Dimensionality Reduction

Algorithms in Machine Learning, ISAE-SUPAERO

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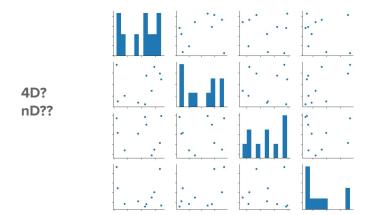
Dimensionality reduction: why?

What is high dimension?

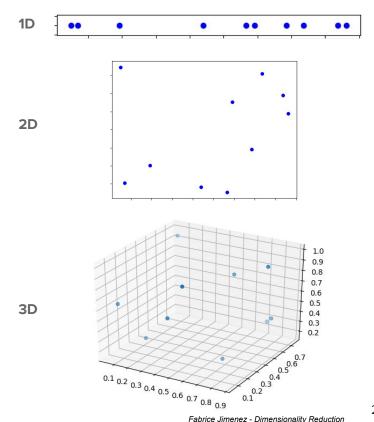
v1	v2	v3	v4	 v365
8.4	15	2.2	0.5	 65.8
9.1	10	5.1	-4.3	 -7

Many, many features... Maybe more features than data points!

Let's consider 10 points in 20 dimensions v1 ... v20



First issue: visualization



One so-called "Curse", several effects...

Effect n°1 - Data Coverage

Statistics, Machine Learning: generalization / estimation of population, from a reduced sample which is representative

Representative = training set covering "enough" portions of feature space

Imagine we need to train a binary classifier (red or blue points)...

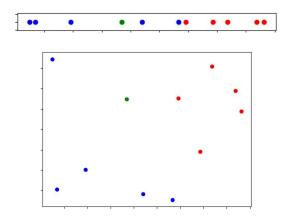


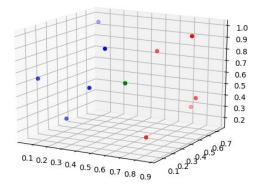
How would you classify the new green point?

1D

2D

3D





One so-called "Curse", several effects...

Effect n°1 - Data Coverage

Statistics, Machine Learning: generalization / estimation of population, from a reduced sample which is representative

Representative = training set covering "enough" portions of feature space

Imagine we need to train a binary classifier (red or blue points)...



Ideally, training samples "uniformly" distributed on the feature space...

If in 1D you need ~10 points:

- -> in 2D you need ~100 points
- -> in 3D you need ~1000 points
- -> in nD you need ~10^n points

For fixed number of samples, many dimensions = poor coverage

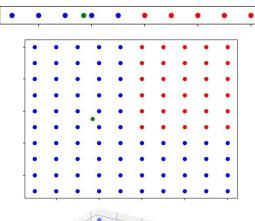
= classifier with poor performance!!!

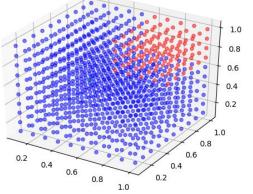
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One so-called "Curse", several effects...

Effect n°2 - Distance concentration

With high dimension, sparsity of points increases, and Euclidean distance becomes more and more equal between all points

Notion of distance important for many ML tasks: clustering, outlier detection...

Many dimensions = distance obsolete = results obsolete...



Let's take our dataset with 10 points in 20 dimensions:

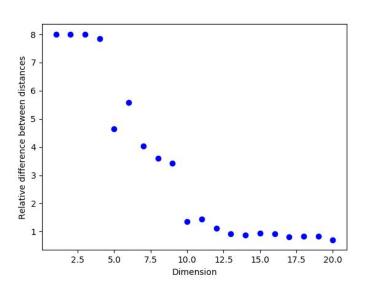
- -> For each dimension d, keep the d first variables
- -> Compute all pairwise distances
- -> Study the relative difference between distances

More details:

On the surprising behavior of distance metrics in high dimensional space

Charu C. Aggarwal et. al.

$$\lim_{d o\infty} E\left(rac{\operatorname{dist}_{\max}(d) - \operatorname{dist}_{\min}(d)}{\operatorname{dist}_{\min}(d)}
ight) o 0$$



Roughly, when d exceeds the number of points, distance becomes useless...

One so-called "Curse", several effects...

