

LECTURE 13:  
KERNEL MACHINES

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FEATURE SPACES

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- The extended representation is called a feature space.
- An algorithm that is linear in the feature space may be highly nonlinear in the original space if the features contain nonlinear mappings of the raw data.
- Thus, we can think of having “promoted” our data  $x$  into a higher-dimensional feature space  $z$  using a nonlinear mapping  $\phi(x)$  and then running our original algorithm in that new space.
- The feature point  $z = \phi(x)$  corresponding to a point  $x$  is called the *image* of  $x$ .
- The point  $x$ , if any, corresponding to a given  $z$  is called the *pre-image* of  $z$ .

ALGORITHMS BASED ON VECTOR DATA

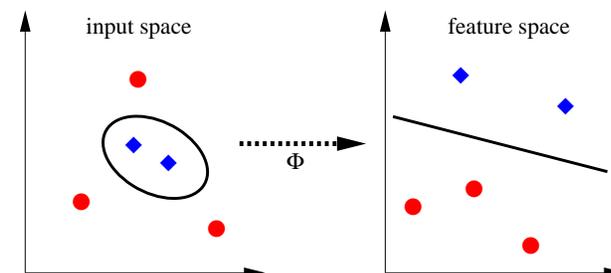
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- Recall our normal approach to many classification, regression, and unsupervised learning problems:  
Embed the input to the problem into a vector space (e.g.  $R^n$ ) and then do some geometric, or linear algebraic operations.
- We can often make our algorithms more powerful by embedding the data into a richer (larger) space which includes some fixed, possibly nonlinear functions of the original measurements.
- For example, if we measure  $x_1, x_2, x_3$  for each datapoint, we might use the representation  $z = [1, x_1, x_2, x_3, x_1^2, x_2^2, x_3^2, x_1x_2, x_2x_3]$  in a regression or classification machine.
- We’ve seen this trick before: adding a bias term, quadratic regression, basis functions, generalized linear models.

BIG FEATURE SPACES

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- Problem: the feature space may be ultra-high dimensional or even infinite dimensional, so direct (explicit) calculations in feature space may not be practical or even possible.
- What should we do?  
Restrict ourselves to manageable feature spaces? No!
- Enter the “kernel trick” .



## THE KERNEL TRICK

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- The key idea of kernel machines is to reduce an algorithm to one which *depends only on dot products between data vectors* and then to *replace* the dot product with another positive definite function  $K(x_1, x_2)$  called a *kernel*.
- This has a very deep interpretation as mapping the data into a much higher dimensional feature space  $z = \phi(x)$ , as determined by the eigenfunctions of the kernel operator, and taking a regular dot product in that space  $K(x_1, x_2) = \langle z_1, z_2 \rangle = \langle \phi(x_1), \phi(x_2) \rangle$ .
- The kernel trick allows us to efficiently compute the dot products in very high dimensional spaces using the kernel function.

## PROPERTIES OF KERNELS

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- Kernels are *symmetric* in their arguments:  $K(x_1, x_2) = K(x_2, x_1)$ .
- They are positive valued for any inputs:  $K(x_1, x_2) \geq 0$ .
- The Cauchy-Schwartz inequality still holds:  
 $K^2(x_1, x_2) \leq K(x_1, x_1)K(x_2, x_2)$ .
- Technically, to use a function as a kernel, it must satisfy “Mercer’s conditions” for a positive-definite operator.
- The intuition is easy to get for finite spaces.
  1. Discretize  $x$  space as densely as you want.
  2. Between each two cells, compute the kernel function, and write these values as a matrix. (Symmetric, obviously.)
  3. If the matrix is positive definite, the kernel is OK.

## WHEN THE KERNEL TRICK WORKS

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- The kernel trick allows us to efficiently compute the dot products in very high dimensional spaces using the kernel function.
- But it only saves us for dot products, not for addition, subtraction, outer products, etc.  
We couldn’t even write down the results of these operations anyway, so not a big loss.
- In general, everything we represent in the high-dimensional feature space must be representable as a *linear combination of the images of training data points*.
- Actually this turns out to be fine for many problems, e.g. the optimal weights in a SVM/kernel perceptron are representable. (There is more theory about this under (surprise surprise) the “Representer’s Theorem”. [originally by Kimeldorf and Wahaba, reproved by Schoelkopf et al]).

