INTRODUCTION KERNELS FOR COMPLEX DATA TYPES MULTI-VIEW LEARNING STRUCTURED OUTPUT PREDICTION CONCLUSIO

# KERNEL METHODS IN MACHINE LEARNING LECTURE 9

# KERNEL METHODS WITH STRUCTURED AND MULTI-VIEW DATA

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### **O**VERVIEW

#### Introduction

KERNELS FOR COMPLEX DATA TYPES
String kernels
Graph kernels

LEARNING WITH MULTIPLE VIEWS

STRUCTURED OUTPUT PREDICTION

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# WHY KERNEL METHODS WHEN THERE IS DEEP LEARNING?

No universal solution to everything; these methods work for different problems.

#### Different complexities:

- ▶ Kernel methods depend on the  $n \times n$  kernel matrix; algorithms often  $\mathcal{O}(n^3)$ . Need to learn  $\mathcal{O}(n)$  parameters.
  - $\Rightarrow$  can tackle high-dimensional and complex features well, can easily tackle medium-size datasets  $n \approx 10^4$
- ▶ Deep networks generally scale  $\mathcal{O}(nd)$  with d the input dimension. Need to learn  $\mathcal{O}(d)$  parameters<sup>1</sup>
  - $\Rightarrow$  can tackle a lot of data, but the dimensionality is a restricting factor if it gets very high.

<sup>&</sup>lt;sup>1</sup>for dense neural networks

# WHY KERNEL METHODS WHEN THERE IS DEEP LEARNING?

Deep networks incorporate less prior knowledge, and use (and overfit) more data

#### Big data vs small data

- ► Automated data gathering from for example social networks; huge datasets ⇒ neural networks
- ► Expensive experimentations done by humans or with expensive machinery; complex, small datasets ⇒ perfect for kernel methods!
  - Life sciences, medicine, ...

## RECAP: KERNELS

RKHS  $\mathcal{H} \ni f: \mathcal{X} \to \mathbb{R}$ 

Kernel function  $k: \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ 

Kernel trick  $k(x,z) = \langle \phi(x), \phi(z) \rangle$ 

Feature map  $\phi: \mathcal{X} \to \mathcal{H}$ 

Representer theorem  $f(x) = \sum_{i} \alpha_{i} k(x, x_{i}), \quad \alpha_{i} \in \mathbb{R}$ 

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Representer theorem  $f(x) = \sum_{i} \alpha_i k(x, x_i), \quad \alpha_i \in \mathbb{R}$ 

 $\mathcal{X}$  doesn't have to be  $\mathbb{R}^d$ !

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#### KERNELS ON TEXT

Learning from natural language documents.



#### Basic concepts:

- ▶ Word: any sequence of basic alphabet surrounded by punctuation or spaces.
- ▶ **Document**: sequence of words
- ► Corpus: full set of documents
- ▶ **Dictionary**: set of terms occurring in corpus

#### BAG-OF-WORDS REPRESENTATION

#### BoW feature:

$$\phi_{BoW}(x) = [tf(t_1, x), tf(t_2, x), ..., tf(t_N, x)]^{\top}$$

In which

- $\triangleright$  x is a document
- $\triangleright$   $t_i$  is a term in the dictionary
- ▶  $tf(t_j, x)$  gives the **term frequency** of  $t_j$  in document x

#### BoW kernel:

$$k_{BoW}(x,z) = \langle \phi_{BoW}(x), \phi_{BoW}(z) \rangle$$

### BAG-OF-WORDS REPRESENTATION

Not very efficient, however there are things to do to improve:

- ► Tokenization: avoid computing over sparse high-dimensional feature vectors
- ▶ Stop word removal: exclude frequent but non-informative words
- ▶ Stemming: remove inflections and word form variation
- Normalization: reducing the effect of the length of documents;  $\phi(x) = \frac{\phi(x)}{\|\phi(x)\|}$

Doesn't capture all useful information:

- ▶ order of words "you are" vs "are you"
- ▶ grammatical information is lost (verb, noun..)

