Dimensionality reduction

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Summer school on Machine Learning in High Energy Physics
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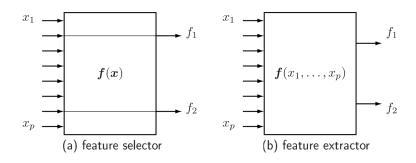
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Definition

Feature selection / Feature extraction



Dimensionality reduction: find transformation of original data to low dimensional space which extracts most relevant information for machine learning task.

Applications

Applications:

- visualization in 2D or 3D
- reduce operational costs (less memory, disc, CPU usage on data transfer)
- remove multi-collinearity to improve performance of machine-learning models

Categorization

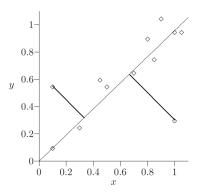
Supervision in dimensionality reduction:

- supervised (such as LDA)
- unsupervied

Mapping to reduced space:

- linear
- non-linear

Example: line of best fit



- In PCA sum of squared of perpendicular distances to line is minimized.
- Not invariant to scale

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Definition

Linear transformation of data:

$$\xi_i = \sum_{j=1}^{D} a_{ij} x_j, \quad i = 1, 2, ... \tilde{D}.$$

Three equivalent ways to derive PCA:

- Find orthogonal transform A yielding new variables ξ_i having stationary values for their variance
- ullet Find orthogonal transform, yielding uncorrelated ξ_j
- Find line of best fit, plane of best fit, etc. where fit is the sum of squares of perpendicular distances.

Method works for $\mathbb{E}x = 0$. So sample observations should be shifted to guarantee: $\sum_{n=1}^{N} x_{nd} = 0$, d = 1, 2, ... D.

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Derivation

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Covariance matrix properties

 $\Sigma = cov[x] \in \mathbb{R}^{D \times D}$ is symmetric positive semidefinite matrix

- has $\lambda_1, \lambda_2, ... \lambda_D$ eigenvalues, satisfying: $\lambda_i \in \mathbb{R}, \ \lambda_i \geq 0$.
- if eigenvalues are unique, corresponding eigenvectors are also unique
- always exists a set of orthogonal eigenvectors $z_1, z_2, ... z_D$: $\sum z_i = \lambda_i z_i$.

later we will assume that $\lambda_1 \geq \lambda_2 \geq ... \geq \lambda_D \geq 0$.

Derivation: 1st component

Consider first component:

$$\xi_1 = \sum_{j=1}^D a_{1j} x_j$$

Optimization problem:

$$\begin{cases} \operatorname{Var} \xi_1 \to \mathsf{max}_{\mathsf{a}} \\ |a_1|^2 = a_1^{\mathsf{T}} a_1 = 1 \end{cases}$$

Variance is equal:

$$Var[\xi_1] = E[\xi_1^2] - (E\xi_1)^2 = E[a_1^T x x^T a_1] - E[a_1^T x] E[x^T a_1]$$

= $a_1^T (E[x x^T] - E[x] E[x^T]) a_1 = a_1^T \Sigma a_1$

Derivation: 1st component

Optimization problem is equivalent to finding unconditional stationary value of

$$\textit{L}(\textit{a}_1, \nu) = \textit{a}_1^{\textit{T}} \Sigma \textit{a}_1 - \nu (\textit{a}_1^{\textit{T}} \textit{a}_1 - 1) \rightarrow \textit{extr}_{\textit{a}_1, \nu}$$

$$\frac{\partial L}{\partial a_1} = 0 : 2\Sigma a_1 - \nu a_1 = 0$$

 a_1 is selected from a set of eigenvectors of A. Since

$$Var[\xi_1] = a_1^T \Sigma a_1 = \lambda_i a_1^T a_1$$

 a_1 is the eigenvector, corresponding to largest eigenvalue λ_i . Eigenvector is not unique if λ_{max} is a repeated root of characteristic equation: $|\Sigma - \nu I| = 0$.

