

# CSCE 990 Lecture 9: Designing Kernels\*

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## Introduction

- We are now very aware of the importance and power of kernels in SVMs
- We also know from Chapter 2 about some basic kernels and simple ways to build new kernels out of old ones
  - Linear scaling, addition, multiplication, etc. of existing kernels
- We'll look at other ways to construct new kernels from existing ones, plus other completely different types of kernels
- Some of them might look familiar ...

## Outline

- Tricks for constructing kernels
- String kernels
- Spectrum kernels
- Locality-improved kernels
- Kernels defined on graphs
- Sections 13.1–13.3, 13.5, assorted papers

## Tricks for Constructing Kernels

- If  $k_1$  and  $k_2$  are kernels, then so are

$$\alpha_1 k_1 + \alpha_2 k_2 \text{ for } \alpha_1, \alpha_2 \geq 0$$

⇒ If input vectors can be partitioned into subvectors of different types (e.g. strings and real values), can apply direct sum:

$$(k_1 \oplus k_2)(x_1, x_2, x'_1, x'_2) = k_1(x_1, x'_1) + k_2(x_2, x'_2)$$

where  $x_1, x'_1 \in \mathcal{X}_1$  (e.g.  $\mathbb{R}^n$ ) and  $x_2, x'_2 \in \mathcal{X}_2$  (e.g. strings)

$$k_1 k_2$$

⇒ Similar to application of direct sum, use tensor product:

$$(k_1 \otimes k_2)(x_1, x_2, x'_1, x'_2) = k_1(x_1, x'_1) k_2(x_2, x'_2)$$

# Tricks for Constructing Kernels

## Conformal Transformations

- For a real-valued function  $f$ ,  $k'(x, x') = f(x)f(x')$  is a kernel
- This leads to conformal transformations:

$$k_f(x, x') = f(x)k(x, x')f(x')$$

- If  $k$  is a kernel, then so is  $k_f$
- Recall that if  $\|x\| = \|x'\| = 1$ , then  $\langle x, x' \rangle = \cos(\angle(x, x'))$ ; thus

$$\begin{aligned}\cos(\angle(\Phi_f(x), \Phi_f(x'))) &= \frac{f(x)k(x, x')f(x')}{\sqrt{f(x)k(x, x)f(x)}\sqrt{f(x')k(x', x')f(x')}} \\ &= \frac{k(x, x')}{\sqrt{k(x, x)}\sqrt{k(x', x')}} \\ &= \cos(\angle(\Phi(x), \Phi(x')))\end{aligned}$$

I.e. angles in feature space are preserved in a conformal transformation

# Tricks for Constructing Kernels

## Convolution Kernels

- Notions of tensor products and direct sums lead to  $R$ -convolution kernels
- E.g. consider partitioning the string  $x = ATG$  into two distinct, contiguous, nonempty substrings:

$$R_1 : \quad x_{1,R_1} = A \quad \text{AND} \quad x_{2,R_1} = TG$$

OR

$$R_2 : \quad x_{1,R_2} = AT \quad \text{AND} \quad x_{2,R_2} = G$$

(similarly, decompose  $x'$ )

- Now can compute a kernel for each substring of each partitioning and combine:

$$\begin{aligned} k(x, x') &= k_1(x_{1,R_1}, x'_{1,R_1})k_2(x_{2,R_1}, x'_{2,R_1}) \\ &\quad + k_1(x_{1,R_2}, x'_{1,R_2})k_2(x_{2,R_2}, x'_{2,R_2}) \end{aligned}$$

# Tricks for Constructing Kernels

## Convolution Kernels (cont'd)

- Generally, define the set of allowed decompositions as a relation  $R(x_1, \dots, x_D, x)$  and define the  $R$ -convolution

$$(k_1 \star \dots \star k_D)(x, x') := \sum_R \prod_{d=1}^D k_d(x_d, x'_d)$$

(i.e. sum over all allowable decompositions of  $x$  into  $x_1, \dots, x_D$ , etc.)

