

Dimensionality reduction and data visualization

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Introduction

Supervised learning (predictive methods)

- Develop predictive models using **labeled training data**
- Ensure that the models perform well on **future data (test data)**



Unsupervised learning (descriptive methods)

- Data exploration
 - Analyze distribution/geometry of the data
 - Goal: acquire or extract knowledge / patterns from data
- **Dimension reduction, visualization, clustering**



Data exploration

d variables

	rcc	wcc	hc	hg	ferr	bmi	ssf	pcBfat	lbm	ht	wt	sex
	4.82	7.6	43.2	14.4	58	22.37	50	11.64	53.11	163.9	60.1	f
	4.32	6.8	40.6									f
	5.16	7.2	44.3									f
	4.66	6.4	40.9									f
	4.19	9	39									f
	4.53	5	40.7									f
	4.42	6.4	42.8									f
	4.32	4.3	41.6									m
	4.73	6.7	42.8									m
	4.71	7.2	43.6									m
	4.93	7.3	46.2									m
	5.21	7.5	47.5									m
Point x_i	5.09	8.9	46.3	15.4	44	29.97	71.1	13.97	88	185.1	102.7	m
	5.11	9.6	48.2	16.7	103	27.39	65.9	11.66	83	185.5	94.2	m
	4.94	6.3	45.7	15.5	50	23.11	34.3	6.43	74	184.9	79	m
	4.86	3.9	44.9	15.4	73	22.83	34.5	6.56	70	181	74.8	m
	4.51	4.4	41.6	12.7	44	19.44	65.1	15.07	53.42	179.9	62.9	f
	4.62	7.3	43.8	14.7	26	21.2	76.8	18.08	61.85	188.7	75.5	f

Data matrix

- sample $x_i = (x_{i,1} \ \cdots \ x_{i,d})^\top$
- $X = \begin{pmatrix} x_{1,1} & \cdots & x_{1,d} \\ \vdots & & \vdots \\ x_{n,1} & \cdots & x_{n,d} \end{pmatrix} \in \mathbb{R}^{n \times d}$

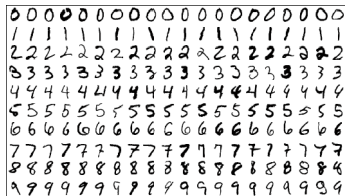
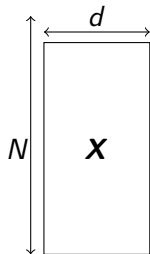
n points

Variable j (hemoglobin)

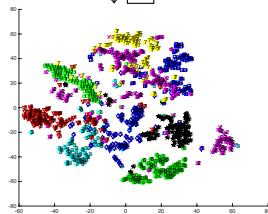
What are the relations between the variables? How close are the samples?

Dimension reduction: the goal

- Let $\mathbf{X} \in \mathbb{R}^{N \times D}$ the data (N samples of dimension d)
- Goal: find a projection of \mathbf{X} onto $\mathbf{Z} \in \mathbb{R}^{N \times q}$ with $q < d$



$d = 784$



$q = 2$

What for?

- Visualization ($q = 2$ ou 3)
 - check the data
 - identify outliers
 - visualize the data according to their categories (if labelled)
- Data representation ($q < d$)
 - Noise reduction
 - pre-processing: computation issue
 - hidden structure in the data (example: manifolds)

Coding/Encoding scheme

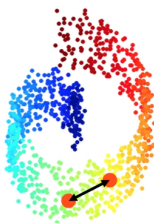
$$\begin{aligned} \text{cod} : \mathbb{R}^d &\longrightarrow \mathbb{R}^q, & \mathbf{x} &\longmapsto \mathbf{z} = \text{cod}(\mathbf{x}) \\ \text{dec} : \mathbb{R}^q &\longrightarrow \mathbb{R}^d, & \mathbf{z} &\longmapsto \mathbf{x} = \text{dec}(\mathbf{z}) \end{aligned}$$

How to assess the quality of the coding?