Derivation: 2nd component

$$\xi_2 = a_2^T x$$

$$\begin{cases} \operatorname{Var}[\xi_2] = a_2^T \Sigma a_2 \to \operatorname{max}_{a_2} \\ a_2^T a_2 = |a_2|^2 = 1 \\ \operatorname{cov}[\xi_1, \xi_2] = a_2^T \Sigma a_1 = \lambda_1 a_2^T a_1 = 0 \end{cases}$$

Lagrangian (assuming $\lambda_1>0$)

$$L(a_2, \nu, \eta) = a_2^T \Sigma a_2 - \nu (a_2^T a_2 - 1) - \eta a_2^T a_1 \rightarrow extr_{a_2, \nu, \eta}$$
$$\frac{\partial L}{\partial a_2} = 0 : 2\Sigma a_2 - 2\nu a_2 - \eta a_1 = 0$$
$$a_1^T \frac{\partial L}{\partial a_2} = 2a_1^T \Sigma a_2 - \eta = 0$$

Derivation: 2nd component

Since $a_1^T \Sigma a_2 = a_2^T \Sigma a_1 = 0$, we obtain $\eta = 0$. Then we have that:

$$\Sigma a_2 = \nu a_2$$

so a_2 is eigenvector of Σ , and since we maximize

$$\operatorname{Var}[\xi_2] = a_2^T \Sigma a_2 = \lambda_i a_2^T a_2$$

this should be eigenvector, corresponding to second largest eigenvalue λ_2 .

Derivation: k-th component

$$\begin{aligned} \xi_k &= a_k^T \, \mathbf{x} \\ \begin{cases} Var[\xi_k] &= a_k^T \boldsymbol{\Sigma} a_k \to \mathsf{max}_{a_k} \\ a_k^T a_k &= |a_k|^2 = 1 \\ cov[\xi_k, \xi_j] &= a_k^T \boldsymbol{\Sigma} a_j = \lambda_j a_k^T a_j = 0, \quad j = 1, 2, ... k - 1. \end{cases} \\ \mathsf{Lagrangian} \text{ (assuming } \lambda_j > 0, \ j = 1, 2, ... k - 1) \\ L(a_k, \nu, \eta) &= a_k^T \boldsymbol{\Sigma} a_k - \nu (a_k^T a_k - 1) - \sum_{i=1}^{k-1} \eta_i a_k^T a_i \to \mathsf{extr}_{a_2, \nu, \eta} \\ \frac{\partial L}{\partial a_k} &= 0: \ 2\boldsymbol{\Sigma} a_k - 2\nu a_k - \sum_{i=1}^{k-1} \eta_i a_i = 0 \\ \forall j = 1, 2, ... k - 1: \ a_j^T \frac{\partial L}{\partial a_2} &= 2a_j^T \boldsymbol{\Sigma} a_k - \eta_j = 0 \end{cases}$$

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Derivation: k-th component

Since
$$a_1^T \Sigma a_k = a_k^T \Sigma a_1 = 0$$
, we obtain $\eta_j = 0$ for all $j = 1, 2, ..., k-1$, so

$$\Sigma a_k = \nu a_k$$

 a_k is then the eigenvector.

Variance of ξ_i is

$$Var[\xi_k] = a_k^T \Sigma a_k = \lambda_i a_k^T a_k = \lambda_i$$

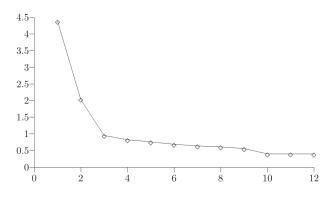
so a_k should be the eigenvector corresponding to the k-th largest eigenvalue λ_k .

