TBSI Seminar Summer 2020

### Seminar in Data Science

# Lecture 8: Kernel Methods; Data Normalization

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A linear regression problem

Linear (second-order) auto-regressive model for time-series: y<sub>t</sub> linear function of  $y_{t-1}, y_{t-2}$ 

$$y_t = w_1 + w_2 y_{t-1} + w_3 y_{t-2}, t = 3, ..., T.$$

This writes  $y_t = w^T x_t$ , with  $x_t$  the "feature vectors"

$$x_t := (1, y_{t-1}, y_{t-2}), t = 3, ..., T.$$

Define  $3 \times (T-2)$  matrix X with t-th column  $x_t$ , and  $(T-2) \times 1$  vector  $y = (y_3, \ldots, y_T).$ 

#### Model fitting via least-squares:

$$\min_{w} \|X^T w - y\|_2^2$$

Prediction rule: at time T + 1,  $\hat{y}_{T+1} = w_1 + w_2 y_T + w_3 y_{T-1} = w^T x_{T+1}$ .

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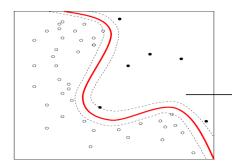
 $y_{t-1}, y_{t-2}$  $y_t = w_1 + w_2 y_{t-1} + w_3 y_{t-2} + w_4 y_{t-1}^2 + w_5 y_{t-1} y_{t-2} + w_6 y_{t-2}^2.$ 

Nonlinear auto-regressive model for time-series: y<sub>t</sub> quadratic function of

This writes  $y_t = w^T \phi(x_t)$ , with  $\phi(x_t)$  the augmented feature vectors  $\phi(x_t) := \left(1, y_{t-1}, y_{t-2}, y_{t-1}^2, y_{t-1}y_{t-2}, y_{t-2}^2\right).$ 

Prediction rule is  $\hat{y}_{T+1} = w^T \phi(x_{T+1})$ .

#### Nonlinear classification



Non-linear (e.g., quadratic) decision boundary

$$w_1x_1 + w_2x_2 + w_3x_1^2 + w_4x_1x_2 + w_5x_2^2 + b = 0.$$

Writes 
$$w^T \phi(x) + b = 0$$
, with  $\phi(x) := (x_1, x_2, x_1^2, x_1 x_2, x_2^2)$ .

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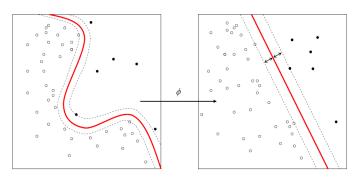
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# Challenges

In principle, it seems we can always augment the dimension of the feature space to make the data linearly separable. (See the video at

http://www.youtube.com/watch?v=3liCbRZPrZA)



How do we do it in a computationally efficient manner?

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$$\min_{\mathbf{w}} \ \mathcal{L}(\mathbf{X}^T \mathbf{w}, \mathbf{y}) + \lambda \|\mathbf{w}\|_2^2$$

where

- $ightharpoonup X = [x_1, \dots, x_m]$  is a  $n \times m$  matrix of data points.
- ▶  $y \in \mathbf{R}^m$  contains a response vector (or labels).
- $\mathbf{v} \in \mathbf{R}^n$  contains classifier or regression coefficients.
- $ightharpoonup \mathcal{L}$  is a "loss" function that depends on the problem considered.
- $\lambda \geq 0$  is a regularization parameter.

*Prediction/classification rule:* depends only on  $w^T x$ , where  $x \in \mathbf{R}^p$  is a new data point.

Note: here, we consider *l*<sub>2</sub>-norm penalty only!

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$$\mathcal{L}(z,y) = \|z-y\|_2^2 = \sum_{i=1}^n (x_i - y_i)^2.$$

► Hinge loss: (for SVMs)

$$\mathcal{L}(z,y) = \sum_{i=1}^{m} \max(0,1-y_i z_i)$$

► Logistic loss: (for logistic regression)

$$\mathcal{L}(z,y) = -\sum_{i=1}^{m} \log(1 + e^{-y_i z_i}).$$

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$$\min_{w} \mathcal{L}(X^T w, y) + \lambda ||w||_2^2$$

the optimal w lies in the span of the data points  $(x_1, \ldots, x_m)$ :

$$w = Xv$$

for some vector  $v \in \mathbf{R}^m$ .

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### Proof

#### A linear algebra result

For any matrix  $X \in \mathbf{R}^{n \times m}$ : every  $w \in \mathbf{R}^n$  can be written as the sum of two *orthogonal* vectors, one in the range of X and the other orthogonal to it:

$$w = Xv + r$$

where  $v \in \mathbf{R}^m$ , and  $X^T r = 0$  (that is, r is in the nullspace  $\mathcal{N}(X^T)$ ).