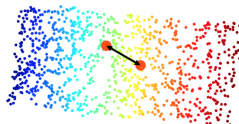
Principle of dimension reduction methods

- Project samples $\{\mathbf{x}_i \in \mathbb{R}^d\}_{i=1}^N$ onto $\{\mathbf{z}_i \in \mathbb{R}^q\}_{i=1}^N$ ($q < d$) such that the data topology is preserved
 - preserve distance between samples
 - preserve the neighborhood ...

$$\mathbf{x}_i \in \mathbb{R}^3$$



$$\mathbf{z}_i \in \mathbb{R}^2: \text{distance preservation}$$



Methods we will study

Linear : PCA, non-linear : SNE and t-SNE variant

Principal Component Analysis (PCA)

Model: data = information + noise

$$\mathbf{X} = \mathbf{Z}\mathbf{P}^\top + \mathbf{B}$$

Linear orthogonal projection:

$$\begin{aligned} \text{cod} : \mathbb{R}^d &\longrightarrow \mathbb{R}^q, & \mathbf{x} &\longmapsto \mathbf{z} = \mathbf{P}^\top \mathbf{x} \\ \text{dec} : \mathbb{R}^q &\longrightarrow \mathbb{R}^d, & \mathbf{z} &\longmapsto \hat{\mathbf{x}} = \mathbf{P}\mathbf{z} \end{aligned}$$

Property: columns of \mathbf{P} are orthogonal

Dimensions:

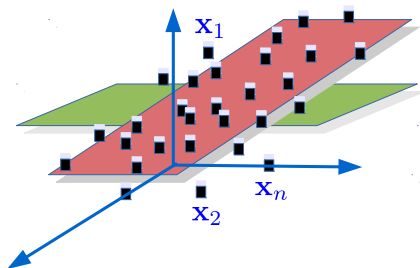
$$\mathbf{X} = \begin{pmatrix} \mathbf{x}_1^\top \\ \vdots \\ \mathbf{x}_N^\top \end{pmatrix} \in \mathbb{R}^{N \times d}, \quad \mathbf{Z} = \begin{pmatrix} \mathbf{z}_1^\top \\ \vdots \\ \mathbf{z}_N^\top \end{pmatrix} \in \mathbb{R}^{N \times q}, \quad \mathbf{P} \in \mathbb{R}^{d \times q}$$

Objective: minimize error between \mathbf{x}_i and its estimation $\hat{\mathbf{x}}_i = \text{dec}(\text{cod}(\mathbf{x}_i))$

$$\min_{\mathbf{P} \in \mathbb{R}^{d \times q}} \sum_{i=1}^N \|\mathbf{x}_i - \mathbf{P}\mathbf{P}^\top \mathbf{x}_i\|^2$$

Another view of PCA

PCA linearly projects $\{\mathbf{x}_i \in \mathbb{R}^d\}_{i=1}^N$ onto a subspace of dimension q ($q < d$) such that the **variance** of the projections $\{\mathbf{z}_i = \mathbf{P}^\top \mathbf{x}_i \in \mathbb{R}^q\}_{i=1}^N$ remains maximal



Variance maximization (case $q = 1$)

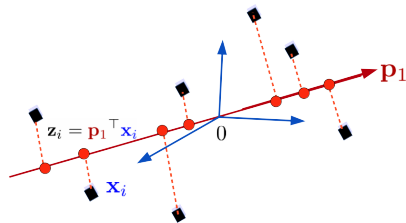
$$\max_{\mathbf{p} \in \mathbb{R}^q} \|\mathbf{X}\mathbf{p}\|_2^2 \quad \text{with} \quad \|\mathbf{p}\|_2^2 = 1 \quad \text{and} \quad \mathbf{Z} = \mathbf{X}\mathbf{p}$$

