

Advanced Machine Learning  
Lecture 2  
Learning with kernel methods

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Fall 2015

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# Learning in RKHS

Motivation : predict the property of a molecule



- **Inputs** : molecule (drug candidate)
- **Output** : activity on a cancer line (or several cancer lines)

A regression problem from structured data.

# Learning in RKHS

## A to-do do list

- 1 Define a PDS kernel :  $k(\cdot, \cdot)$
- 2 Define a (unique) RKHS,  $\mathcal{H}$  from  $k$  with an appropriate norm  $\|\cdot\|_{\mathcal{H}}$
- 3 Define a loss functional with two terms : a local loss function  $\ell$  and a penalty function  $\Omega$
- 4 Prove/use a representer theorem to get the form of the minimizer of this functional :  $\sum_i \alpha_i h(\cdot, x_i)$  **We learn this today**
- 5 Solve the optimization problem with this minimizer **we do that for kernel ridge regression today**

# Learning in RKHS

To solve the molecular property pb

- $S_n = \{(x_1, y_1), \dots, (x_n, y_n)\}$
- Assume we have defined a kernel over labeled graphs  $k$  and its associated RKHS  $\mathcal{F}$
- Choose and minimize the loss function :
  - $\arg \min_{f \in \mathcal{F}} \frac{1}{2} \sum_{i=1}^n \|y_i - f(x_i)\|^2 + \lambda \|f\|_{\mathcal{F}}^2$

### Theorem

Let  $k : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R}$  be a positive definite symmetric kernel and  $\mathcal{H}_k$ , its corresponding RKHS, then, for any non-decreasing function  $\Omega : \mathbb{R} \rightarrow \mathbb{R}$  and any loss function  $L : \mathbb{R}^n \rightarrow \mathbb{R} \cup \{+\infty\}$ , any minimizer of :

$$J(f) = L(f(x_1), \dots, f(x_n)) + \lambda \Omega(\|f\|_{\mathcal{H}}^2) \quad (1)$$

admits an expansion of the form :

$$f^*(\cdot) = \sum_{i=1}^n \alpha_i k(x_i, \cdot).$$

Moreover if  $\Omega$  is strictly increasing, then any minimizer of 1 has exactly this form.

# Learning in RKHS

## Proof of the Representer theorem

Let us define :  $\mathcal{H}_1 = \text{span} \{k(x_i, \cdot), i = 1, \dots, n\}$

Any  $f \in \mathcal{H}$  writes as :  $f = f_1 + f^\perp$ , with  $f_1 \in \mathcal{H}_1$  and  $f^\perp \in \mathcal{H}_1^\perp$   
where  $\mathcal{H}$  = direct sum of  $\mathcal{H}_1$  and  $\mathcal{H}_1^\perp$ .

By orthogonality,  $\|f\|^2 = \|f_1\|^2 + \|f_1^\perp\|^2$

Hence, by property of  $\Omega$ ,

$$\Omega(\|f\|^2) = \Omega(\|f_1\|^2) + \Omega(\|f_1^\perp\|^2) \geq \Omega(\|f_1\|^2)$$

By the reproducing property, we get :

$$f(x_i) = \langle f_1(\cdot) + f_1^\perp(\cdot), k(x_i, \cdot) \rangle = \langle f_1(\cdot), k(x_i, \cdot) \rangle = f_1(x_i)$$

Hence,  $L(f(x_1), \dots, f(x_n)) = L(f_1(x_1), \dots, f_1(x_n))$  and

$$J(f) \leq J(f_1)$$

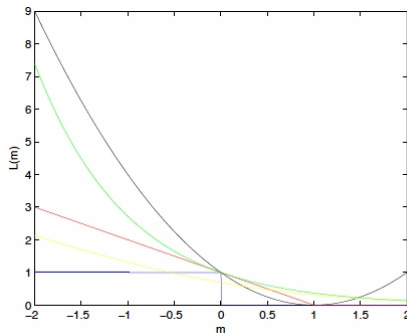
To recap, if  $f$  is a minimizer of  $J(f)$ , then  $f_1$  is also a minimizer of  $J$ . Moreover if  $\Omega$  is strictly increasing,  $J(f_1) < J(f)$ , then any  $f = f_1 + f_1^\perp$  exactly equals to  $f_1$ .