- Based on earlier results, we know this to be a valid kernel
- A special case: ANOVA kernel of order  $D$

$$k_D(x, x') := \sum_{1 \leq i_1 < \dots < i_D \leq N} \prod_{d=1}^D k^{(i_d)}(x_{i_d}, x'_{i_d})$$

( $D = N \Rightarrow$  tensor prod,  $D = 1 \Rightarrow$  direct sum)

## String Kernels

- To apply SVMs to text classification, can map documents to bag-of-words representation and use kernels defined on  $\mathbb{R}^n$ 
  - Each dimension is one word, value in that dimension is word frequency
  - Ignores word ordering
- Alternatively, can use a string kernel, which computes similarities between two strings based on their common substrings
- Related to  $R$ -convolution kernel



# String Kernels

(cont'd)

- Let  $\Sigma$  be a finite alphabet,  $\Sigma^n$  be set of all length- $n$  strings over  $\Sigma$ , and  $\Sigma^* = \bigcup_{n=0}^{\infty} \Sigma^n$
- Given  $s \in \Sigma^*$ , let  $\mathbf{i} := (i_1, \dots, i_{|u|})$  be an index sequence with  $1 \leq i_1 < \dots < i_{|u|} \leq |s|$  and  $u := s(\mathbf{i}) := s(i_1) \dots s(i_{|u|})$  be a (possibly non-contiguous) subsequence of  $s$
- $l(\mathbf{i}) := i_{|u|} - i_1 + 1$  is the length of  $u$  in  $s$ 
  - E.g. if  $s = ABBA$ , then  $l(1, 2, 3) = 3$  (for  $ABB$ ),  $l(1, 4) = 4$  (for  $AA$ )
  - $\Phi_n(s)$  defines one dimension per substring  $u \in \Sigma^n$ , and the  $u$ th component of  $\Phi_n(s)$  is

$$[\Phi_n(s)]_u := \sum_{\mathbf{i}: s(\mathbf{i})=u} \lambda^{l(\mathbf{i})}$$

for  $0 < \lambda \leq 1$

## String Kernels

(cont'd)

- E.g. if  $s = ABBA$ , then  $[\Phi_2(s)]_{AB} = \lambda^2 + \lambda^3$
- $[\Phi_n(s)]_u$  larger if  $u$  (nearly) contiguous and common in  $s$
- The string kernel is then

$$\begin{aligned} k_n(s, t) &= \sum_{u \in \Sigma^n} [\Phi_n(s)]_u [\Phi_n(t)]_u \\ &= \sum_{u \in \Sigma^n} \sum_{(i,j): s(i)=t(j)=u} \lambda^{l(i)} \lambda^{l(j)} \end{aligned}$$

- If want to vary  $n$ , use  $k := \sum_n c_n k_n$
- Since value of  $k_n$  (and therefore  $k$ ) depend on lengths of  $s$  and  $t$ , normalize  $k$  in feature space

## String Kernels

(cont'd)

- To efficiently compute the kernel, define for  $i = 1, \dots, n - 1$

$$k'_i(s, t) = \sum_{u \in \Sigma^i} \sum_{(\mathbf{i}, \mathbf{j}) : s(\mathbf{i}) = t(\mathbf{j}) = u} \lambda^{|s| + |t| - i_1 - j_1 + 2}$$

- Then if  $x \in \Sigma^1$ , can recursively compute  $k_n(s, t)$ :

$$\begin{aligned} k'_0(s, t) &= 1 \quad \text{for all } s, t \\ k'_i(s, t) &= 0 \quad \text{if } \min(|s|, |t|) < i \\ k_i(s, t) &= 0 \quad \text{if } \min(|s|, |t|) < i \end{aligned}$$

$$k'_i(sx, t) = \lambda k'_i(s, t) + \sum_{j: t_j = x} k'_{i-1}(s, t[1, \dots, j-1]) \lambda^{|t| - j + 2}$$

$$k_n(sx, t) = k_n(s, t) + \sum_{j: t_j = x} k'_{n-1}(s, t[1, \dots, j-1]) \lambda^2$$

