The curse of dimensionality



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Introduction

Modern data are often high dimensional.

 computational biology: DNA, few observations and huge number of variables;



Introduction

 images or videos: an image from a digital camera has millions of pixels, 1h of video contains more than 130000 images



 data coming from consumer preferences: Netflix for instance owns a huge (but sparse) database of ratings given by millions of users on thousands of movies or TV shows.



The curse of dimensionality:

 this term was first used by R. Bellman in the introduction of his book "Dynamic programming" in 1957:

All [problems due to high dimension] *may be subsumed under the heading "the curse of dimensionality". Since this is a curse,* [...], *there is no need to feel discouraged about the possibility of obtaining significant results despite it.*

- he used this term to talk about the difficulties to find an optimum in a high-dimensional space using an exhaustive search,
- in order to promote dynamic approaches in programming.

In high dimensional spaces, nobody can hear you scream

Concentration phenomena

Surprising asymptotic properties for covariance matrices



Supervised classification or regression often rely on local averages:

Classification : you know the classes of n points from your learning database, you can classify a new point x by computing the most represented class in the neighborhood of x.



15-Nearest Neighbor Classifier

Regression : you observe n i.i.d observations (x^i, y^i) from the model

$$y^i = f(x^i) + \epsilon_i,$$

and you want to estimate f. If you assume f is smooth, a simple solution consists in estimating f(x) as the average of all y_i corresponding to the k nearest neighbors x_i of x.



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Makes sense in small dimension. Unfortunately, not so much when the dimension p increases...



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, $p = 10$, $s^{1/p} = 0.63$



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To overcome this limitation, you need a number of sample which grows exponentially with $p\ldots$



Nearest neighbors

X,Y two independent variables, with uniform distribution on $[0,1]^p.$ The mean square distance $\|X-Y\|^2$ satisfies

$$\mathbb{E}[\|X - Y\|^2] = p/6$$
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The notion of nearest neighbors vanishes.

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Volume of the ball

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Consequence: if you want to cover $[0,1]^p$ with a union of n unit balls, you need

$$n \geq \frac{1}{V_p} = \frac{\Gamma(p/2+1)}{\pi^{p/2}} \overset{p \to \infty}{\sim} \left(\frac{p}{2\pi e}\right)^{\frac{p}{2}} \sqrt{p\tau}$$

For p = 100, $n = 42 \ 10^{39}$.

Assume you draw n samples with uniform law in the hypercube, most sample points will be in corners of the hypercube :



Volume of the shell

Probability that a uniform variable on the unit sphere belongs to the shell between the spheres of radius 0.9 and 1 is

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Fig. Probability that X belongs to the shell $S_{0.9}$ regarding to the dimension p.

 $X_1, \ldots X_n$ i.i.d. in dimension p, with uniform distribution on the unit ball. Median distance from the origin to the closest data point is given by

$$med(p,n) = \left(1 - \frac{1}{2^{\frac{1}{n}}}\right)^{\frac{1}{p}}.$$

For n = 500 and p = 10, med = 0.52, which means that most data points are closer to the edge of the ball than to the center.

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Exemple. Assume n data sampled independently with a uniform law on $[-1, 1]^p$. You want to estimate $e^{-||x||^2/8}$ in 0 from your data. You choose as an estimator the observed value in x_i , the nearest neighbor of 0. For n = 1000 samples and p = 10, the probability that this nearest neighbor is at a distance larger than $\frac{1}{2}$ from 0 is around 0.99.

Where is located the mass of the Gaussian distribution ?

Mass of the standard Gaussian distribution in the ring between radius r and r+dr

$$\begin{split} \mathbb{P}[r \le \|X\| \le r + dr] \simeq \frac{e^{-r^2/2}}{(2\pi)^{p/2}} (V_p(r + dr) - V_p(r)) \simeq \frac{e^{-r^2/2}}{(2\pi)^{p/2}} r^{p-1} \ pdr \ V_p(1). \\ \longrightarrow \text{ maximum for } r = \sqrt{p-1} \end{split}$$



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Most of the mass of a Gaussian distribution is located in areas where the density is extremely small compared to its maximum value. 19

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Sample Covariance Matrices appear everywhere in statistics

- classification with gaussian mixture models
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Problems:

- often necessary to invert Σ
- if n is not large enough, the estimates of Σ are ill-conditionned or singular
- \blacksquare sometimes necessary to estimate the eigenvalues of Σ

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Random matrices

• If
$$n,p \to \infty$$
 with $p/n \to c > 0,$ then

 $\|\widehat{\Sigma}_p - I_p\|_2 \not\to 0 \quad (\|\|_2 \text{ denotes the spectral norm}).$

• Even false for
$$p/n = 1/100$$
.

Context $x_1 \dots, x_n \in \mathbb{R}^p$ i.i.d. samples from a gaussian multivariate distribution $\mathcal{N}(0, I_p)$. Note $X = (x_1, \dots, x_p)$.

- $\bullet \ p/n = c > 1$
- Convergence in $\|\|_{\infty}$

$$\max_{i,j} |\widehat{\Sigma}_{i,j} - \delta_{i,j}| \xrightarrow{a.s.} 0$$

However, we lose the convergence in spectral norm since

 $\operatorname{rank}(X) \le p \implies \lambda_{\min}(\widehat{\Sigma_p}) = 0 < 1 = \lambda_{\min}(\Sigma_p)$

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No contradiction with the fact that all norms are equivalent in finite dimension.

More precisely, the random matrices theory tells us that when $p, n \to \infty$ with $p/n \to c > 0$, then [Marčenko-Pastur Theorem, 1967]

$$\frac{1}{p}\sum_{k=1}^p \delta_{\lambda_k(\widehat{\Sigma_p})} \xrightarrow{a.s.} \mu \quad \text{weakly},$$

with μ the Marčenko-Pastur law of parameter $c_{\rm r}$ which satisfies

•
$$\mu(\{0\}) = max(0, 1 - c^{-1})$$

• on $(0,\infty)$, μ has a continuous density supported on $[(1-\sqrt{c})^2, (1+\sqrt{c})^2].$









Figure: Marčenko-Pastur law for different limit ratios $c = \lim_{p \to \infty} p/n$.



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Classical ways to avoid the curse of dimensionality

Dimension reduction:

- the problem comes from that p is too large,
- therefore, reduce the data dimension to $d \ll p$,
- such that the curse of dimensionality vanishes!

Regularization:

- the problem comes from that parameter estimates are unstable,
- therefore, regularize these estimates,
- such that the parameter are correctly estimated!

Parsimonious models:

- the problem comes from that the number of parameters to estimate is too large,
- therefore, make restrictive assumptions on the model,
- such that the number of parameters to estimate becomes more "decent"!