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Number of components

- Data visualization: 2 or 3 components.
- Take most significant components until their variance falls sharply down:



Number of components

Remind that $A = [a_1|a_2|...|a_D], A^T A = I, \xi = A^T x.$ Denote $S_k = [\xi_1, \xi_2, ... \xi_k, 0, 0, ..., 0] \in \mathbb{R}^D$

$$\mathbb{E}[\|S_k\|^2] = \mathbb{E}[\xi_1^2 + \xi_2^2 + \dots + \xi_k^2] = \sum_{i=1}^k \operatorname{var} \xi_i = \sum_{i=1}^k \lambda_i$$

$$\mathbb{E}[\|S_D\|^2] = \mathbb{E}[\xi^T \xi] =$$

$$= \mathbb{E}x^T A A^T x = \mathbb{E}\left[x^T x\right] = \mathbb{E}[\|x\|^2]$$

Select such k^* that

$$\frac{\mathbb{E}[\|S_k\|^2]}{\mathbb{E}[\|x\|^2]} = \frac{\mathbb{E}[\|S_k\|^2]}{\mathbb{E}[\|S_D\|^2]} = \frac{\sum_{i=1}^k \lambda_i}{\sum_{i=1}^D \lambda_i} > threshold$$

We may select k^* to account for 90%, 95% or 99% of total variance.

Transformation $\xi \rightleftharpoons x$

Dependence between original and transformed features:

$$\xi = A^T(x-\mu), \ x = A\xi + \mu, \quad \mu$$
 is the mean of original non-shifted data

Taking first r components - $A_r = [a_1|a_2|...|a_r]$, we get the image of the reduced transformation:

$$\xi_r = A_r^T (x - \mu)$$

 ξ_r will correspond to

$$x_r = A \begin{pmatrix} \xi_r \\ 0 \end{pmatrix} + \mu = A_r \xi_r + \mu$$

$$x_r = A_r A_r^T (x - \mu) + \mu$$

 $A_r A_r^T$ is projection matrix with rank r.

Properties of PCA

- Covariance matrix replaced with sample-covariance.
- Depends on scaling (unit deviation transforms covariance matrix to correlation matrix).
- Does not require distribution assumptions
- ullet Eigenvectors may be obtained from SVD decomposition of X.

Application details

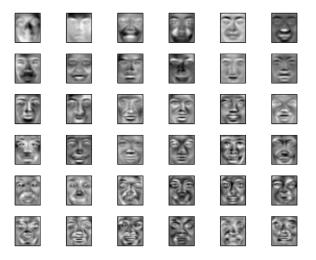
Example

Faces database:



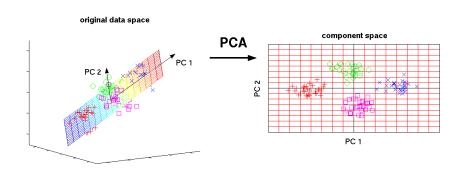
Application details

Eigenfaces



PCA for visualization

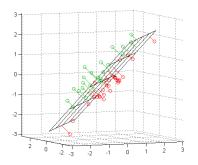
Uncorrelatedness does not imply independence.



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Principal component analysis

Application details

Best hyperplane fit



Subspace L_k or rank k best fits points $x_1, x_2, ... x_D$ if sum of squared distances of these points to this plane is maximized over all planes of rank k.

Best hyperplane fit

For point x_i denote p_i the projection on plane L_k and h_i orthogonal component. Then $\|x_i\|^2 = \|p_i\|^2 + \|h_i\|^2$. For set of points:

$$\sum_{i} \|x_{i}\|^{2} = \sum_{i} \|p_{i}\|^{2} + \sum_{i} \|h_{i}\|^{2}$$

Since sum of squares is constant, minimization of $\sum_i \|h_i\|^2$ is equivalent to maximization of $\sum_i \|p_i\|^2$.