Minimization of error / maximization of variance

$$\begin{aligned}
 J(\mathbf{P}) &= \sum_{i=1}^N \|\mathbf{x}_i - \hat{\mathbf{x}}_i\|^2 = \sum_{i=1}^N (\mathbf{x}_i - \mathbf{P}\mathbf{P}^\top \mathbf{x}_i)^\top (\mathbf{x}_i - \mathbf{P}\mathbf{P}^\top \mathbf{x}_i) \\
 &= \sum_{i=1}^N (\mathbf{x}_i^\top \mathbf{x}_i - 2\mathbf{x}_i^\top \mathbf{P}\mathbf{P}^\top \mathbf{x}_i + \mathbf{x}_i^\top \mathbf{P}\mathbf{P}^\top \mathbf{P}\mathbf{P}^\top \mathbf{x}_i) \\
 &= \sum_{i=1}^N \mathbf{x}_i^\top \mathbf{x}_i - \sum_{i=1}^N \mathbf{x}_i^\top \mathbf{P}\mathbf{P}^\top \mathbf{x}_i = \sum_{i=1}^N \mathbf{x}_i^\top \mathbf{x}_i - \sum_{i=1}^N \mathbf{z}_i^\top \mathbf{z}_i \\
 &= \text{trace} \left(\sum_{i=1}^N \mathbf{x}_i \mathbf{x}_i^\top - \frac{1}{N} \sum_{i=1}^N \mathbf{z}_i \mathbf{z}_i^\top \right) = \text{trace} \left(\sum_{i=1}^N \mathbf{x}_i \mathbf{x}_i^\top - \sum_{i=1}^N \mathbf{P}^\top \mathbf{x}_i \mathbf{x}_i^\top \mathbf{P} \right) \\
 J(\mathbf{P}) &= \text{trace}(\mathbf{X}^\top \mathbf{X}) - \text{trace}(\mathbf{P}^\top \mathbf{X}^\top \mathbf{X} \mathbf{P})
 \end{aligned}$$

$\Rightarrow \min J(\mathbf{P}) \Leftrightarrow$ maximizing the variance of the projections w.r.t. \mathbf{P}

First projection vector \mathbf{p}_1 of \mathbf{P}



- Data: $\{\mathbf{x}_i \in \mathbb{R}^{N \times d}\}_{i=1}^N$
- Assume the \mathbf{x}_i are **normalized**
- Projections onto \mathbf{p}_1 : $\{\mathbf{z}_i = \mathbf{p}_1^\top \mathbf{x}_i \in \mathbb{R}\}_{i=1}^N$

Computing $\mathbf{p}_1 \in \mathbb{R}^d$

\mathbf{p}_1 : a **unit vector** that maximizes the variance of the $\{\mathbf{z}_i\}_{i=1}^N$

$$\max_{\mathbf{p}_1 \in \mathbb{R}^d} \frac{1}{N} \sum_{i=1}^N (\mathbf{p}_1^\top \mathbf{x}_i)^2 \quad \text{s.t.} \quad \|\mathbf{p}_1\|^2 = 1$$

→ Solve a constrained optimization problem

Computing \mathbf{p}_1

$$\max_{\mathbf{p}_1 \in \mathbb{R}^d} \mathbf{p}_1^\top \mathbf{C} \mathbf{p}_1 \quad \text{s.t.} \quad \|\mathbf{p}_1\|^2 = 1$$

$\mathbf{C} = \frac{1}{N} \sum_{i=1}^N \mathbf{x}_i \mathbf{x}_i^\top = \frac{1}{N} \mathbf{X}^\top \mathbf{X}$ is the correlation matrix

Solution derivation

- Lagrangian: $\mathcal{L}(\mathbf{p}_1, \lambda_1) = -\mathbf{p}_1^\top \mathbf{C} \mathbf{p}_1 + \lambda_1(\mathbf{p}_1^\top \mathbf{p}_1 - 1)$
- Optimality conditions :

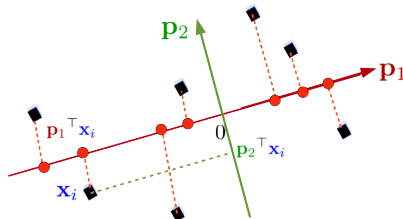
$$\nabla_{\mathbf{p}_1} \mathcal{L} = -2\mathbf{C} \mathbf{p}_1 + 2\lambda_1 \mathbf{p}_1 = 0 \quad \text{and} \quad \nabla_{\lambda_1} \mathcal{L} = \mathbf{p}_1^\top \mathbf{p}_1 - 1 = 0$$

$$\implies \mathbf{C} \mathbf{p}_1 = \lambda_1 \mathbf{p}_1 \quad \text{and} \quad \mathbf{p}_1^\top \mathbf{C} \mathbf{p}_1 = \lambda_1$$