## EXAMPLES OF KERNELS

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- Linear:  $K(x_1, x_2) = \mathbf{x}_1^\top \mathbf{x}_2$
- Gaussian:  $K(x_1, x_2) = \exp[-.5 * \|\mathbf{x}_1 - \mathbf{x}_2\|^2]$
- Polynomial:  $K(x_1, x_2) = (1 + \mathbf{x}_1^\top \mathbf{x}_2)^k$  (watch scaling!)
- Tanh:  $K(x_1, x_2) = \tanh(a * \mathbf{x}_1^\top \mathbf{x}_2 + b)$
- Closure rules:
  - The sum of any two kernels is a kernel.
  - The product of any two kernels is a kernel.
  - A kernel plus a constant is a kernel.
  - A scalar times a kernel is a kernel.

## GEOMETRY OF FEATURE SPACE

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- Dot product between two points =  $K(x_i, x_j) > 0$ , and so all points lie in a single orthant in feature space.

- Length of a point in feature space:

$$\|z_i\|^2 = K(x_i, x_i)$$

(so for Gaussian kernels, everybody lies on surface of unit sphere)

- Distance between two points in feature space:

$$\|z_1 - z_2\|^2 = K(x_1, x_1) + K(x_2, x_2) - 2K(x_1, x_2)$$

- Distance between a point and the mean of all others:

$$\|z_k - \bar{z}\|^2 = K(x_k, x_k) + \frac{1}{N^2} \sum_{ij} K(x_i, x_j) - 2 \sum_i K(x_k, x_i)$$

- Zero mean calculations in feature space:

$$\langle z_i - \bar{z}, z_j - \bar{z} \rangle = K(x_i, x_j) + \frac{1}{N^2} \sum_{k\ell} K(x_k, x_\ell) - \sum_k K(x_k, x_i) \sum_k K(x_j, x_k)$$

## GRAM MATRIX

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- But first, some notation...
- The “Gram Matrix” is the  $N$  by  $N$  symmetric matrix of all pairwise kernel evaluations:  $G_{ij} = K(x_i, x_j)$ .
- If you successfully “kernelize” an algorithm, then you can build the Gram matrix and throw away the original data.
- Your algorithm will only need to consult entries of the Gram matrix as it runs, because it depends only on dot products between the data points.
- An equivalent characterization to Mercer’s conditions is that a valid kernel generates symmetric positive definite Gram matrices for any finite sample of raw data  $x_n$ . (Saitoh)

## DOT-PRODUCTING AN ALGORITHM

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- The art of designing a kernel machine is to take a standard algorithm and massage it so that all references to the original data vectors  $x$  appear only in dot products  $\langle x_i, x_j \rangle$ .
- Often you can do this and obtain an exactly equivalent algorithm to the one you started with. Sometimes you need to make small modifications.
- Thus, a kernel machine contains two modules: the algorithm and the kernel function. Choosing the kernel function is a hard problem which we won’t discuss today.
- “Kernelizing” an algorithm can actually be pretty easy. How about a few examples...

## EXAMPLE: K-NN CLASSIFICATION

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- Distance between two points in feature space:

$$\|z_1 - z_2\|^2 = K(\mathbf{x}_1, \mathbf{x}_1) + K(\mathbf{x}_2, \mathbf{x}_2) - 2K(\mathbf{x}_1, \mathbf{x}_2)$$

### EXAMPLE: K-MEANS CLUSTERING

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- Distance between two points in feature space:

$$\|\mathbf{z}_1 - \mathbf{z}_2\|^2 = K(\mathbf{x}_1, \mathbf{x}_1) + K(\mathbf{x}_2, \mathbf{x}_2) - 2K(\mathbf{x}_1, \mathbf{x}_2)$$