#### SEMANTIC KERNELS

Add semantic content by considering transformations

$$\phi_S(x) = \mathbf{S} \, \phi_{BoW}(x),$$

where the new features are now linear combinations of the old ones. (e.g. group related words, documents have non-zero similarity if they use terms from the same group.)

$$k_S(x,z) = \phi_{BoW}(x)^{\top} \mathbf{S}^{\top} \mathbf{S} \phi_{BoW}(z)$$

#### SEMANTIC KERNELS

$$k_S(x,z) = \phi_{BoW}(x)^{\top} \mathbf{S}^{\top} \mathbf{S} \phi_{BoW}(z)$$

Semantic kernel can be decomposed further: write S = RP;

- ▶ R is diagonal matrix containing term weights or relevances
  - ▶ Rare words might carry more meaning; inverse weighting
- ▶ P is proximity matrix defining semantic spread between the terms
  - ▶  $\mathbf{P}_{ij} > 0$  when term i is related to term j; e.g. "tree", "spruce" and "pine"
  - ▶ Load this from a database (e.g. WordNet), or analyse co-occurrences from large documents.

# STRING KERNELS<sup>2</sup>

Kernel on words/strings instead of collections of words.

Based on counting common subsequences; underlying feature map contains a feature for each possible substring.

- x AAACAAATAAGTAACTAATCTTTTAGGAAGAACGTTTCAACCATTTTGAG
- x' TACCTAATTATGAAATTAAATTTC $\overline{ ext{AGTGTGCTGATGGAAACGGAGAAGTC}}$

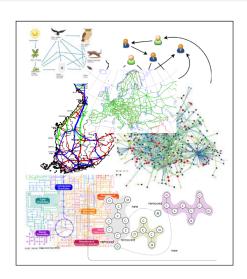
Multiple choices to tune the kernel:

- ▶ when do two subsequences match
- what kind of subsequences to consider
- ▶ how to weight matches

<sup>&</sup>lt;sup>2</sup>Shawe-Taylor & Cristianini, chapter 11

# Graphs are everywhere

Social networks, maps, communication channels, protein interaction networks, molecules, ...



#### Graphs are everywhere

Machine learning tasks on graphs:

- Given a graph, predict labels for its nodes
- Link prediction: given a set of nodes, predict which ones should be connected
- Graph classification
  - ▶ Drug discovery: given a candidate drug molecule (graph), predict if it will be active against a given type of cancer cell
  - ▶ Protein function prediction: given a 3D protein structure, predict its functional role

### GRAPH KERNELS

A broad idea: try to find common elements (labels, paths, subgraphs, ...) and count their occurrences

Subgraphs & neighbourhoods:

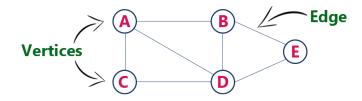
▶ Weisfeiler-Lehman kernel

#### Labels:

▶ Optimal assignment kernel

Paths / walks:

▶ Random walk kernel



# Weisfeiler-Lehman kernel

"How often the nodes' neighbourhoods match, up to depth d?"

$$k(G, G') = \sum_{i=0}^{d} \lambda_d \sum_{v \in V} \sum_{v' \in V'} \delta(relabel(v, i), relabel(v', i))$$

#### In which:

- ightharpoonup G, G' are the graphs to compare
- $\triangleright$  d is depth parameter
- $\triangleright$   $\lambda_d$  is parameter controlling importance of the level
- ▶  $\delta(\cdot, \cdot)$  returns 1 if its arguments are the same, 0 otherwise.
- ▶  $relabel(\cdot,i)$  returns a new label for the node based on it's neighbour's labels on level i-1: its own label & a sorted list of neighbouring node's labels
  - $\triangleright$  i = 0 just use original labels