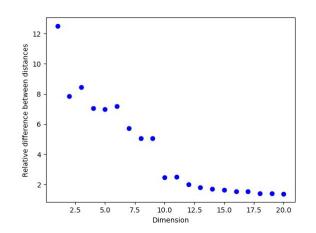
Effect n°3 - Noise pollution

When looking for a pattern in low dimension, if you add many dimensions irrelevant to this pattern: noise conceals the actual information

Relative difference between distances:

- → Main jump between 1D and 2D (introduction of noise)
- → Second jump roughly when d > number of points (effect n°2)





Can you identify the pattern? 1D 2D **3D** 2.6 2.4 2.2 $1.4_{1.2} \underbrace{1.0_{0.8}_{0.6}}_{0.4} \underbrace{0.2_{0.0}}_{0.375} \underbrace{3.50}_{3.75} \underbrace{3.00}^{2.75} \underbrace{2.25}^{2.00}$

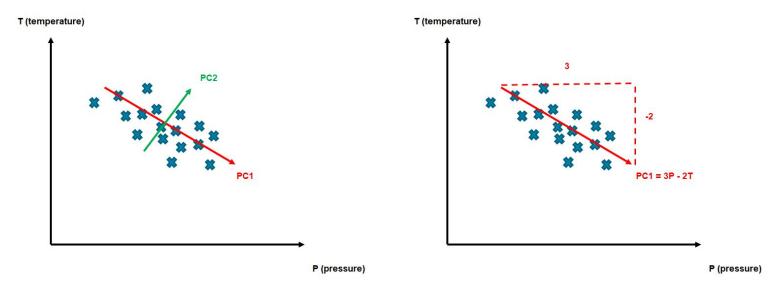
PCA and information pattern

Effects especially valid if dimensions are drawn from independent distributions

Correlations reinforce patterns visibility, but how to distinguish correlation information, noise, and pattern information?

Principal Components Analysis → Linear combination of initial features

- → First components maximize the projected variability of the dataset = spread distances = often main information patterns
- → Correlated features are gathered in the same components: removes correlation redundancy



Defeat the Curse of Dimensionality? Compute PCA and keep the first principal components might be a solution...

PCA Computation

Dataset Matrix

Centerina

Scaling

$$M = \begin{bmatrix} X_{1\,1} & \dots & X_{1\,p} \\ \dots & \dots & \dots \\ X_{n\,1} & \dots & X_{n\,p} \end{bmatrix}$$

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$$\tilde{M} = \begin{bmatrix} \frac{X_{11} - \bar{X}_{1}}{\sigma(X_{1})} & \dots & \frac{X_{1p} - \bar{X}_{p}}{\sigma(X_{p})} \\ \dots & \dots & \dots \\ \frac{X_{n1} - \bar{X}_{1}}{\sigma(X_{1})} & \dots & \frac{X_{np} \bar{X}_{p}}{\sigma(X_{p})} \end{bmatrix}$$

Covariance matrix:

$$cov(M) = \frac{1}{n} \times \bar{M}^T.\bar{M}$$

Correlation matrix:

$$cor(M) = \frac{1}{n} \times \tilde{M}^T.\tilde{M}$$

Eigenvalue decomposition:

$$A = V^T.D.V$$

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 where $D = diag(\lambda_1, \lambda_2, ..., \lambda_n)$

- applies to square matrices n x n which are diagonalizable
- V: columns of V are eigenvectors Vi for the corresponding eigenvalues λ_i i.e $AV_i=\lambda_i V_i$

$$\lambda_i$$
 i.e $AV_i = \lambda_i V_i$

PCA Computation:

- Correlation matrix is square, symmetric, real: diagonalizable in orthonormal basis (Spectral theorem...)
- **Principal Components:** eigenvectors of correlation matrix
- Explained variance by component i: ith eigenvalue

Exercise: PCA and diagonalization

Demonstrate that principal components of the PCA are the eigenvectors of the correlation matrix



1/ Consider principal component with direction vector p
Our dataset M projected on this component becomes F such as:

$$F = \tilde{M}p \qquad \begin{cases} \bar{F} = 0 \\ \|p\| = 1 \end{cases}$$

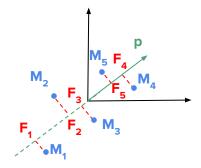
2/ Write the explained variance on this component:

$$var(F) = \frac{1}{n} \sum_{i=1}^{n} (Fi - \bar{F})^{2} = \frac{1}{n} \sum_{i=1}^{n} Fi^{2} = \frac{1}{n} F^{T} F$$

$$= \frac{1}{n} (\tilde{M}p)^{T} \tilde{M}p$$

$$= \frac{1}{n} p^{T} (\tilde{M}^{T} \tilde{M}) p$$

$$var(F) = p^{T} cor(M) p$$



3/ We want the component to explain a maximum of variance, we need to formulate the following maximization problem:

$$\begin{cases} maximize \left[p^{T}cor\left(M \right)p \right] \\ \|p\| = p^{T}p = 1 \end{cases} \iff \begin{cases} \mathcal{L}\left(p \right) = p^{T}cor\left(M \right)p - \mu\left(p^{T}p - 1 \right) \\ \frac{\partial \mathcal{L}}{\partial p}\left(p \right) = 2cor\left(M \right)p - 2\mu p = 0 \\ \iff cor\left(M \right)p = \mu p \end{cases}$$