- 1 $(\lambda_1, \mathbf{p}_1)$ is the couple (eigenvalue , eigenvector) of the \mathbf{C}
- 2 $\mathbf{p}_1^\top \mathbf{C} \mathbf{p}_1 = \lambda_1$ is the objective to maximize

\mathbf{p}_1 is the eigenvector associated to the highest eigenvalue of \mathbf{C} .

Computing \mathbf{p}_2 and beyond



- \mathbf{p}_2 : unit vector orthogonal to \mathbf{p}_1 that maximizes the variance of the projections $\{\mathbf{p}_2^\top \mathbf{x}_i\}_{i=1}^N$ onto \mathbf{p}_2

Solution

- \mathbf{p}_2 is the eigenvector associated to λ_2 , the 2nd highest eigenvalue of \mathbf{C}

Lemma

The sub-space of size k that maximizes the variance of the projection necessarily includes the sub-space of size $k - 1$.

PCA algorithm

- 1 Normalize the data : $\{\mathbf{x}_i \in \mathbb{R}^d\}_{i=1}^N \longrightarrow \{x_{ij} = \frac{x_{ij} - \bar{x}_j}{\sigma_j}, j = 1, d\}_{i=1}^N$
- 2 Compute the correlation matrix $\mathbf{C} = \frac{1}{N} \mathbf{X}^\top \mathbf{X}$
- 3 Find the eigenvalue decomposition $\{\mathbf{p}_j \in \mathbb{R}^d, \lambda_j \in \mathbb{R}\}_{j=1}^d$ of \mathbf{C}
- 4 Order the eigenvalues λ_j by decreasing order
- 5 The projection matrix is:

$$\mathbf{P} = (\mathbf{p}_1, \dots, \mathbf{p}_q) \in \mathbb{R}^{d \times q}$$

$\{\mathbf{p}_1, \dots, \mathbf{p}_q\}$ are the q eigenvectors associated to the q highest eigenvalues.

Application

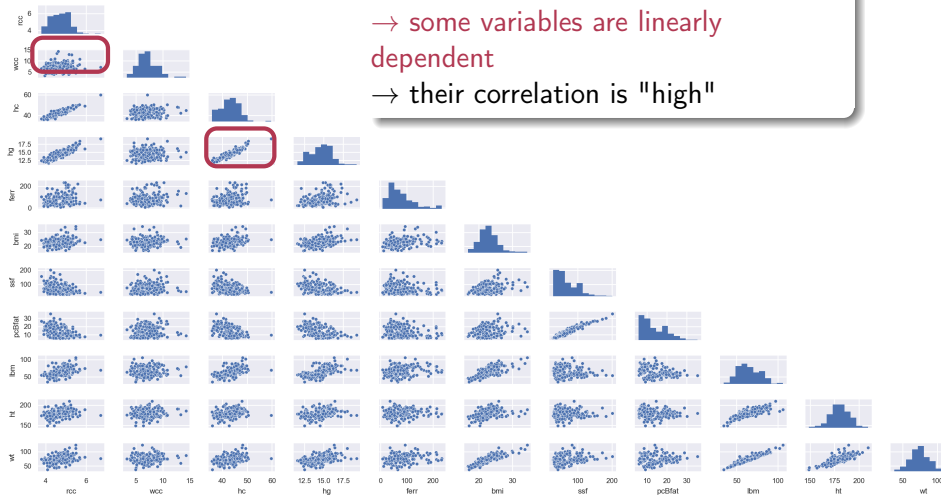
rcc	wcc	hc	hg	ferr	bmi	ssf	pcBfat	lbm	ht	wt	sex
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5.16	7.2	44.3	14.5	88	18.29	61.9	12.92	48.76	175	56	f
4.66	6.4	40.9	13.9	109	18.37	38.2	8.45	41.93	157.9	45.8	f
4.19	9	39	13.4	69	18.93	43.5	10.16	42.95	158.9	47.8	f
4.53	5	40.7	14	41	17.79	56.8	12.55	38.3	156.9	43.8	f
4.42	6.4	42.8	14.5	63	20.31	58.9	13.46	39.03	149	45.1	f
4.32	4.3	41.6	14	177	26.73	35.2	6.46	91	190.4	96.9	m
4.73	6.7	42.8	14.9	8	19.81	41.8	7.19	70	195.2	75.5	m
4.71	7.2	43.6	14	32	20.39	30.5	5.63	67	186.6	71	m
4.93	7.3	46.2	15.1	41	21.12	34	6.59	67	184.4	71.8	m
5.21	7.5	47.5	16.5	20	21.89	46.7	9.5	70	187.3	76.8	m
5.09	8.9	46.3	15.4	44	29.97	71.1	13.97	88	185.1	102.7	m
5.11	9.6	48.2	16.7	103	27.39	65.9	11.66	83	185.5	94.2	m
4.94	6.3	45.7	15.5	50	23.11	34.3	6.43	74	184.9	79	m
4.86	3.9	44.9	15.4	73	22.83	34.5	6.56	70	181	74.8	m
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4.62	7.3	43.8	14.7	26	21.2	76.8	18.08	61.85	188.7	75.5	f