Another view on PCA directions

k-th step optimization problem for $\xi_k = a_k^T x$:

$$\begin{cases} \mathit{Var}[\xi_k] = a_k^T \Sigma a_k \to \max_{a_k} \\ a_k^T a_k = |a_k|^2 = 1 \\ \mathit{cov}[\xi_k, \xi_j] = a_k^T \Sigma a_j = \lambda_j a_k^T a_j = 0, \quad j = 1, 2, ... k - 1. \end{cases}$$

can be equivalently represented as:

$$\begin{cases} \|Xa_k\| \to \max_{a_k} \\ \|a_k\| = 1 \\ a_k \perp a_1, a_k \perp a_2, \dots a_k \perp a_{k-1} \text{ if } k \geq 2 \end{cases}$$
 (1)

since maximization of $\|Xa_k\|$ is equivalent to maximization of $\frac{1}{N} \|Xa_k\|^2 = \frac{1}{N} (Xa_k)^T (Xa_k) = \frac{1}{N} a_k^T X^T X a_k = a_k^T \Sigma a_k$.

Property of PCA

Theorem 1

For $1 \le k \le r$ let L_r be the subspace spanned by $a_1, a_2, ... a_r$. Then for each k L_k is the best-fit k-dimensional subspace for X.

Proof: use induction. For r=1 the statement is true by definition since projection maximization is equivalent to distance minimization.

Suppose theorem holds for r-1. Let L_r be the plane of best-fit of dimension with dim L=r. We can always choose a orthonormal basis of L_r b_1 , b_2 , ... b_r so that

$$\begin{cases} ||b_r|| = 1 \\ b_r \perp a_1, b_r \perp a_2, ... b_r \perp a_{r-1} \end{cases}$$

by setting b_r perpendicular to projections of $a_1, a_2, ... a_{r-1}$ on L_r .

Property of PCA

Consider the sum of squared projections:

$$||Xb_1||^2 + ||Xb_2||^2 + ... + ||Xb_{r-1}||^2 + ||Xb_r||^2$$

By induction proposition:

$$||Xb_1||^2 + ||Xb_2||^2 + ... + ||Xb_{r-1}||^2 \le ||Xa_1||^2 + ||Xa_2||^2 + ... + ||Xa_{r-1}||^2$$

and

$$||Xb_r|| \leq ||Xa_r||$$

since b_r by (28) satisfies constraints of optimization problem (27) and a_r is its optimal solution.

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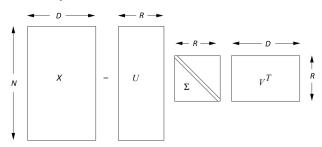
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SVD decomosition

Every matrix $X \in \mathbb{R}^{N \times D}$ of rank R can be decomposed into the product of three matrices:

$$X = U\Sigma V^T$$

where $U \in \mathbb{R}^{N \times D}$, $\Sigma \in \mathbb{R}^{R \times R}$, $V \in \mathbb{R}^{R \times D}$, and $\Sigma = diag\{\sigma_1, \sigma_2, ... \sigma_R\}$, $\sigma_1 \geq \sigma_2 \geq ... \geq \sigma_R \geq 0$, $U^T U = I$, $V^T V = I$. I denotes identity matrix.



Applications of SVD

For square matrix X:

- U, V^T represent rotations, Σ represents scaling (with projection and reflection),
 every square matrix may be represented as superposition of rotation, scaling and another rotation.
- For full rank X:

$$X^{-1} = V \Sigma^{-1} U^T,$$

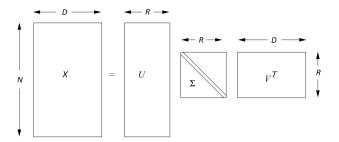
since
$$XX^{-1} = U\Sigma V^T V\Sigma^{-1} U^T = I$$
.

• For not-full-rank X: generalized inverse matrix equals

$$X^+ = V \Sigma^+ U^T$$

 (A^+) is such matrix that $AA^+A=A$

Interpretation of SVD



For X_{ij} let i denote objects and j denote properties.

