Recitation 5 Kernels

DS-GA 1003 Machine Learning

CDS

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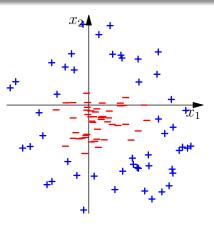
Outline

- Motivation
- Kernel
- Incorporating kernels into ridge regression and SVM
- RBF Kernel
- IPython Demo: MNIST with RBF kernel

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Question

Consider applying SVM to the data set. What is the issue?



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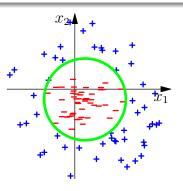
Consider applying SVM to the data set. What is the issue?

Solution

We want to allow for non-linear regression functions, but we would like to reuse the same fitting procedures we have already developed. To do this we will expand our feature set by adding non-linear functions of old features.

Solution

For the SVM we expand our feature vector from $(1, x_1, x_2)$ to $(1, x_1, x_2, x_1x_2, x_1^2, x_2^2)$. Using w = (-1.875, 2.5, -2.5, 0, 1, 1) gives $-1.875 + 2.5x_1 - 2.5x_2 + x_1^2 + x_2^2 = (x_1 + 1.25)^2 + (x_2 - 1.25)^2 - 5 = 0$ as our decision boundary.



- Linear model is clearly insufficient to represent these problems.
- The most intuitive solution is to expand the input space
 - Adding features
- We can define a **feature map function** $\varphi(x) : \mathcal{X} \mapsto \mathcal{H}$
 - $dim(\mathcal{H}) > dim(\mathcal{X})$
 - For SVM example above, $\varphi(1, x_1, x_2) = [1, x_1, x_2, x_1x_2, x_1^2, x_2^2]$.
- We then find a linear separator on the feature space \mathcal{H} .

Adding Features

- Polynomials can approximate any function (Taylor's Theorem).
- We can linearly model any problem perfectly if we add enough terms.
- But adding features obviously comes with a cost.
- The cost grows exponentially as we increase the degree.

Adding Features

Question

Suppose we begin with *d*-dimensional inputs $x = (x_1, \ldots, x_d)$. We add all features up to degree *M*. More precisely, all terms of the form

$$x_1^{p_1}\cdots x_d^{p_d}$$
 $p_i\geq 0$ and $p_1+\cdots+p_d\leq M$

How many features will we have in total?

• There will be $\binom{M+d}{M}$ terms total. Grows very quickly!

• For example, if d = 40 and M = 8 we get $\binom{40+8}{8} = 377348994$.

- Both *M* and *d* impacts the cost of adding features.
- If we stick with polynomial features up to order M, it's takes exponential time $O(d^M)$ to compute all features.
- How do we make the computation feasible?

Representer Theorem

Theorem

Suppose you have a loss function of the form

$$J(w) = L(w^{T}\varphi(x_1), \ldots, w^{T}\varphi(x_n)) + R(||w||_2)$$

where

•
$$x_i \in \mathbb{R}^d, w \in \mathbb{R}^{d'}, \varphi(x) : \mathbb{R}^d \mapsto \mathbb{R}^{d'}.$$

•
$$L: \mathbb{R}^n \to \mathbb{R}$$
 is an arbitrary function (loss term).

• $R : \mathbb{R}_{\geq 0} \to \mathbb{R}$ is increasing (regularization term).

Assume J has at least one minimizer. Then J has a minimizer w^* of the form $w^* = \sum_{i=1}^n \alpha_i \varphi(x_i)$ for some $\alpha \in \mathbb{R}^n$. If R is strictly increasing, then all minimizers have this form.

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Representer Theorem: Proof

Proof.

- Let $w^* \in \mathbb{R}^{d'}$ and let $S = \text{Span}(\varphi(x_1), \dots, \varphi(x_n)).$
- Suppose w^* is the optimal parameter, and it **does not lie in** S.
- Then we can write w^{*} = u + v where u ∈ S and v ∈ S[⊥]. (Here u is the orthogonal projection of w^{*} onto S, and S[⊥] is the subspace of all vectors orthogonal to S.)
- Then $(w^*)^T \varphi(x_i) = (u + v)^T \varphi(x_i) = u^T \varphi(x_i) + v^T \varphi(x_i) = u^T \varphi(x_i)$. So the prediction only depends on $u^T \varphi(x_i)$.
- But $||w^*||_2^2 = ||u+v||_2^2 = ||u||_2^2 + ||v||_2^2 + 2u^T v = ||u||_2^2 + ||v||_2^2 \ge ||u||_2^2$.
- Thus $R(||w^*||_2) \ge R(||u||_2)$ showing $J(w^*) \ge J(u)$.

