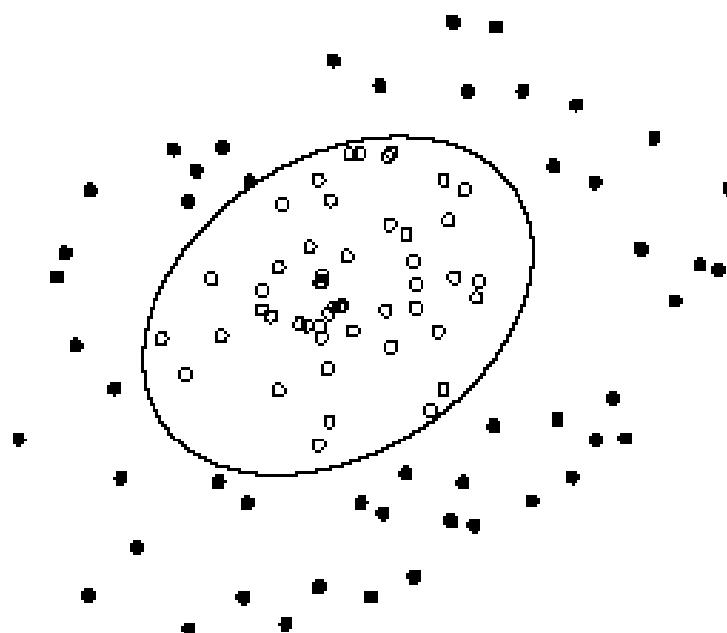
## Separating Hyperplane

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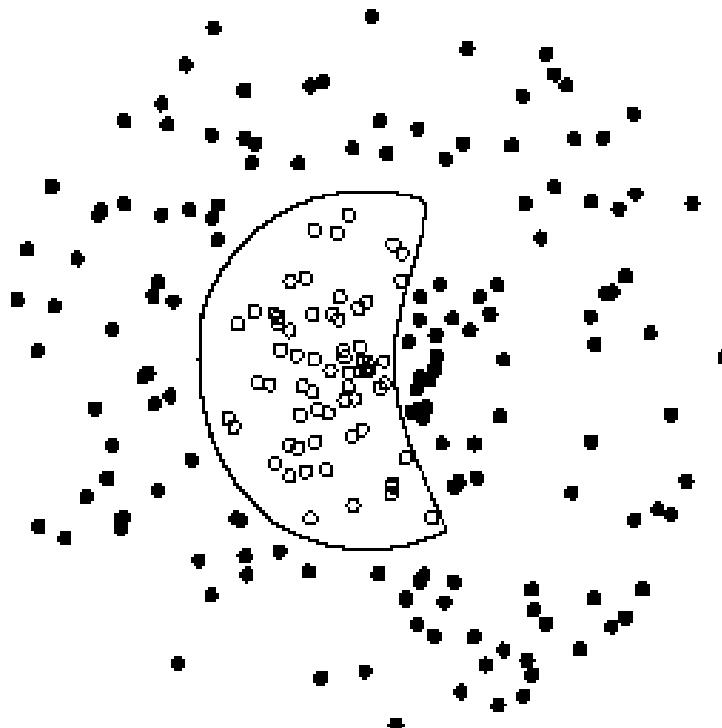


B. Schölkopf, Canberra, February 2002

# Non-linear Classification



separation by ellipsoid



separation by 4th degree polynomial

## Kernel trick

*kernel trick*: dot-products in feature space can be computed as a kernel function  $K(\mathbf{x}_i, \mathbf{x}_j) = \Phi(\mathbf{x}_i) \cdot \Phi(\mathbf{x}_j)$

idea: work directly on  $\mathbf{x}$ , avoid having to compute  $\Phi(\mathbf{x})$  at all

example:

$$\begin{aligned} K(\mathbf{a}, \mathbf{b}) &= (\mathbf{a} \cdot \mathbf{b})^3 = ((a_1, a_2) \cdot (b_1, b_2))^3 \\ &= (a_1 b_1 + a_2 b_2)^3 \\ &= a_1^3 b_1^3 + 3a_1^2 b_1^2 a_2 b_2 + 3a_1 b_1 a_2^2 b_2^2 + a_2^3 b_2^3 \\ &= ((a_1^3, \sqrt{3}a_1^2 a_2, \sqrt{3}a_1 a_2^2, a_2^3) \cdot (b_1^3, \sqrt{3}b_1^2 b_2, \sqrt{3}b_1 b_2^2, b_2^3)) \\ &= \Phi(\mathbf{a}) \cdot \Phi(\mathbf{b}) \end{aligned}$$

## Kernels

examples:

1. polynomial kernel:  $K(\mathbf{x}_i, \mathbf{x}_j) = (\mathbf{x}_i \cdot \mathbf{x}_j + 1)^z$
2. Gaussian kernel:  $K(\mathbf{x}_i, \mathbf{x}_j) = \exp(-||\mathbf{x}_i - \mathbf{x}_j||^2/2\sigma^2)$
3. sigmoid kernel:  $K(\mathbf{x}_i, \mathbf{x}_j) = \tanh(\kappa(\mathbf{x}_i \cdot \mathbf{x}_j) + a)$

each kernel computation corresponds to dot product calculation  
for particular mapping  $\Phi(\mathbf{x})$  – implicitly maps to high-dim space

why useful?

- rewrite training examples using more complex features
- dataset not linearly separable in original space may be linearly separable in higher-dimensional space