#### Weisfeiler-Lehman Kernel

# Original labels i = 0



 $\Sigma = \{A, B\}$ 

#### Relabeled

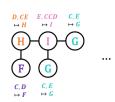
$$i = 1$$

 $\begin{array}{cccc}
 & D & \rightarrow E & \rightarrow C \\
\hline
D & E & \hline
C & C \\
A,B & A,B
\end{array}$ 

$$\Sigma = \{A, B, \mathbf{C}, \mathbf{D}, \mathbf{E}\}$$

#### Relabeled

$$i = 2$$



$$\Sigma = \{A, B, C, D, E, F, G, H, I\}$$

Image: Kriege, Johansson and Morris: A Survey on Graph Kernels, 2020

# OPTIMAL ASSIGNMENT(OA) KERNEL<sup>4</sup>

Find the best mapping between the nodes of two graphs.

$$k(G, G') = \max_{\pi \in \Pi_n} \sum_{i=1}^n \kappa(x_i, y_{\pi(i)})$$

In which:

- ▶ G, G' are the graphs to compare which have label sets  $X = \{x_1, x_2, ..., x_n\}$  and  $Y = \{y_1, y_2, ..., y_n\}^3$
- $\blacktriangleright$   $\pi$  is a permutation from set of all possible permutations of n elements,  $\Pi_n$
- $\triangleright$   $\kappa$  is a kernel on labels

 $<sup>^3{\</sup>rm If}$  one graph has less labels than the other, fill the it up with dummies and define similarity to a dummy be always 0.

<sup>&</sup>lt;sup>4</sup>Fröchlich et al, ICML 2005

# OPTIMAL ASSIGNMENT(OA) KERNEL

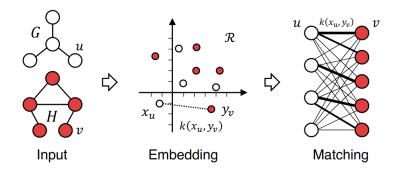


Image: Kriege, Johansson and Morris: A Survey on Graph Kernels, 2020

# RANDOM WALK KERNELS<sup>5</sup>

Count common walks (sequences of adjacent nodes) in two graphs

- ▶ In an unlabeled graph two walks match if they have the same length
- ▶ In a labeled graph also the node and edge labels need to match

Number of walks of length k can be computed with adjacency matrix A raised to the kth power

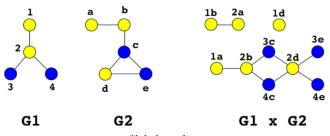
$$A^2 = \left( \begin{array}{c} 2 & 0 & 1 & 0 & 0 & 1 & 0 & 1 \\ 0 & 2 & 0 & 1 & 0 & 0 & 1 & 1 \\ 1 & 0 & 3 & 1 & 1 & 1 & 1 & 1 \\ 0 & 1 & 1 & 4 & 1 & 1 & 2 & 1 \\ 0 & 0 & 1 & 1 & 2 & 1 & 1 & 1 \\ 1 & 0 & 1 & 1 & 1 & 3 & 0 & 2 \\ 0 & 1 & 1 & 2 & 1 & 0 & 3 & 0 \\ 1 & 1 & 1 & 1 & 1 & 2 & 0 & 3 \end{array} \right)$$

<sup>&</sup>lt;sup>5</sup>Kashima et al., ICML 2003, Gärtner et al., COLT 2003

$$G_1 = (V_1, E_1)$$
 and  $G_2 = (V_2, E_2) \Rightarrow \text{product graph } G_{\times} = (V_{\times}, E_{\times});$ 

- ▶  $V_{\times} = \{(u, v) : u \in V_1, v \in V_2, label(u) = label(v)\}$ Make a new node for all pairs for which the labels match.
- $E_{\times} = \{((u_1, v_1), (u_2, v_2)) \in E_1 \times E_2 : (u_1, u_2) \in V_{\times}, \\ (v_1, v_2) \in V_{\times}, label((u_1, u_2)) = label((v_1, v_2))\}$

Two vertices in the direct product graph are adjacent iff the associated pairs of vertices are adjacent in original graphs.