# Spectrum Kernel

- Another type of string kernel
- For a fixed integer  $\gamma \geq 1$ , define the  $\gamma$ -spectrum of a sequence to be the set of all length- $\gamma$  contiguous sequences it contains
- Feature map for spectrum kernel is indexed by all possible length- $\gamma$  subsequences from alphabet  $\Sigma$  (similar to bag of words)
- For each  $a \in \Sigma^\gamma$ , let  $\phi_a(x)$  = number of times  $a$  occurs in  $x$  contiguously
- Now define  $\Phi_\gamma(x) = (\phi_a(x))_{a \in \Sigma^\gamma}$ 
  - This is a weighted representation of  $x$ 's  $\gamma$ -spectrum
  - A sparse vector

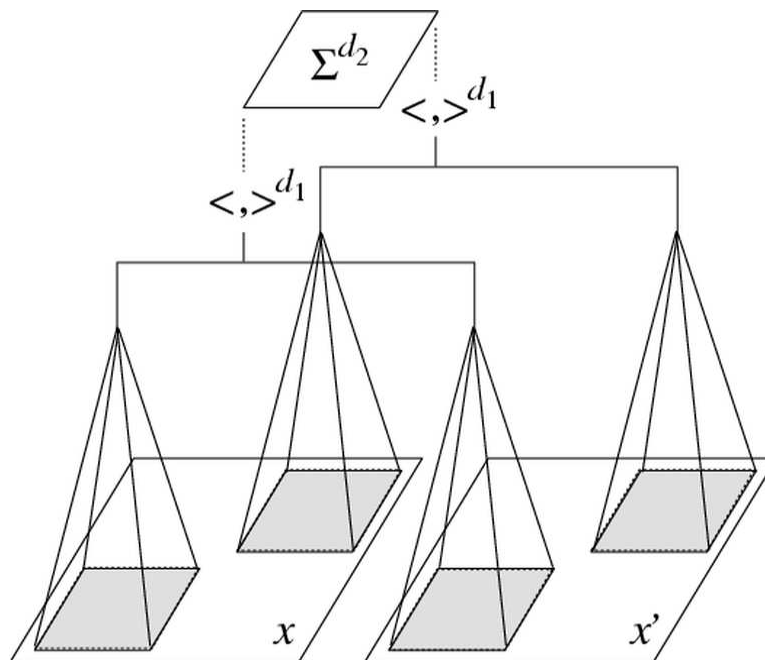
## Spectrum Kernels

(cont'd)

- Can compute  $k_\gamma(x, x') = \langle \Phi_\gamma(x), \Phi_\gamma(x') \rangle$  in time  $O(|x| + |x'|)$ 
  1. Collect set of length- $\gamma$  subsequences of  $x$  into array  $A_x$  and sort it (same with  $x'$ )
    - $A_x$  contains non-zero entries of  $\Phi_\gamma(x)$
  2. Scan  $A_x$  and  $A_{x'}$ , multiplying entries that match, and sum the products

# Locality-Improved Kernels

- A variation on existing kernels to emphasize local correlations over long-range (global) ones
- E.g. in image processing, replace polynomial kernel  $\langle x, x' \rangle^d$  with a variant that focuses on subimages first
- Generally, take the dot product over all corresponding subimages of the two images, raise to the  $d_1$  power, sum these values, then raise to the  $d_2$  power



# Locality-Improved Kernels

## Image Processing (cont'd)

- Specifically:

1. Compute  $(x.*x')$ , the pixel-wise product of  $x$  and  $x'$
2. Sample  $(x.*x')$  with pyramidal receptive fields:

$$z_{ij} := \sum_{i',j'} w(\max(|i - i'|, |j - j'|)) (x.*x')_{i'j'}$$

where e.g. weighting function  $w(n) = \max(q - n, 0)$ ; i.e. only include pixels in a width- $p$  window ( $p = 2q + 1$ ) centered at  $(i, j)$