Pair plots of the variables

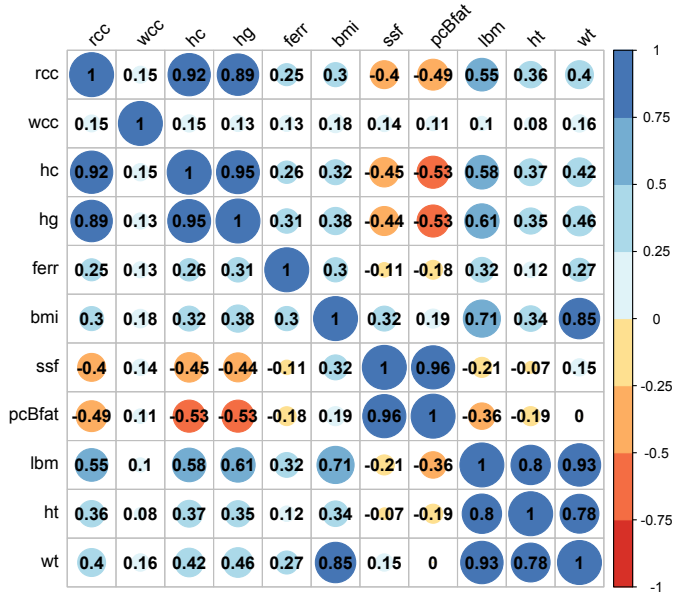
Bivariate representation

→ some variables are linearly dependent

→ their correlation is "high"