- $L(f(x_1), \dots, f(x_n)) = \sum_i (y_i - h(x_i))^2$  and  $\Omega(\|f\|) = \|f\|^2$ 
  - Kernel ridge regression :  $\hat{\alpha} = (K + \lambda Id)^{-1} \mathbf{y}$  (proved during course)
- SVM without bias  $b$
- $L(f(x_1), \dots, f(x_n)) = \max(0, 1 - y_i f(x_i))$  (hinge loss) and  $\Omega(\|f\|) = \|f\|^2$ 
  - If you want to introduce  $b$ , you need to refer to the semi-parametric representer theorem.



# Learning in RKHS

## Losses



Various convexifications of the 0 – 1 loss, including the hinge loss.

# Learning in RKHS

Solving our problem with labeled graphs (molecule) in the input space

A molecule  $x$  is seen as a labeled graph defined on an alphabet  $\mathcal{A}$ , a finite set of symbols.

For a given length  $L$ , let us first enumerate all the paths of length  $\ell \leq L$  in the training dataset (data are molecule = labeled graphs). Let  $m$  be the size of this set ( $m$  is the sum of the number of all possible combinations of  $\ell \leq L$  symbols of  $\mathcal{A}$ ). Let us build a dictionary of possible paths of length less or equal to  $L$ .

For a graph  $x$ , define the feature map :

$\varphi(x) = (\varphi_1(x), \dots, \varphi_m(x))^T$  where  $\varphi_i(x)$  is 1 if the  $i^{th}$  path appears in the labeled graph  $G$ , and 0 otherwise. **Definition 1 :**

$$k_L(x, x') = \langle \varphi(x), \varphi(x') \rangle$$

**Tanimoto kernel**

$$k_L^t(x, x') = \frac{k_L(x, x')}{k_L(x, x) + k_L(x', x') - k_L(x, x')}$$

**idea :**  $k_L^t$  calculates the ratio between the number of elements of the intersection of the two sets of paths ( $G$  and  $G'$  are seen as bags of paths) and the number of elements of the union of the two sets.

**Reference :** Ralaivola et al. 2005, Su et al. 2011

# Learning in RKHS

## Solution for Kernel Ridge regression

Let  $K_L = (k_L(x_i, x_{jij}))$  be the Gram matrix of our training molecules  $\{x_1, \dots, x_n\}$ . Let  $\mathbf{y}$  the vector of  $n$  dimensions with the target values. Using the representer theorem for the ridge loss, and using the proof on the blackboard we get :

$$\hat{\alpha} = (K + \lambda Id)^{-1} \mathbf{y}$$

We are now able to predict for a new molecule its score with respect with a given cancer (cell line) by using :

$$f(\cdot) = \sum_{i=1}^n \hat{\alpha}_i k_L(x, x_i)$$

- 1 Learning in RKHS
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- Use closure properties to build new kernels from existing ones
- Kernels can be defined for various objects :
  - **Structured objects** : (sets), graphs, trees, sequences, ...
  - Unstructured data with underlying structure : texts, images, documents, signal, biological objects (gene, mRNA,protein, ...)
- **Kernel learning** :
  - Hyperparameter learning : see Chapelle et al. 2002
  - Multiple Kernel Learning : given  $k_1, \dots, k_m$ , learn a convex combination  $\sum_i \beta_i k_i$  of kernels (see SimpleMKL Rakotomamonjy et al. 2008, unifying view in Kloft et al. 2010)

# Kernel design

Kernel closure properties : a few rules

- The sum of two kernels is a kernel
- The product of two kernels is a kernel
- $k(x, x') = \exp(-\gamma D(x, x')^2)$  is a kernel if  $D : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R}^+$  is a distance.

Exercise : prove it....

# Kernel design

## Examples of kernels for non-vectorial data and applications

- Convolution kernels
- Graph kernel
- Fisher kernels
- Sequence kernels



### *Definition :*

Suppose that  $x \in \mathcal{X}$  is a **composite structure** and  $x_1, \dots, x_D$  are its "parts" according a relation  $R$  such that  $(R(x, x_1, x_2, \dots, x_D))$  is true, with  $x_d \in \mathcal{X}_d$  for each  $1 \leq d \leq D$ ,  $D$  being a positive integer.  $k_d$  be a PDS kernel on a set  $\mathcal{X} \times \mathcal{X}$ , for all  $(x, x')$ , we define :

$$k_{conv}(x, x') = \sum_{(x_1, \dots, x_d) \in R^{-1}(x), (x'_1, \dots, x'_d) \in R^{-1}(x')} \prod_{d=1}^D k_d(x_d, x'_d)$$

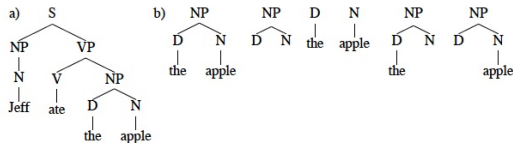
$R^{-1}(x)$  = all decompositions  $(x_1, \dots, x_D)$  such that  $(R(x, x_1, x_2, \dots, x_D))$ .  $k_{conv}$  is a PDS kernel as well. Intuitive kernel, used as a building principle for a lot of other kernels. Next, we will see two examples.