- U represents standardized coordinates of concepts
- V^T represents standardized coordinates of concepts representations
- \bullet Σ shows the magnitude of standardized concept presence in X.

Example

	The lord of the rings	Harry Potter	Avatar	Titanic	Love story	A walk to remember
Andrew	4	5	5	0	0	0
Andrew John	4	5 4	5 5	0	0	0
John	4	4	5	0	0	0
John Matthew	4 5	4 5	5 4	0	0	0

Example

$$U = \begin{pmatrix} 0. & 0.6 & -0.3 & 0. & 0. & -0.8 \\ 0. & 0.5 & -0.5 & 0. & 0. & 0.6 \\ 0. & 0.6 & 0.8 & 0. & 0. & 0.2 \\ 0.6 & 0. & 0. & -0.8 & -0.2 & 0. \\ 0.6 & 0. & 0. & 0.2 & 0.8 & 0. \\ 0.5 & 0. & 0. & 0.6 & -0.6 & 0. \end{pmatrix}$$

$$\Sigma = \text{diag}\{(14. 13.7 1.2 0.6 0.6 0.5)\}$$

$$V^{T} = \begin{pmatrix} 0. & 0. & 0. & 0.6 & 0.6 & 0.5 \\ 0.5 & 0.6 & 0.6 & 0. & 0. & 0. \\ 0.5 & 0.3 & -0.8 & 0. & 0. & 0. \\ 0. & 0. & 0. & -0.2 & 0.8 & -0.6 \\ -0. & -0. & -0. & 0.8 & -0.2 & -0.6 \\ 0.6 & -0.8 & 0.2 & 0. & 0. & 0. \end{pmatrix}$$

Example (excluded insignificant concepts)

$$U_2 = \begin{pmatrix} 0. & 0.6 \\ 0. & 0.5 \\ 0. & 0.6 \\ 0.6 & 0. \\ 0.6 & 0. \\ 0.5 & 0. \end{pmatrix}$$

$$\Sigma_2 = \mathsf{diag}\{ \begin{pmatrix} 14. & 13.7 \end{pmatrix} \}$$

$$V_2^T = \begin{pmatrix} 0. & 0. & 0. & 0.6 & 0.6 & 0.5 \\ 0.5 & 0.6 & 0.6 & 0. & 0. & 0. \end{pmatrix}$$

Concepts may be

- patterns among movies (along j) fantasy/romance
- patterns among people (along i) boys/girls

Dimensionality reduction case: patterns along j axis.

Applications

• Example: new movie rating by new person

$$x = (5 \ 0 \ 0 \ 0 \ 0 \ 0)$$

• Dimensionality reduction: map x into concept space:

$$y = V_2^T x = (0 \ 2.7)$$

• **Recommendation system:** map y back to original movies space:

$$\hat{x} = yV_2^T = \begin{pmatrix} 1.5 & 1.6 & 1.6 & 0 & 0 \end{pmatrix}$$

Fronebius norm

- Fronebius norm of matrix X is $\|X\|_F \stackrel{df}{=} \sqrt{\sum_{n=1}^N \sum_{d=1}^D x_{nd}^2}$
- Using properties $||X||_F = \operatorname{tr} XX^T$ and $\operatorname{tr} AB = \operatorname{tr} BA$, we obtain:

$$||X||_{F} = \operatorname{tr}[U\Sigma V^{T}V\Sigma U^{T}] = \operatorname{tr}[U\Sigma^{2}U^{T}] =$$

$$= \operatorname{tr}[\Sigma^{2}U^{T}U] = \operatorname{tr}[\Sigma^{2}] = \sum_{r=1}^{R} \sigma_{r}^{2}$$
(2)

Matrix approximation

Consider approximation $X_k = U\Sigma_k V^T$, where $\Sigma_k = \text{diag}\{\sigma_1, \sigma_2, ... \sigma_k, 0, 0, ..., 0\} \in \mathbb{R}^{R \times R}$.