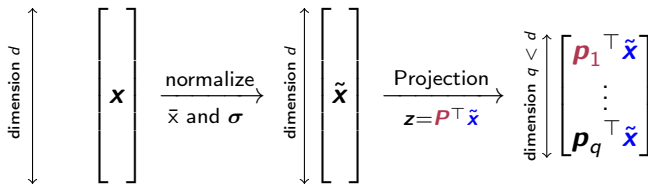


Correlation matrix



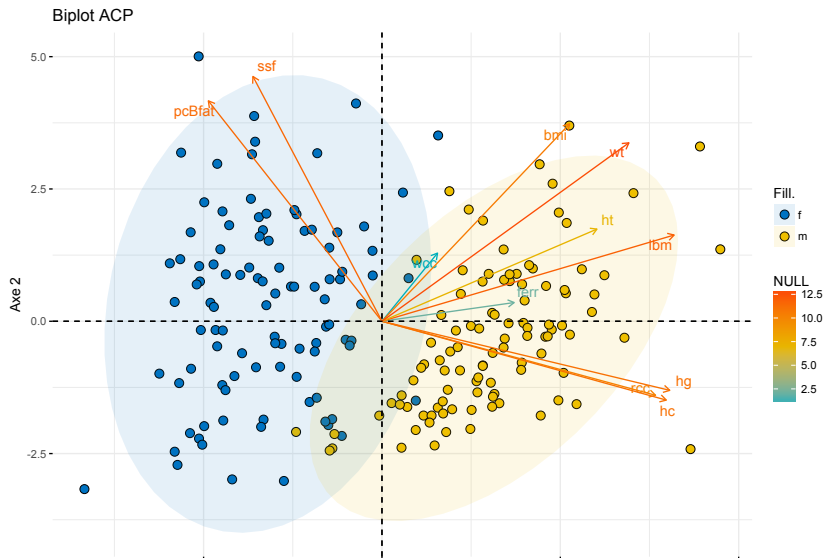
Dimension reduction

Knowing $\mathbf{P} \in \mathbb{R}^{d \times q}$, the mean $\bar{\mathbf{x}}$ and std σ

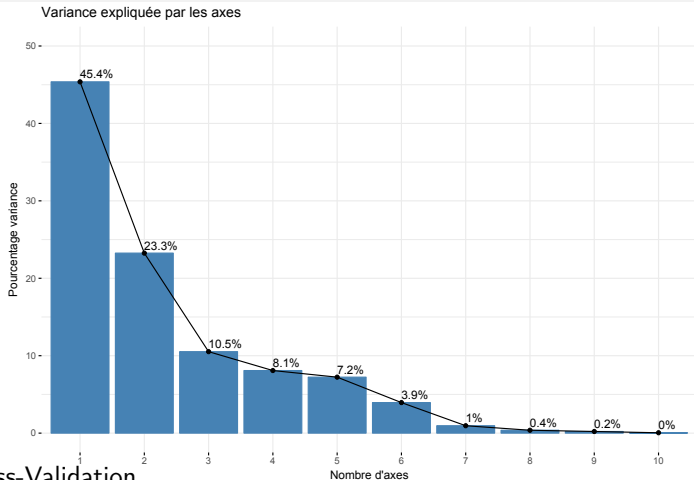


→ we achieve dimensionality reduction

Data visualization ($q = 2$)

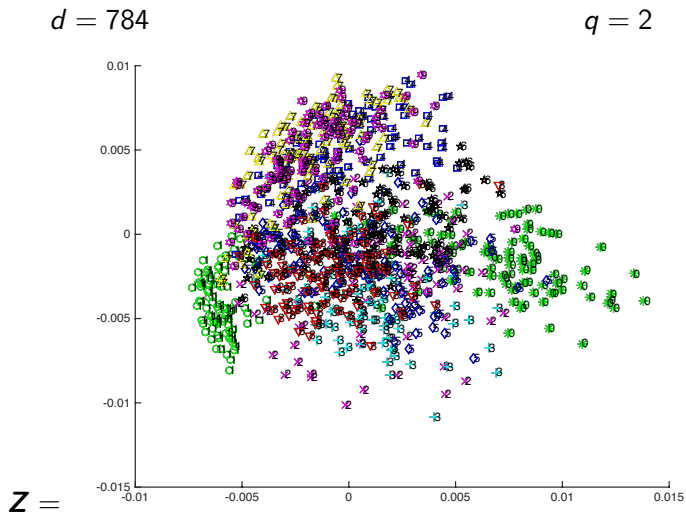


How to choose q ?