Use the adjacency matrix,  $A_{\times}^{k}$ , to simultaneously trace common **labeled** walks in the two original graphs with  $G_{\times}!$ 

- ► Tracing a walk in the product graph corresponds to simultaneously tracing common walks in the two original graphs
- ▶ Ignore the labels in product graph; yet count only walks with matching labels.

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Random walk graph kernel: count all pairs of matching walks

$$k_{\times}(G_1, G_2) = \sum_{i,j=1}^{|V_{\times}|} \left[ \sum_{n=0}^{\infty} \lambda^n A_{\times}^n \right]_{ij}$$

- Considers walks of any length!  $[A_{\times}^n]_{ij}$  contains the number of walks of length n between the nodes i and j
- ▶  $0 \le \lambda \le 1$  is a decaying factor for the sum to converge
- ► Corresponds to an infinite-dimensional feature space

$$k_{\times}(G_1, G_2) = \sum_{i,j=1}^{|V_{\times}|} \left[ \sum_{n=0}^{\infty} \lambda^n A_{\times}^n \right]_{ij} = \sum_{i,j=1}^{|V_{\times}|} s_{ij}$$

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Geometric matrix series:

$$S = [s_{ij}]_{i,j=1}^{|V_{\times}|} = \sum_{n=0}^{\infty} \lambda^n A_{\times}^n = I_{|V_{\times}|} + \lambda A_{\times} + \lambda^2 A_{\times}^2 + \dots$$

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Multiply both sides with  $\lambda A_{\times}$ :

$$\lambda A_{\times}S = \lambda A_{\times} \sum_{\mathbf{n}=\mathbf{0}}^{\infty} \lambda^n A_{\times}^n$$

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$$\Rightarrow S = \left(I_{|V_{\times}|} - \lambda A_{\times}\right)^{-1}$$

$$\Rightarrow k_{\times}(G_{1}, G_{2}) = \sum_{i,j=1}^{|V_{\times}|} \left[ \left(I_{|V_{\times}|} - \lambda A_{\times}\right)^{-1} \right]_{ij}$$

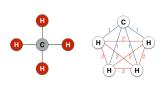
#### SHORTEST PATH KERNELS

RW kernels are not very efficient  $(\mathcal{O}(n^6))$  and suffer from tottering  $\Rightarrow$  from walks to paths





Shortest path kernels<sup>6</sup> much more efficient;  $\mathcal{O}(n^4)$ . (Uses Floyd-Warshall to find the paths).



Transform input graph to shortest-path graph.

Kernel is obtained by comparing node and edge labels in the new graph.

<sup>&</sup>lt;sup>6</sup>Borgwardt and Kriegel, 2005

## **O**VERVIEW

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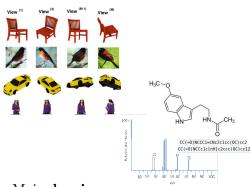
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#### LEARNING WITH MULTIPLE VIEWS

STRUCTURED OUTPUT PREDICTION

#### Multi-view data





Moi **bonjour** hello Guten Tag

#### Integrating data sources

#### Early fusion:

- ► Combine the data sources first (human expert, simple vector concatenation..)
- Learn a single model from combined data

#### Intermediate fusion:

- ► Input all data sources separately
- ▶ Learn how to combine the sources during learning the model
  - ► Multiple Kernel Learning

#### Late fusion:

- Learn a model for each data source independently
- ► Combine the results after learning e.g. majority voting

## Multiple Kernel Learning

Simple input data concatenation in feature spaces:

$$k(x,z) = \sum_{v=1}^{V} k^{v}(x,z)$$

Multiple Kernel Learning (MKL):

$$k(x,z) = \sum_{v=1}^{V} \alpha_v k^v(x,z),$$

in which the weights  $\alpha_v > 0$  are learnt.

#### Multiple Kernel Learning

How to learn the combination weights?