- Represent cluster centres as linear combinations of data points:

$$\mathbf{c}_k = \sum_i \alpha_{ik} \mathbf{z}_i$$

- Distance between a new point and a cluster centre in feature space:

$$\|\mathbf{z} - \mathbf{c}_k\|^2 = K(\mathbf{x}, \mathbf{x}) + \sum_{ij} \alpha_{ik} \alpha_{jk} K(\mathbf{x}_i, \mathbf{x}_j) - 2 \sum_i \alpha_{ik} K(\mathbf{x}, \mathbf{x}_i)$$

### KERNEL PERCEPTRON

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- The update rule for the weight vector in the perceptron can also be rewritten in terms of dot products only.
- Old rule: if  $y_i \mathbf{w}^\top \mathbf{x}_i + b \leq 0$  then  $\mathbf{w} \leftarrow \mathbf{w} + y_i \mathbf{x}_i$ .
- Recall our new representation:  $\mathbf{w} = \sum_i (\alpha_i y_i) \mathbf{z}_i$
- Equivalent new update rule:  
if  $y_i \sum_j \alpha_j y_j \mathbf{z}_i^\top \mathbf{z}_j + b \leq 0$  then  $\alpha_i \leftarrow \alpha_i + 1$ .

### EXAMPLE: PERCEPTRON CLASSIFICATION

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- The regular perceptron (hyperplane classifier) was:

$$f(\mathbf{x}) = \text{sign}[\mathbf{w}^\top \mathbf{x} + b]$$

- To kernelize, we must represent the weights as linear combinations of the input vector images (but this is OK):

$$\mathbf{w} = \sum_i (\alpha_i y_i) \mathbf{z}_i$$

- The original can be rewritten in terms of dot products:

$$f(\mathbf{z}) = \text{sign}\left[\sum_i (\alpha_i y_i) \mathbf{z}^\top \mathbf{z}_i\right]$$

### EXAMPLE: RIDGE REGRESSION

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- Think of the ridge regression cost function:  
 $\sum_i (y_i - \mathbf{w}^\top \mathbf{x}_i)^2 + \lambda \|\mathbf{w}\|^2$   
minimizing this is equivalent to minimizing:  
 $\sum_i \eta_i^2 + \lambda \|\mathbf{w}\|^2$  subject to  $\eta_i = (y_i - \mathbf{w}^\top \mathbf{x}_i)$ .
- Let's introduce Lagrange multipliers to enforce the constraints:  
 $\min \sum_i \eta_i^2 + \lambda \|\mathbf{w}\|^2 + \sum_i \alpha_i (\eta_i - (y_i - \mathbf{w}^\top \mathbf{x}_i))$
- Setting partial derivatives to zero gives:  
 $\mathbf{w}^* = (1/2\lambda) \mathbf{1} + \sum_i \alpha_i \mathbf{x}_i$  and  $\eta_i = \alpha_i / 2$
- Plugging back into the original cost gives:  
 $\min \sum_i y_i \alpha_i - \frac{1}{4\lambda} \sum_{ij} \alpha_i \alpha_j \mathbf{x}_i^\top \mathbf{x}_j - \frac{1}{4} - \sum_i \alpha_i^2$
- In matrix form:  
 $\min \mathbf{y}^\top \alpha - \frac{1}{4\lambda} \alpha^\top G \alpha - \frac{1}{4}$
- Completely Kernelized!

### EXAMPLE: PCA

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- Amazing!
- Standard PCA (assume data is zero mean):

$$C = \frac{1}{N} = \sum_i \mathbf{x}_i \mathbf{x}_i^\top$$
$$\lambda \mathbf{v} = C \mathbf{v}$$

- All eigenvectors with nonzero eigenvalues must lie in the span of the data, and thus can be written as linear combinations of the data (why?):

$$\mathbf{v} = \sum_i \alpha_i \mathbf{x}_i$$

- Now we can rewrite the eigenvector condition:

$$\lambda \sum_i \alpha_i \mathbf{x}_i = \frac{1}{N} \sum_{ij} \alpha_i \mathbf{x}_j \mathbf{x}_i^\top \mathbf{x}_j$$