- Cross-Validation
- "Elbow trick" on the graph of eigenvalues
- Set a proportion (for instance 95%) of the recovered variance

Visualizing Mnist dataset



Drawbacks of PCA

- Only **linear projection**
- PCA solely relies on order 2 statistics (mean and variance)

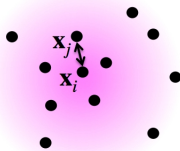
Beyond PCA

- Non-linear PCA
- ISOMAP, LLE, MVU, SNE, t-SNE ...
- Neural networks
 - auto-encoders
 - embeddings for custom data: word2vec, doc2vec (text), signal2vec (time series)

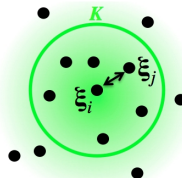
SNE (Stochastic Neighbor Embedding) and t-SNE

Intuition

- Transform pairwise distances $dist(\mathbf{x}_i, \mathbf{x}_j)$ into probability $\mathbb{P}_X(\mathbf{x}_i|\mathbf{x}_j)$ (that \mathbf{x}_i and \mathbf{x}_j are close)
 - low distance \rightarrow high probability to be close
- Same for the projections: $dist(\mathbf{z}_i, \mathbf{z}_j) \rightarrow \mathbb{P}_Z(\mathbf{z}_i|\mathbf{z}_j)$
- Find $\{\mathbf{z}_i\}$ that minimize the distance between the distributions \mathbb{P}_X and \mathbb{P}_Z



$$dist(\mathbf{x}_i, \mathbf{x}_j) = \|\mathbf{x}_i - \mathbf{x}_j\|_2$$



$$\delta_{ij} = \|\xi_i - \xi_j\|_2$$

SNE: the maths

for $\{\mathbf{x}_i\}_{i=1}^n$

- define the probability that \mathbf{x}_j and \mathbf{x}_i are close

$$\mathbb{P}_X(\mathbf{x}_i|\mathbf{x}_j) = \frac{\exp^{-d_{ij}^2}}{\sum_{k=1}^N \sum_{l \neq k} \exp^{-d_{kl}^2}}$$

with
$$d_{ij} = \frac{\text{dist}(\mathbf{x}_i, \mathbf{x}_j)^2}{2\sigma_i^2}$$

σ_i defines the number of neighbors of sample \mathbf{x}_i . It is selected such that

$$\log(K) = -\sum_{j=1}^N \mathbb{P}_X(\mathbf{x}_j|\mathbf{x}_j) \log \mathbb{P}_X(\mathbf{x}_j|\mathbf{x}_j)$$

for $\{\mathbf{z}_i\}_{i=1}^n$ (the unknowns)

- Prob. that \mathbf{z}_j is neighbor of \mathbf{z}_i

$$\mathbb{P}_Z(\mathbf{z}_i|\mathbf{z}_j) = \frac{\exp^{-\delta_{ij}^2}}{\sum_{k=1}^N \sum_{l \neq k} \exp^{-\delta_{kl}^2}}$$

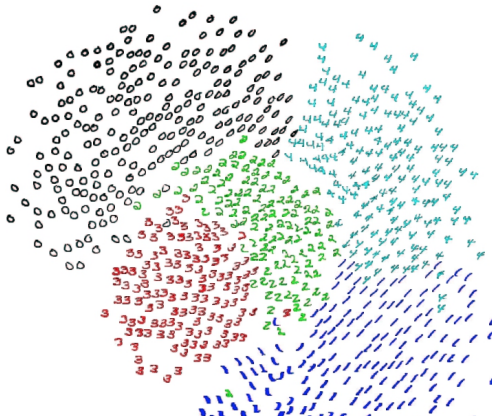
with
$$\delta_{ij} = \|\mathbf{z}_i - \mathbf{z}_j\|$$

Computing the $\{\mathbf{z}_i\}_{i=1}^n$

- Minimize the Kullback-Leibler divergence between \mathbb{P}_X et \mathbb{P}_Z
- $\min_{\mathbf{z}_1, \dots, \mathbf{z}_N} \sum_{i,j=1}^N \mathbb{P}_X(\mathbf{x}_i|\mathbf{x}_j) \log \frac{\mathbb{P}_X(\mathbf{x}_i|\mathbf{x}_j)}{\mathbb{P}_Z(\mathbf{z}_i|\mathbf{z}_j)}$
- Solution via numerical methods