- ► Two-step approach with kernel alignment, where weights are learned before the predictive model
- ▶ One-step approaches: learn the weights jointly with the learning problem
  - Note: usually iterates over learning the kernel weights and learning the decision function

# MKL WITH KERNEL ALIGNMENT<sup>7</sup>

Kernel alignment:

$$A(\mathbf{K}, \mathbf{G}) = \frac{\langle \mathbf{K}_c, \mathbf{G}_c \rangle_F}{\|\mathbf{K}_c\|_F \|\mathbf{G}_c\|_F}$$

in which 
$$\mathbf{K}_c = \left(\mathbf{I} - \frac{1}{n} \mathbb{1} \mathbb{1}^{\top}\right) \mathbf{K} \left(\mathbf{I} - \frac{1}{n} \mathbb{1} \mathbb{1}^{\top}\right)$$

Ideal (target) kernel for binary classification:  $\mathbf{y}\mathbf{y}^{\top}$  with  $y_i = -1.1.$ 

A good kernel **K** for binary classification has large  $A(\mathbf{K}, \mathbf{y}\mathbf{y}^{\top})$ .

<sup>&</sup>lt;sup>7</sup>Cortes et al 2012

### MKL WITH KERNEL ALIGNMENT

ALIGN assigns the weights in MKL kernel to be the scores  $A(\mathbf{K}^v, \mathbf{y}\mathbf{y}^\top)$ :

$$k(x,z) = \sum_{v=1}^{V} \alpha_v k^v(x,z) = \sum_{v=1}^{V} A(\mathbf{K}^v, \mathbf{y} \mathbf{y}^\top) k^v(x,z),$$

ALIGNF problem:

$$\max_{\mathbf{d}} \frac{\left\langle \sum_{v=1}^{V} d_v \mathbf{K}^v, \mathbf{y} \mathbf{y}^\top \right\rangle_F}{\left\| \sum_{v=1}^{V} d_v \mathbf{K}^v \right\|_F \|\mathbf{y} \mathbf{y}^\top \|_F}, \quad s.t \|\mathbf{d}\|_2 = 1, d_i \ge 0$$

solved as a quadratic programming problem.

# SIMPLEMKL<sup>8</sup>

- ▶ One-step approach: learn the model (such as SVM classifier) simultaneously with the kernel weights.
- ► Learns a linear combination

$$k_d(x_i, x_j) = \sum_{m=1}^{P} d_m k_m(x_i, x_j)$$

where kernel weights are constrained to convex combination

$$\sum_{m} d_m = 1, d_m \ge 0$$

▶ Bi-level optimization scheme with SVM solver as a wrapper

<sup>&</sup>lt;sup>8</sup>Rakotomamonjy et al 2008

#### SIMPLEMKL: THE IDEA

- Can be interpreted as learning a mixture of classifiers
- ▶ One base classifier for each kernel, with kernel weight  $d_m$ , example weights  $\alpha_i$ :

$$f_m(\mathbf{x}) = \langle \mathbf{w}_m, \phi_m(\mathbf{x}) \rangle = \sum_i \alpha_i y_i d_m \kappa_m(\mathbf{x}, \mathbf{x}_i)$$

- ▶ Above  $\mathbf{w}_m = \sum_i \alpha_i y_i d_m \phi_m(\mathbf{x}_i)$  obtained from Lagrangian duality
- ▶ The full model will use a mixture of base classifiers:

$$f(\mathbf{x}) = \sum_{m=1}^{P} f_m(\mathbf{x}) = \sum_{i} \alpha_i y_i d_m \kappa_m(\mathbf{x}, \mathbf{x}_i)$$