3. Raise each  $z_{ij}$  to the  $d_1$  power (this gives local correlations)
  4. Sum  $z_{ij}^{d_1}$  over entire image and raise this sum to the  $d_2$  power (long-range correlations)
- If  $d_1 = 1$ , get standard polynomial kernel  $\langle x, x' \rangle^{d_2}$

## Locality-Improved Kernels

Image Processing (cont'd)

Classifier	Error on MNIST (%)
$k^{1,4}$	4.0
$k_9^{2,2}$	3.1
$k_9^{4,1}$	3.4
Virt SV	2.8
VSV $k_9^{2,2}$	2.0



## Locality-Improved Kernels

### DNA Start Codon Recognition

- Problem: in a DNA sequence (from alphabet  $\{A, C, T, G\}$ ), identify subsequences that encode genes
  - Typically such a coding region begins with *ATG*
  - But not all *ATG* occurrences imply a coding region
  - Thus the learning problem is to take a length-200 window centered at an *ATG* and predict if it's a coding region
- For this problem, long-range dependencies aren't very important, so use a kernel to emphasize local correlations

## Locality-Improved Kernels

### DNA Start Codon Recognition (cont'd)

- We'll consider correlations inside small windows of length  $2\ell + 1$ :

$$\text{win}_p(x, x') = \left( \sum_{j=-\ell}^{+\ell} v_j \text{match}_{p+j}(x, x') \right)^{d_1}$$

where  $\text{match}_{p+j}(x, x') = 1$  if  $x$  and  $x'$  match at position  $p + j$  and 0 otherwise, and  $v_j$  is a weight for window position  $j$  (larger near 0)

- Now we sum the values of  $\text{win}_p$ :

$$k(x, x') = \left( \sum_{p=1}^{\ell} \text{win}_p(x, x') \right)^{d_2}$$

(Should summation really be only to  $\ell$ ?)

Classifier	Error (%)
ANN	15.4
Poly kernel, $d = 1$	13.8
L-I kernel, $d_1 = 4, \ell = 4$	11.9
Codon-improved kernel, $d_1 = 2, \ell = 3$	12.2

## Kernels on Graphs

- Very general form of structured data
- Can represent many data types, including chemical structures
- Will consider directed graphs with labels on edges and nodes
- Let  $\mathcal{G}$  be the space of all graphs, modulo isomorphism