### EXAMPLE: FISHER DISCRIMINANT

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- See if you can figure this one out on your own...  
(or look it up)

### EXAMPLE: PCA

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- Eigenvector condition:

$$\lambda \sum_i \alpha_i \mathbf{x}_i = \frac{1}{N} \sum_{ij} \alpha_i \mathbf{x}_j \mathbf{x}_i^\top \mathbf{x}_j$$

- Take the dot product with  $\mathbf{x}_k$  on left and right:

$$\lambda \sum_i \alpha_i \mathbf{x}_k^\top \mathbf{x}_i = \frac{1}{N} \sum_{ij} \alpha_i \mathbf{x}_k^\top \mathbf{x}_j \mathbf{x}_i^\top \mathbf{x}_j$$

- The above is true for all  $k$ , so write it as a vector equation in  $\alpha$ :

$$N \lambda G \alpha = G^2 \alpha$$
$$N \lambda \alpha = G \alpha$$

- Result: Form  $G$ , find its eigenvectors  $\alpha$ , and use these to construct linear combinations of original data points which are the eigenvectors of the original covariance matrix.  
(Careful! Zero mean and normalization of eigenvectors.)

### THE DESIRE FOR SPARSITY

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- In kernel machines, the principal trick is to convert the problem into a *dual* form, which usually involves representing everything in the feature space as a linear combination of images of the training points:  $\mathbf{z} = \sum_i \alpha_i \phi(\mathbf{x}_i)$
- Then we do all our calculations with the dual variables  $\alpha_i$  and we never have to “touch” feature space directly.
- For very large datasets, it is desirable to have many of the coefficients  $\alpha_i$  be exactly zero (sparsity) to reduce computational load, especially at test time.
- As a *separate trick*, different from the kernel trick, we can look for ways to make things sparse.
- These tricks are often confused, because the most famous kernel machine (the SVM) used them both.

## CONTROLLING OVERFITTING

- A *third* aspect to kernel machines is how to control overfitting.
- For example, in the perceptron, if we use a very nonlinear kernel, we might always be able to separate our data exactly. Then we could be seriously overfitting.
- We can use *weight decay* to prevent this, by penalizing  $\|\mathbf{w}\|^2$  in addition to trying to separate our training sample in the feature space. This is equivalent to *maximum margin*.
- A deep motivation for weight decay in this context comes from minimizing error bounds based on the “VC dimension” theory.
- This idea is often confused with the kernel & sparsity tricks because the most famous kernel machine (the SVM) also used weight decay to control overfitting and discussed the VC motivation.

## MAXIMUM MARGIN = MINIMUM NORM

- Maximizing the margin is equivalent to picking the separating hyperplane that minimizes the norm of the weight vector:  

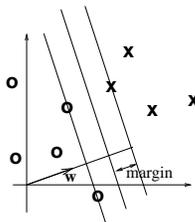
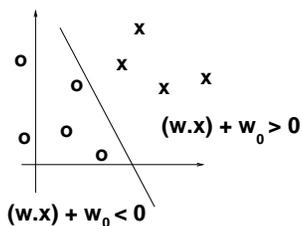
$$\min \|\mathbf{w}\|^2 \quad \text{subject to} \quad y_i[\mathbf{w}^\top \mathbf{x}_i + b] \geq 1$$
- Use Lagrange multipliers to enforce the constraint:  

$$\min \|\mathbf{w}\|^2 - \sum_i \alpha_i (y_i[\mathbf{w}^\top \mathbf{x}_i + b] - 1) \quad \alpha_i \geq 0$$
- We can convert it to *dual form* by setting partial derivatives to zero and substituting.