## SIMPLEMKL: PRIMAL PROBLEM

$$\min_{\mathbf{w},b,\xi,d} \frac{1}{2} \sum_{m} \frac{1}{d_{m}} \|\mathbf{w}_{m}\|^{2} + C \sum_{i} \xi_{i}$$

$$s.t. \ y_{i} \left( \sum_{m} \langle \mathbf{w}_{m}, \phi_{m}(\mathbf{x}_{i}) \rangle \ (+b) \right) \geq 1 - \xi_{i}, i = 1 \dots, \ell$$

$$\xi_{i} \geq 0, i = 1 \dots, \ell$$

$$\sum_{m} d_{m} = 1, d_{m} \geq 0, m = 1, \dots, P$$

- Objective minimizes a linear combination of norms of weight vectors corresponding to different kernels plus slack for examples
- ► Constraints declare that the mixture classifier should achieve a large margin
- ► It is a convex problem (Rakotomamonjy, 2008)

# SIMPLEMKL: BI-LEVEL OPTIMIZATION, PRIMAL

Reformulation as a bi-level optimization problem

$$\min_{\boldsymbol{d} \in \mathcal{H}} J(\boldsymbol{d})$$

$$s.t. J(\boldsymbol{d}) = \begin{cases} \min_{\mathbf{w}, b, \xi} & \frac{1}{2} \sum_{m} \frac{1}{d_{m}} \|\mathbf{w}_{m}\|^{2} + C \sum_{i} \xi_{i} \\ s.t. & y_{i} \left( \sum_{m} \langle \mathbf{w}_{m}, \phi_{m}(\mathbf{x}_{i}) \rangle \right. (+b) \right) \geq 1 - \xi_{i} \\ & \xi_{i} \geq 0, i = 1 \dots, \ell \end{cases}$$

- ▶ In outer loop optimize kernel weights d
- ► Inner loop corresponds to a primal soft-margin SVM using a combined kernel with current kernel weights

## SIMPLEMKL: BI-LEVEL OPTIMIZATION, DUAL

▶ We can plug in the dual of the SVM problem in the inner loop

$$\min_{\boldsymbol{d} \in \mathcal{H}} J(\boldsymbol{d})$$

$$s.t. \ J(\boldsymbol{d}) = \begin{cases} \max_{\alpha} & \sum_{i} \alpha_{i} - \frac{1}{2} \sum_{i,j} \alpha_{i} \alpha_{j} \sum_{m} d_{m} \kappa_{m}(\mathbf{x}_{i}, \mathbf{x}_{j}) \\ s.t. & 0 \leq \alpha_{i} \leq C \\ & (\sum_{i} \alpha_{i} y_{i} = 0) \end{cases}$$

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It's easy to predict functions  $f: \mathcal{X} \to \mathbb{R}$  with kernel methods;  $f \in \mathcal{H}$ .

By extension also vector-valued functions  $f: \mathcal{X} \to \mathbb{R}^p$  are easy.

We have seen that it's perfectly fine for  $\mathcal{X}$  to be structured data and not just subset of  $\mathbb{R}^d$ .

What if output space  $\mathcal{Y}$  in  $f: \mathcal{X} \to \mathcal{Y}$  is structured?

### STRUCTURED OUTPUT PREDICTION

- Want to find f
- $\triangleright$  Know how to map  $\mathcal{Y}$  to RKHS  $\mathcal{F}_{u}$  with  $\phi_{u}$
- $\rightarrow$  Learn  $h: \mathcal{X} \rightarrow \mathcal{F}_{\eta}$
- $\rightarrow$  In general q cannot be found explicitly: search the set of  $\mathcal{Y}$ for element that would give the predicted value in  $\mathcal{F}_{v}$ ;

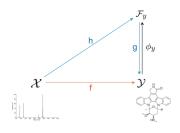


Image: Brouard et al: Fast metabolite identification with Input Output Kernel Regression, 2016

### CONCLUSION

- ► Kernel methods are useful for many real-world learning problems that neural networks are not well applicable to.
- ► Kernels can handle structured inputs; kernels for text, graphs..
- ► Kernels are a natural choice for incorporating multiple views in a learning problem.
- ► Kernels can also be used in prediction problems when target of a learning problem is not vectorial, but structured.