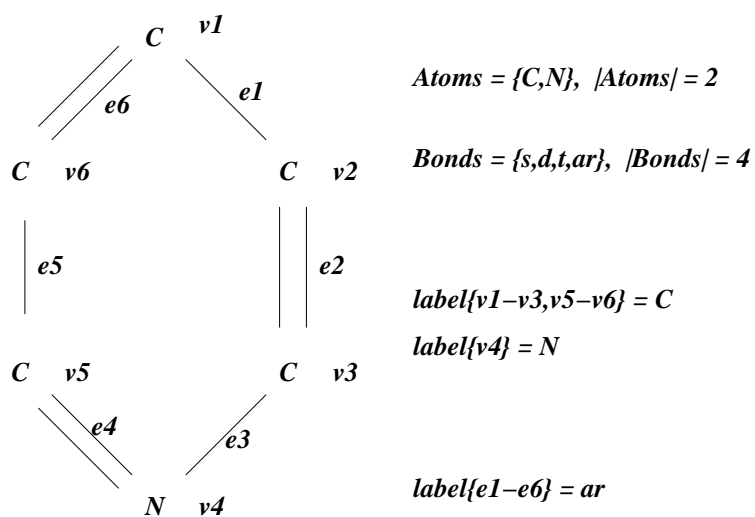
## Complete Graph Kernels

- A complete graph kernel  $k$  is one whose implicit remapping  $\Phi : \mathcal{G} \rightarrow \mathcal{H}$  distinguishes all pairs of graphs  $(G, G') \in \mathcal{G} \times \mathcal{G}$ , i.e.  $\Phi$  is injective
- Example (Subgraph feature space): Let each dimension in  $\Phi(G)$  correspond to a distinct connected graph  $H \in \mathcal{G}$ . Then  $[\Phi(G)]_H =$  number of times an isomorphism of  $H$  appears in  $G$ .
- Gärtner et al. [2003] showed that for injective  $\Phi$ ,  $k(G, G) + k(G', G') - 2k(G, G') = \langle \Phi(G) - \Phi(G'), \Phi(G) - \Phi(G') \rangle = 0$  iff  $G \simeq G'$   
 $\Rightarrow$  Computing  $k$  is as hard as graph isomorphism, for which no efficient algorithm is currently known
- Further, the kernel for the subgraph mapping is in fact NP-hard to compute (reduce from Hamiltonian path), even to approximate and/or if  $H$  comes from a restricted class of graphs

## Kernels Based on Label Pairs

- Now consider more restrictive kernels that can be efficiently considered
- Focus on graphs with labels on nodes but not edges; labels come from  $\mathcal{L} = \{\ell_1, \dots, \ell_m\}$
- Let label matrix  $L$  be such that  $[L]_{ri} = 1$  if node  $v_i$ 's label is  $\ell_r$  and  $[L]_{ri} = 0$  otherwise
- Let adjacency matrix  $E$  be such that  $[E]_{ij} = 1$  if directed edge  $(v_i, v_j)$  exists in graph  $G$  and  $[E]_{ij} = 0$  otherwise;  $[E^n]_{ij}$  is number of length- $n$  walks from  $v_i$  to  $v_j$
- $[LL^\top]_{rr}$  = number of times label  $\ell_r$  is assigned to a vertex in  $G$
- $[LE^nL^\top]_{ij}$  = number of walks of length  $n$  between vertices labeled  $\ell_i$  and vertices labeled  $\ell_j$

# Matrix Example



$$L = \begin{bmatrix} 1 & 1 & 1 & 0 & 1 & 1 \\ 0 & 0 & 0 & 1 & 0 & 0 \end{bmatrix} \quad E = \begin{bmatrix} 0 & 1 & 0 & 0 & 0 & 1 \\ 1 & 0 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 & 0 & 1 \\ 1 & 0 & 0 & 0 & 1 & 0 \end{bmatrix}$$

$$LEL^\top = \begin{bmatrix} 8 & 2 \\ 2 & 0 \end{bmatrix} \quad LE^2L^\top = \begin{bmatrix} 18 & 2 \\ 2 & 2 \end{bmatrix}$$

## Kernels Based on Label Pairs

(cont'd)

- $\mathcal{W}_n(G)$  = set of all  $n$ -edge walks in  $G$
- For walk  $w \in \mathcal{W}_n(G)$ ,  $l_1(w)$  = label of first vertex of  $w$  and  $l_{n+1}(w)$  = label of last vertex
- $\lambda$  = sequence of nonnegative weights  $\lambda_0, \lambda_1, \dots$
- Define mapping  $\Phi(G)$  to have one feature per pair of labels  $(\ell_i, \ell_j)$ :  $[\Phi(G)]_{\ell_i, \ell_j} =$

$$\sum_{n=0}^{\infty} \lambda_n \left| \left\{ w \in \mathcal{W}_n(G) : l_1(w) = \ell_i \wedge l_{n+1}(w) = \ell_j \right\} \right|$$

i.e. the weighted sum of the number of length- $n$  walks from an  $\ell_i$ -labeled vertex to an  $\ell_j$ -labeled vertex, weighted by  $\lambda_n$ , summed over all  $n \rightarrow \infty$

## Kernels Based on Label Pairs

(cont'd)

- Thus kernel is  $\langle \Phi(G), \Phi(G') \rangle =$

$$\left\langle L \left( \sum_{i=0}^{\infty} \lambda_i E^i \right) L^\top, L' \left( \sum_{i=0}^{\infty} \lambda_i E'^i \right) L'^\top \right\rangle$$