4/ Explained variance on the component is then:

$$var(F) = p^{T}cor(M)p = p^{T}\mu p = \mu$$

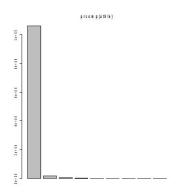
Explained variance is eigenvalue of the correlation matrix

PCA Interpretation

Be careful with quick interpretations!

Example 1:

One of the variables "draws" all the variance of the dataset: careful with scaling!

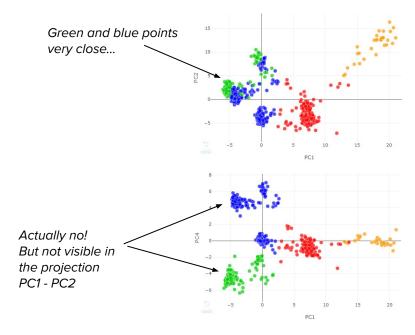


No scaling: 1 variable with high variance "draws" all PCA effect to itself **Scaling:** 1 noise variable will have same variance as information variable

If different units: scaling mandatory, otherwise does not make sense!

Example 2:

Be careful with proximity of points in projection graph

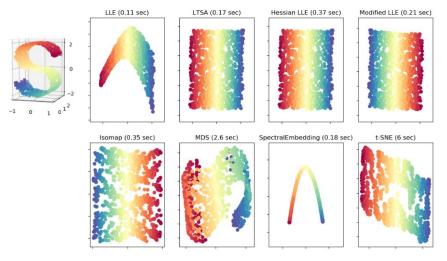


Think about these examples when you get your results...

PCA vs Manifold Learning

PCA is nice when target pattern is linked to variance in a linear direction...

Unfortunately, it's not always the case: patterns may be non-linear. Non-linear projection methods exist to reduce dimension:



(source: scikit-learn)

Manifold Learning

We will see 1 example in this course: t-SNE (t-distributed Stochastic Neighbor Embedding)

t-distributed Stochastic Neighbor Embedding (t-SNE)

Particular technique of dimensionality reduction: it transcribes the similarities between points in a lower dimension

Similarities in initial space: normal probability density around each point

Gaussian Distribution Around Data Point

More details on t-SNE:

Visualizing data using t-SNE Laurens van der Maaten et. al.

Perplexity: parameter ruling the coverage
→ to change sigma

$$P_{ij} = \frac{exp(-||x_i - x_j||^2/2\sigma_i^2)}{\sum_{k \neq i} exp(-||x_i - x_k||^2/2\sigma_i^2)}$$

then

$$P_{ij} = \frac{P_{ij} + P_j}{2n}$$

t-distributed Stochastic Neighbor Embedding (t-SNE)

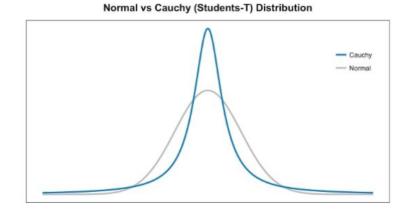
Particular technique of dimensionality reduction: it transcribes the similarities between points in a lower dimension

Similarities in reduced space: Cauchy probability density around each point

 $Q_{ij} = \frac{(1+||y_i - y_j||^2)^{-1}}{\sum_{k \neq i} (1+||y_i - y_k||^2)^{-1}}$

Learning rate and stopping criteria:

parameters impacting accuracy of the projection



Objective:

minimize difference between distance distributions in initial and reduced space

How:

optimization (gradient descent or others) on the Kullback-Leibler divergence KL(P,Q)

Going from normal to Cauchy distribution: stretches the distances → highly differentiated clusters (sometimes too much!)

Pros: high transcription capability of complex patterns

Cons: sensitive to tuning (optimization algorithm)

Dimensionality Reduction: Application

It's time to play on your own...

Main interest = discover the dataset, play with PCA, t-SNE parameters, and take a step back for interpretation



Questions?