Representer Theorem

- If your loss function only depends on w via its inner products with the inputs, and the regularization is an increasing function of the ℓ_2 norm, then we can write w^* as a linear combination of the training data.
- This applies to ridge regression and SVM.

Representer Theorem: Ridge Regression

• By adding features to ridge regression we had

$$\begin{split} J(\tilde{w}) &= \frac{1}{n} \sum_{i=1}^{n} (\tilde{w}^{T} \varphi(x_{i}) - y_{i})^{2} + \lambda \|\tilde{w}\|_{2}^{2} \\ &= \frac{1}{n} \|\Phi \tilde{w} - y\|_{2}^{2} + \lambda \tilde{w}^{T} \tilde{w}, \end{split}$$

where $\Phi \in \mathbb{R}^{n \times d'}$ is the matrix with $\varphi(x_i)^T$ as its *i*th row.

Representer Theorem applies giving w̃ = ∑_{j=1}ⁿ α_jφ(x_j) = Φ^Tα.
Plugging in gives

$$J(\alpha) = \frac{1}{n} \left\| \Phi \Phi^{T} \alpha - y \right\|_{2}^{2} + \lambda \alpha^{T} \Phi \Phi^{T} \alpha.$$

• Define $K = \Phi \Phi^T$

Representer Theorem: Dual SVM

• The dual SVM problem (with features) is given by

$$\begin{aligned} \text{maximize}_{\alpha} & \sum_{i=1}^{n} \alpha_{i} - \frac{1}{2} \sum_{i,j=1}^{n} \alpha_{i} \alpha_{j} y_{i} y_{j} \varphi(x_{i})^{T} \varphi(x_{j}) \\ \text{subject to} & \sum_{i=1}^{n} \alpha_{i} y_{i} = 0 \\ & \alpha_{i} \in \left[0, \frac{c}{n}\right] \quad \text{for } i = 1, \dots, n. \end{aligned}$$

- We can immediately kernelize (no representer theorem needed) by replacing $\varphi(x_i)^T \varphi(x_j) = k(x_i, x_j)$.
- Recall that we were able to derive the conclusion of the representer theorem using strong duality for SVMs.

The Kernel Function

Definition (Kernel)

Given a feautre map $\varphi(x) : \mathcal{X} \mapsto \mathcal{Z}$, the **kernel function** corresponding to $\varphi(x)$ is

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

where $\langle \cdot, \cdot \rangle$ is an inner product operator.

- So a kernel function computes the inner product of applying the feature map φ(x) for two inputs x, x' ∈ X.
- We only need to know the output of the kernel to find the parameters.
- Predictor function is:

$$f(x^*) = \sum_i \alpha_i k(x_i, x^*)$$

Efficiency of Kernel

Consider the polynomial kernel $k(x, y) = \langle \varphi(x), \varphi(y) \rangle = (1 + x^T y)^M$ where $x, y \in \mathbb{R}^d$. For example, if M = 2 we have

$$(1 + x^T y)^2 = 1 + 2x^T y + x^T y x^T y = 1 + 2 \sum_{i=1}^d x_i y_i + \sum_{i,j=1}^d x_i y_i x_j y_j.$$

Option 1: First explicitly evaluate $\varphi(x)$ and $\varphi(y)$, and then compute $\langle \varphi(x), \varphi(y) \rangle$.

- $\varphi(x) = (1, \sqrt{2}x_1, \dots, \sqrt{2}x_d, x_1^2, \dots, x_d^2, \sqrt{2}x_1x_2, \sqrt{2}x_1x_3, \dots, \sqrt{2}x_{d-1}x_d)$
- Takes $O(d^M)$ times to evaluate $\varphi(x)$ and $\varphi(y)$.
- Takes another $O(d^M)$ times to compute the inner product.
- Time complexity is $O(d^M)$.

Efficiency of Kernel

Consider the polynomial kernel $k(x, y) = \langle \varphi(x), \varphi(y) \rangle = (1 + x^T y)^M$ where $x, y \in \mathbb{R}^d$. This computes the inner product of all monomials up to degree M in time O(d). For example, if M = 2 we have

$$(1 + x^T y)^2 = 1 + 2x^T y + x^T y x^T y = 1 + 2 \sum_{i=1}^d x_i y_i + \sum_{i,j=1}^d x_i y_i x_j y_j.$$

Option 2: First calculate $1 + x^T y$, then calculate $(1 + x^T y)^M$.

- Takes O(d) time to evaluate $1 + x^T y$.
- Takes O(1) time to calculate $(1 + x^T y)^M$
- Time complexity is O(d)

Recap on what we achieved

- Start with a low dimensional model
 - Due to limited input data size
 - Number of parameters is d
- Want to increase the model capacity by adding features $x_i \rightarrow \varphi(x_i)$
 - The cost is too high as we increase degrees
 - Number of parameters is d', d' >> d
- Realize the optimal parameter is a linear combination of $\varphi(x_i)$
 - Representer Theorem
 - Number of parameters becomes N, d' >> N > d
- Realize we only need the inner product of two $\varphi(x_i)$, $k(\cdot, \cdot)$
 - There are more efficient methods to compute the inner product
 - We don't need to explicitly compute $\varphi(\cdot)$
- The rephrased problem becomes a linear problem
 - But the solution still has high dimensional expressive power!