- Under certain conditions, can efficiently compute the matrix power series
- E.g. if  $\lambda_i = \beta^i / i!$  for some  $\beta > 0$  and if  $E$  can be diagonalized such that  $E = T^{-1}DT$ , then  $E^n = T^{-1}D^nT$  and  $[D^n]_{ii} = [D_{ii}]^n$  since  $D$  is diagonal
- Now we can compute

$$\lim_{n \rightarrow \infty} \sum_{i=0}^n \frac{(\beta E)^i}{i!}$$

as

$$T^{-1} \left( \lim_{n \rightarrow \infty} \sum_{i=0}^n \frac{\beta^i D^i}{i!} \right) T ,$$

where limits are taken component-wise



## Kernels Based on Contiguous Label Sequences

- Previous kernel's mapping  $\Phi$  has a low-dimensional feature space:  $|\mathcal{L}|^2$   
  
 $\Rightarrow$  E.g. if all node labels are C or N, then feature space has dimension 4
- For a more expressive feature mapping, will use mapping with one dimension per label sequence rather than label pair
- Assume we have labels for both nodes and edges; if nodes or edges are not labeled, use generic symbol '#'

## Kernels Based on Contiguous Label Sequences

(cont'd)

- Let  $\mathcal{S}_n$  be set of all possible label sequences of walks with  $n$  edges and let  $\lambda$ ,  $\mathcal{W}_n(G)$ , and  $l_i(w)$  be as before
- Define mapping  $\Phi(G)$  to have one feature per possible label sequence  $s \in \bigcup_n \mathcal{S}_n$ :

$$[\Phi(G)]_s = \sqrt{\lambda_n} |\{w \in \mathcal{W}_n(G) : \forall i \ s_i = l_i(w)\}|$$

i.e. the number of walks in  $G$  with  $n$  edges whose (vertex and edge) label sequences match  $s = s_1, s_2, \dots, s_{2n+1} \in \mathcal{S}_n$ , weighted by  $\sqrt{\lambda_n}$

# Kernels Based on Contiguous Label Sequences

(cont'd)

- To compute the kernel, use the notion of a product graph: given  $G_1 = (\mathcal{V}_1, \mathcal{E}_1)$  and  $G_2 = (\mathcal{V}_2, \mathcal{E}_2)$ ,  $G_{\times} = G_1 \times G_2$  is defined as

$$\mathcal{V}_{\times} = \{(v_1, v_2) \in \mathcal{V}_1 \times \mathcal{V}_2 : \text{label}(v_1) = \text{label}(v_2)\}$$

$$\mathcal{E}_{\times} = \{((u_1, u_2), (v_1, v_2)) \in \mathcal{V}_{\times}^2 : (u_1, v_1) \in \mathcal{E}_1$$

$$\wedge (u_2, v_2) \in \mathcal{E}_2 \wedge \text{label}(u_1, v_1) = \text{label}(u_2, v_2)\}$$

- One can show that

$$|\{w \in \mathcal{W}_n(G_1 \times G_2) : \forall i \ s_i = l_i(w)\}|$$

$$= |\{w \in \mathcal{W}_n(G_1) : \forall i \ s_i = l_i(w)\}|$$

$$\cdot |\{w \in \mathcal{W}_n(G_2) : \forall i \ s_i = l_i(w)\}|$$

- Since an  $n$ -edge walk in  $G_1 \times G_2$  corresponds to a walk in each of  $G_1$  and  $G_2$ , each with same label sequence, the dot product  $\langle \Phi(G_1), \Phi(G_2) \rangle$  can be computed as

$$k_{\times}(G_1, G_2) = \sum_{i,j=1}^{\mathcal{V}_{\times}} \left[ \sum_{n=0}^{\infty} \lambda_n E_{\times}^n \right]_{ij}$$

**Topic summary due in 1 week!**