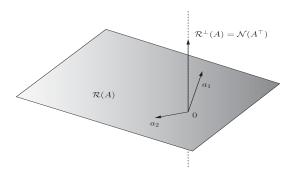


Figure shows the case  $X = A = (a_1, a_2)$ .

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$$\min_{w} \mathcal{L}(X^{T}w, y) + \lambda ||w||_{2}^{2}$$

the optimal w can be written as w = Xv for some vector  $v \in \mathbf{R}^m$ .

Hence training problem depends only on the  $m \times m$  (PSD) matrix  $K := X^T X$ :  $\min_{v} \ \mathcal{L}(Kv,y) + \lambda v^T Kv.$ 

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The training problem depends only on the "kernel matrix"  $K = X^T X$ 

$$\textit{K}_{ij} = \textit{x}_i^T \textit{x}_j, \ 1 \leq i, j \leq \textit{n}.$$

That is, K contains the scalar products between all data point pairs.

The prediction/classification rule depends on the scalar products between new point x and the training data points  $x_1, \ldots, x_m$ :

$$w^T x = v^T X^T x = v^T k, \quad k := X^T x = (x^T x_1, \dots, x^T x_m).$$

Computational advantage: Once K is formed (this takes  $O(n^2p)$ ), then the training problem has only n variables. When p >> n, this leads to a dramatic reduction in problem size.

#### How about the nonlinear case?

In the nonlinear case, we simply replace the feature vectors  $x_i$  by some "augmented" feature vectors  $\phi(x_i)$ , with  $\phi$  a non-linear mapping.

Example: in classification with quadratic decision boundary, we use

$$\phi(x) := (1, x_1, x_2, x_1^2, x_1 x_2, x_2^2).$$

This leads to the modified kernel matrix

$$K_{ij} = \phi(x_i)^T \phi(x_i), \quad 1 \leq i, j \leq m.$$

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$$k(x,z) = \phi(x)^T \phi(z).$$

It provides information about the metric in the feature space, e.g.:

$$\|\phi(x) - \phi(z)\|_2^2 = k(x,x) - 2k(x,z) + k(z,z).$$

The computational effort involved in

- solving the training problem;
- making a prediction,

depends only on our ability to quickly evaluate such scalar products.

We can't choose k arbitrarily; it has to satisfy the above for some  $\phi$ .

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#### Quadratic kernels

Classification with quadratic boundaries involves feature vectors

$$\phi(x) = (1, x_1, x_2, x_1^2, x_1 x_2, x_2^2).$$

*Fact*: given two vectors  $x, z \in \mathbf{R}^2$ , we have

$$\phi(x)^T\phi(z)=(1+x^Tz)^2.$$

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More generally when  $\phi(x)$  is the vector formed with all the products between the components of  $x \in \mathbf{R}^n$ , up to degree d, then for any two vectors  $x, z \in \mathbf{R}^n$ ,

 $\phi(x)^T \phi(z) = (1 + x^T z)^d$ .

Computational effort grows linearly in *n*.

This represents a dramatic reduction in speed over the "brute force" approach:

- Form  $\phi(x)$ ,  $\phi(z)$ ;
- evaluate  $\phi(x)^T \phi(z)$ .

Computational effort grows as  $n^d$ .

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

where  $\sigma>0$  is a scale parameter. Allows to ignore points that are too far apart. Corresponds to a non-linear mapping  $\phi$  to infinite-dimensional feature space.

There is a large variety (a zoo?) of other kernels, some adapted to structure of data (text, images, etc).

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## Kernels and sparsity

Kernel methods essentially lift the space of features, to pose the original problem in higher-dimensional space.

Sparse methods seek to improve the interpretability of the results, by selecting a few features that are relevant to the prediction problem at hand.

Unfortunately kernels and sparsity don't mix well, since we loose the interpretation of the original features in the "lifting" process.

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# Sparse methods: basic idea

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In some cases we are interested in the *interpretability* of the regression parameters:

- Which few features are responsible for most of the predictive power?
- Can we find those features in a reliable way?

Sparse methods attempt to trade-off sparsity of the optimal regression or classification coefficients with predictive power.

In such methods, the  $l_1$ -norm plays an important role.

Consider the *cardinality-constrained* regularized LS:

$$\min_{w} \ \|X^{T}w - y\|_{2}^{2} + \lambda \|w\|_{2}^{2} \ : \ \mathbf{Card}(w) = k$$

where  $\lambda>0$  is the regularization parameter,  $\mathbf{Card}(w)$  denotes cardinality of w (number of nonzero elements in w), and k imposes a bound on it.