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Mercer's Theorem

- Not all function f(x, y) are valid kernels.
- How can we know if k(x, y) is a valid kernel or not?

Theorem (Mercer's Theorem)

Fix a kernel $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$. There is a Hilbert space H and a feature map $\varphi : \mathcal{X} \to H$ such that $k(x, y) = \langle \varphi(x), \varphi(y) \rangle_H$ if and only if for any $x_1, \ldots, x_n \in \mathcal{X}$ the associated matrix K is positive semi-definite:

$$\mathcal{K} = \begin{pmatrix} k(x_1, x_1) & \cdots & k(x_1, x_n) \\ \vdots & \ddots & \vdots \\ k(x_n, x_1) & \cdots & k(x_n, x_n) \end{pmatrix}$$

Such a kernel k is called **positive semi-definite**.

Positive Semi-Definite

Definition (Positive Semi-Definite)

A matrix $A \in \mathbb{R}^{n \times n}$ is **positive semi-definite** if it is symmetric and

$$x^T A x \ge 0$$

for all $x \in \mathbb{R}^n$.

• Equivalent to saying the matrix is symmetric with non-negative eigenvalues.

Kernel Examples

- Dot Product
 - $k(x_i, x_j) = x_i^T x_j$
- Mth Polynomial Kernels
 - $k(x_i, x_j) = (1 + x_i^T x_j)^M$
- RBF Kernels

•
$$k(x_i, x_j) = exp(-\frac{||x_i - x_j||^2}{2\sigma^2})$$

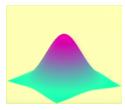
- Sigmoid kernel
 - $k(x_i, x_j) = tanh(\alpha x_i^T x_j + c)$

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RBF Kernels

$$k(w,x) = \exp\left(-\frac{\|w-x\|_2^2}{2\sigma^2}\right).$$

- 2d RBF kernel looks like the following.
- Let's say we fix w. The k(w, x) is high when x is very close to w. The value decays as x is moving away from w.
- σ controls the spread of the kernel. The higher σ is the wider / flatter the landscape is for k(w, x).



RBF Kernels

- As we saw earlier for ridge regression and SVM classification, the decision function has the form $f_{\alpha}(x) = \sum_{i=1}^{n} \alpha_i k(x_i, x)$.
- For ridge regression, this means that using the RBF kernel amounts to approximating our data by a linear combination of Gaussian bumps.
- For SVM classification, each $k(x_i, x) = \exp(-||x_i x||_2^2/(2\sigma^2))$ represents a exponentially decaying distance between x_i and x. Thus our decisions depend on our proximities to data points.

Going to infinite dimension

- What is the polynomial expression of $\varphi(\cdot)$ for RBF and Sigmoid Kernel?
 - There are no finite expression, they are sum of infinite polynomials

•
$$\varphi(x) = e^{-x^2/2\sigma^2} \left[1, \sqrt{\frac{1}{1!\sigma^2}} x, \sqrt{\frac{1}{2!\sigma^4}} x^2, \sqrt{\frac{1}{3!\sigma^6}} x^3, \ldots \right]$$

- This implies we have essentially modeled the problem using a infinite degree polynomial!
- At this point, the factor limiting our model capacity is the amount of training data.

Finding Your Own Kernels

Let $k_1, k_2 : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ be positive semi-definite kernels. Then so are the following:

- $k_3(w,x) = k_1(w,x) + k_2(w,x)$
- $k_4(w,x) = \alpha k_1(w,x)$ for $\alpha \ge 0$
- $k_5(w,x) = f(w)f(x)$ for any function $f : \mathcal{X} \to \mathbb{R}$
- $k_6(w, x) = k_1(w, x)k_2(w, x)$

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Remarks

- With Representer Theorem, we can re-parameterize our prediction function from $f_w(x) = w^T \varphi(x)$ to $f_\alpha(x) = \sum_{i=1}^n \alpha_i k(x_i, x)$.
- The feature representation φ(x) only appears in inner product form in both the loss function and the prediction function.
- Therefore, we just need to evaluate the kernel function k(x, y) and never need to explicitly evaluate φ(x). It's much easier to compute the kernel k(x, y) than the inner product.
- The kernel k(x, y), to some extent, represents a similarity score between two data points.
- We are almost guaranteed to overfit on training data, so regularization is very important.

References

• DS-GA 1003 Machine Learning Spring 2021

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