### Learning task :

- **Input** : sentence  $\rightarrow$  syntax tree
- **Output** : question class
- For instance, in economical news articles, classes are ORGANIZATION, LOCATION,

# Kernel design

## Kernel for Natural Language Processing



Let us first enumerate all tree fragments that occur in the training data. Let  $m$  be the size of this set. For a tree, define  $\varphi(T) = (\varphi_1(T), \dots, \varphi_m(T))^T$  where  $\varphi_i(T)$  is the number of occurrences of the  $i^{th}$  subtree.

### Definition :

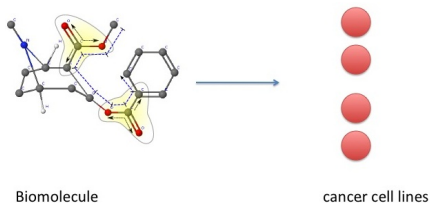
$$k_{conv}(T, T') = k(\varphi(T), \varphi(T'))$$

NB : the kernel can be normalized. In NLP,  $k$  is often chosen as the linear kernel. Efficient implementations are available.

Sequences can be processed in the same way.

References : Collins and Duffy, 2001 ; Suzuki et al. 2003

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### Kernel for labeled graphs

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## Kernel design

### Kernel between vertices in a graph

Let  $x_1, \dots, x_n$ ,  $n$  objects associated with a non oriented graph of size  $n$  and adjacency matrix  $W$ . Define the graph Laplacian :  
 $L = D - W$ ,  $D$  is the diagonal matrix of degrees

$$K = \exp(-\lambda L)$$

We will see applications of this kernel in the unsupervised course.

**Reference : Kondor and Lafferty, 2003**



Combine the advantages of graphical models and discriminative methods

Let  $\mathbf{x} \in \mathbb{R}^p$  be the input vector of a classifier.

- Learn a generative model  $p_\theta(\mathbf{x})$  from unlabeled data  $\mathbf{x}_1, \dots, \mathbf{x}_n$
- Define the Fisher vector as :  $\mathbf{u}_\theta(\mathbf{x}) = \nabla_\theta \log p_\theta(\mathbf{x})$
- Estimate the Fisher Information matrix of  $p_\theta$  :  
$$F_\theta = \mathbb{E}_{\mathbf{x} \sim p_\theta} [\mathbf{u}_\theta(\mathbf{x}) \mathbf{u}_\theta(\mathbf{x})^T]$$
- **Definition** :  $k_{Fisher}(\mathbf{x}, \mathbf{x}') = \mathbf{u}_\theta(\mathbf{x})^T F_\theta \mathbf{u}_\theta(\mathbf{x})$

### Applications

Classification of secondary structure of proteins, topic modeling in documents, image classification and object recognition, audio signal classification ...

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We would like to use an appropriate theorem for learning classification function of the following form :

$$f(x) = \sum_i \alpha_i k(x, x_i) + b \quad (2)$$

whose sign will give the class. For that purpose, the RKHS associated to a given PDS kernel  $k$  is not exactly the functional space we want to work in. We would like to learn two functions :  $f \in$  the RKHS  $\mathcal{F}_k$  and  $g$  a constant function such that  $f(x) = h(x) + g(x)$

- Write a Representer theorem for semi-parametric models for models of the form  $f(\cdot) + \sum_{\beta j} \psi_j(\cdot)$  where the family of functions  $(\psi_j(\cdot))_{j=1, \dots, m}$  is given.
- Apply it with the hinge loss :  
 $L(f(x_1), \dots, f(x_n)) = \sum_{i=1}^n \max(0, 1 - y_i f(x_i))$  and  
 $\Omega(z) = z^2$

- Prove that  $k(x, y) = \exp(-\gamma D(x, y)^2)$  is a kernel for  $D$  a distance function over a set  $\mathcal{X} \times \mathcal{X}$
- Prove other simple rules of construction of kernels (product, sum, ...)

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