Problem above is very difficult to solve. We can approximate it as

$$\min_{w} \|X^{T}w - y\|_{2}^{2} + \frac{\lambda}{k} \|w\|_{1}^{2}.$$

**Proof:** Cauchy-Schwartz inequality implies

$$\sqrt{\text{Card}(w)}\cdot \|w\|_2 \geq \|w\|_1.$$

When  $\lambda$  varies, the set of solutions is the same as that of the "LASSO" problem  $\min \|X^T w - y\|_2^2 + \mu \|w\|_1$ 

where  $\mu > 0$  spans the positive real line.

- Not solvable by direct linear algebra methods.
- Many efficient algorithms have bee proposed for this type of convex problem
- When the number of variables is not too high, coordinate descent (solving the problem one variable at a time, all the others being fixed) works well.

# Example: image of Microsoft in NYT headlines

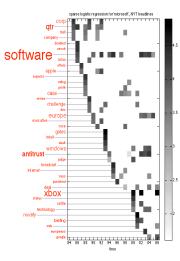


Image of microsoft in NYT headlines (1981-2007).

 Data matrix X records occurrences of words in

documents.

- Response ±1 vector y records presence/absence of term microsoft
- Solve LASSO to regress occurrence of Microsoft against other word occurrences.
- Sparsity allows to find just a few words whose occurrence predicts that of the term "Microsoft" in headlines.

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This is due to the fact that the solution to the problem

$$\min_{w} \ \| \boldsymbol{X}^T \boldsymbol{w} - \boldsymbol{y} \|_2^2 + \mu \| \boldsymbol{w} \|_1,$$

does not depend only on  $K := X^T X$ , but on X itself. This is in contrast with the Ridge regression problem

$$\min_{w} \|X^T w - y\|_2^2 + \mu \|w\|_2^2.$$

The main reason is the lack of rotational invariance of the  $l_1$ -norm.

## Kernel methods: summary

- Kernels need to be chosen by the user.
- Choice not always obvious; Gaussian or polynomial kernels are popular.
- Control over-fitting via cross validation (wrt say, scale parameter of Gaussian kernel, or degree of polynomial kernel).
- Kernel methods not well adapted to sparsity / I<sub>1</sub>-norm regularization (seen next).

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#### **Pre-Processing**

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#### Pre-processing

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Before we start processing data via a supervised learning algorithm, we need to prepare data:

- Remove missing values;
- Remove outlier values:
- Normalize (offset and rescale) data points;
- Normalize features.

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Consider the penalized least-squares problem:

$$\min_{w} \|X^{T}w - y\|_{2}^{2} + \rho^{2}\|w\|_{2}^{2}. \tag{1}$$

The prediction rule for a new data point x is  $\hat{y}(x) = w^T x$ .

It is common to center the  $n \times m$  data matrix  $X = [x_1, \dots, x_m]$  and response y, and modify it to  $X^c := [x_1 - \bar{x}, \dots, x_m - \bar{x}], \ y^c = y - \bar{y}$  with

$$\bar{x} = \frac{1}{m} \sum_{i=1}^{m} x_i, \quad \bar{y} = \frac{1}{m} \sum_{i=1}^{m} y_i.$$

This is equivalent to solving the problem

$$\min_{w,b} \|X^T w + b\mathbf{1} - y\|_2^2 + \rho^2 \|w\|_2^2,$$

since the optimal *b* is  $b = \bar{y} - \bar{x}^T w$ .

The prediction rule is now  $\hat{y}(x) = w^T x + b$ .

In effect, for the LS case, centering implicitly allows to consider a more general affine rule (as opposed to linear, as in problem (1)).

not have a natural interpretation as an affine rule.

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It is thus always better to work with affine rules (arguably more general than linear ones):

$$\min_{w,b} \mathcal{L}(X^T w + b\mathbf{1}, y) + p(w)$$

Such an approach provides an implicit "centering".

Sometimes, scaling the features provides better performance. Just as with centering, scaling should be done differently with different loss functions.

- The standard approach is to center, and then normalize data by variance:  $X \to D^{-1}(X - \hat{x}\mathbf{1}^T)$ , where  $D = \operatorname{diag}(\sigma_1, \dots, \sigma_n)$ .
- The problem becomes

$$\min_{w,b} \mathcal{L}((X - \hat{x} \mathbf{1}^T)^T D^{-1} w, y) + \rho(w) = \mathcal{L}(X^T D^{-1} w - \mathbf{1}(\hat{x}^T D^{-1} w), y) + \rho(w)$$

which is equivalent, under the change of variable  $w \to D^{-1} w$ , to

$$\min_{w,b} \mathcal{L}(X^T w + b\mathbf{1}, y) + p(Dw).$$

 One may use robust methods, where mean and standard deviation are replaced with median and mean absolute deviation.

## Scaling data points

In practice, it may be advisable to scale data points differently.

- In time-series prediction, we may use exponential smoothing in order to give more importance to recent time-series values than older ones.
- In imbalanced classification, we may need to scale the negatively labelled data differently from the positively labelled data.

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