## MAXIMUM MARGIN HYPERPLANE

- Margin = minimum distance to the plane of any point.
- Principle: of all the hyperplanes that separate the data perfectly, pick the one which maximizes the margin
- Since the scale is arbitrary, we will set the numerical value of the margin to be 1.
- Now maximizing the margin is equivalent to picking the separating hyperplane that minimizes the norm of the weight vector:  

$$\min \|\mathbf{w}\|^2 \quad \text{subject to} \quad y_i[\mathbf{w}^\top \mathbf{x}_i + b] \geq 1$$



## PRIMAL $\rightarrow$ DUAL

- Use Lagrange multipliers to enforce the constraint:  

$$\min \|\mathbf{w}\|^2 - \sum_i \alpha_i (y_i[\mathbf{w}^\top \mathbf{x}_i + b] - 1) \quad \alpha_i \geq 0$$
- set  $\partial/\partial \mathbf{w} = 0$  and  $\partial/\partial b = 0$ :  $\mathbf{w}^* = \sum_i y_i \alpha_i \mathbf{x}_i \quad \sum_i y_i \alpha_i = 0$
- The dual problem is now:  $\min \sum_i \alpha_i - \frac{1}{2} \sum_{i,j} \alpha_i \alpha_j y_i y_j \mathbf{x}_i^\top \mathbf{x}_j$   

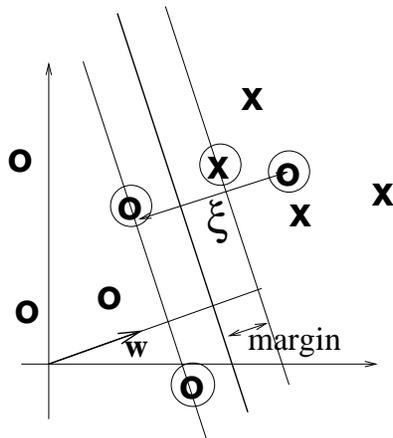
$$\alpha_i \geq 0 \quad \sum_i y_i \alpha_i = 0$$
- This is a *quadratic programming* problem.
- It is convex. Unique solution!

## SPARSITY OF SOLUTION

- Not only is the solution unique, but it is also sparse.
- Only the training points nearest to the separating hyperplane (ie with margin exactly 1) have  $\alpha_i > 0$ . These points are called the “active” points, or the *support vectors* since the final weight vector depends only on them:

$$\mathbf{w}^* = \sum_i y_i \alpha_i \mathbf{x}_i$$

- This is a lucky coincidence that has confused many people: in the case of SVM classification the two goals of controlling overfitting and inducing sparsity can both be achieved simultaneously with only a single trick: maximum margin (minimum weight norm).
- But it is not always like this.



## SUPPORT VECTOR MACHINES

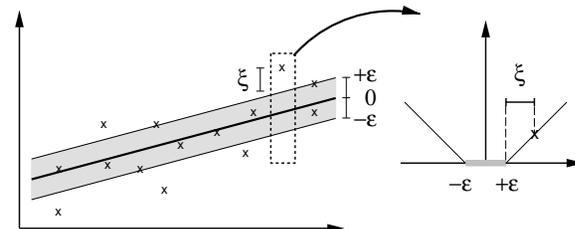
- A *support vector machine* (SVM) is nothing more than a kernelized maximum-margin hyperplane classifier.
- You train it by solving the dual quadratic programming problem.
- You run it by finding dot products of the test point with all the training cases.
- Easy!

## SPARSITY IN REGRESSION

- To introduce sparsity in regression, Vapnik introduced the *epsilon-insensitive loss function*:

$$l(\hat{y}) = 0 \quad \text{if} \quad |y - \hat{y}| \leq \epsilon$$

$$l(\hat{y}) = |y - \hat{y}| \quad \text{if} \quad |y - \hat{y}| > \epsilon$$



## LOTS MORE

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- VC Dimension and Error Bounds
- Other kernel machines (e.g. Gaussian processes)
- see <http://www.kernel-machines.org>  
for lots of papers/tutorials, etc

## THANKS! IT'S BEEN FUN!

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- Last class.
- Thanks for sticking with it.
- Hope you learned something, and had fun also.
- Sorry about all the math.
- Please send me comments/corrections for my book and for next year.