Illustration of SNE

$$X = \begin{pmatrix} 8 & 2 & 5 & 1 & 2 & 4 & 1 & 9 & 7 \\ 1 & 8 & 9 & 2 & 4 & 7 & 7 & 6 & 0 & 1 \\ 6 & 9 & 0 & 0 & 9 & 8 & 1 & 2 & 2 & 6 \\ 3 & 6 & 5 & 7 & 2 & 4 & 4 & 4 & 0 & 3 \\ 9 & 4 & 4 & 8 & 3 & 3 & 3 & 7 & 0 & 7 \\ 7 & 2 & 4 & 3 & 9 & 9 & 6 & 1 & 2 & 8 \\ 7 & 6 & 0 & 0 & 3 & 9 & 8 & 9 & 7 & 4 \\ 9 & 5 & 1 & 8 & 5 & 9 & 9 & 5 & 0 & 9 \\ 5 & 8 & 9 & 0 & 6 & 7 & 3 & 3 & 0 & 1 \\ 0 & 1 & 2 & 1 & 4 & 0 & 1 & 8 & 0 & 2 \end{pmatrix} \quad Z =$$



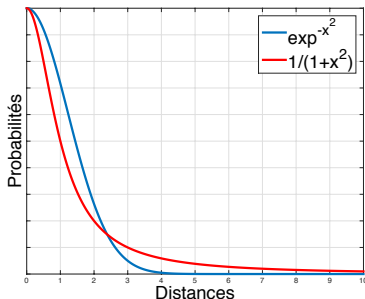
t-SNE variant

SNE

Prob. that \mathbf{z}_j is neighbor of \mathbf{z}_i

$$\mathbb{P}_Z(\mathbf{z}_i|\mathbf{z}_j) = \frac{\exp^{-\delta_{ij}^2}}{\sum_{k=1}^N \sum_{l \neq k} \exp^{-\delta_{lk}^2}}$$

with $\delta_{ij} = \|\mathbf{z}_i - \mathbf{z}_j\|$



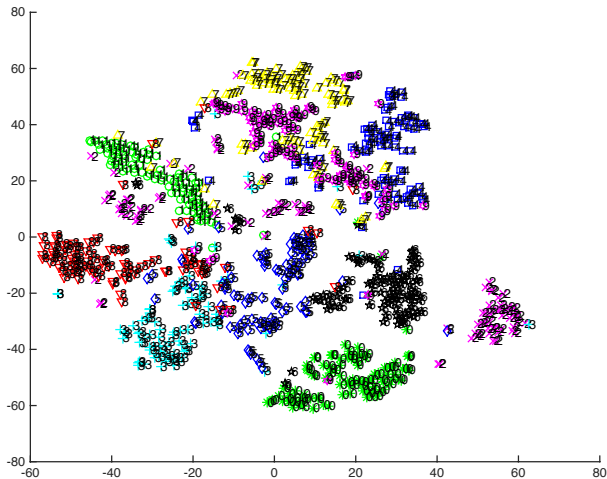
t-SNE

Prob. that \mathbf{z}_j is neighbor of \mathbf{z}_i

$$\mathbb{P}_Z(\mathbf{z}_i|\mathbf{z}_j) = \frac{(1 + \delta_{ij}^2)^{-1}}{\sum_{k=1}^N \sum_{l \neq k} (1 + \delta_{lk}^2)^{-1}}$$

- $\mathbb{P}_x = \mathbb{P}_Z$ large $\Rightarrow \delta_Z < d_X$ (attraction)
- $\mathbb{P}_x = \mathbb{P}_Z$ low $\Rightarrow \delta_Z > d_X$ (repulsion)

Illustration of t-SNE



<https://lvdmaaten.github.io/tsne/>

Conclusions

- PCA: linear dimensionality reduction method
- Several non-linear methods (t-SNE, UMAP, auto-encoder ...)
- They involve advanced optimization methods
- Useful for data visualization and dimension reduction
- Some toolboxes
 - Matlab : <https://lvdmaaten.github.io/drtoolbox/>
 - Python : <http://scikit-learn.org/stable/modules/manifold.html#manifold>
 - Graphical tool : <http://divvy.ucsd.edu/>