Theorem 2

 X_k is the best approximation of X retaining k concepts.

Proof: consider matrix $Y_k = U\Sigma'V^T$, where Σ' is equal to Σ except some R-k elements set to zero:

$$\sigma'_{i_1}=\sigma'_{i_2}=...=\sigma'_{i_{R-k}}=$$
 0. Then, using (2)

$$\|X - Y_k\|_F = \|U(\Sigma - \Sigma')V^T\|_F = \sum_{p=1}^{R-k} \sigma_{i_p}^2 \le \sum_{p=1}^{R-k} \sigma_p^2 = \|X - X_k\|_F$$

since $\sigma_1 \geq \sigma_2 \geq ... \geq \sigma_R \geq 0$.

Matrix approximation

How many components to retain?

General case: Since

$$||X - X_k||_F = \left\| U(\Sigma - \Sigma_k) V^T \right\|_F = \sum_{i=k+1}^R \sigma_i^2$$

a reasonable choice is k^* such that

$$\frac{\|X - X_{k^*}\|_F}{\|X\|_F} = \frac{\sum_{i=k^*+1}^R \sigma_i^2}{\sum_{i=1}^R \sigma_i^2} < threshold$$

Visualization: 2 or 3 components.

Theorem 3

For any matrix Y_k with rank $Y_k = k \colon \|X - X_k\|_F \le \|X - Y_k\|_F$

Finding U and V

• Finding V $X^TX = (U\Sigma V^T)^T U\Sigma V^T = (V\Sigma U^T)U\Sigma V^T = V\Sigma^2 V^T.$ It follows that

$$X^T X V = V \Sigma^2 V^T V = V \Sigma^2$$

So V consists of eigenvectors of X^TX with corresponding eignvalues $\sigma_1^2, \sigma_2^2, ... \sigma_R^2$.

• Finding *U*:

$$XX^T = U\Sigma V^T (U\Sigma V^T)^T = U\Sigma V^T V\Sigma U^T = U\Sigma^2 U^T$$
. So $XX^T U = U\Sigma^2 U^T U = U\Sigma^2$.

So *U* consists of eigenvectors of XX^T with corresponding eigenvalues $\sigma_1^2, \sigma_2^2, ... \sigma_R^2$.

Comments

- ullet Denote the average $ar{X} \in \mathbb{R}^D: ar{X}_j = \sum_{i=1}^N x_{ij}$
- ullet Denote the n-th row of X be $X_n \in \mathbb{R}^D$: $X_{nj} = x_{nj}$
- For centered X sample covariance matrix $\widehat{\Sigma}$ equals:

$$\widehat{\Sigma} = \frac{1}{N} \sum_{n=1}^{N} (X_n - \bar{X})(X_n - \bar{X})^T = \frac{1}{N} \sum_{n=1}^{N} X_n X_n^T$$
$$= \frac{1}{N} X^T X$$

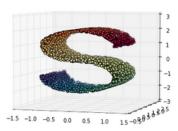
- V consists of principal components since
 - V consists of eigenvectors of X^TX ,
 - ullet principal components are eignevectors of $\widehat{\Sigma}$ and
 - $\widehat{\Sigma} \propto X^T X$

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Non-linear dimensionality reduction

- Based on assumption that original data $x \in \mathbb{R}^D$ is distributed compactly on non-linear surface with dimensionality d < D.
- Let $y \in \mathbb{R}^d$ denote the coordinates of x on the surface.
- d is usually unknown.
- Sample dataset:



• Linear dimensionality reduction techniques will fail here.

Categorization

Non-linear approaches of dimensionality reduction:

- preserving global properties
 - kernel PCA, autoencoders, MDS, ISOMAP, diffusion maps, MVU
- preserving local properties
 - LLE, LTSA
- global alignment of local linear models (not considered here)

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Multi-dimensional scaling

Multi-dimensional scaling

Map $x \rightarrow y$ preserving distances as much as possible.

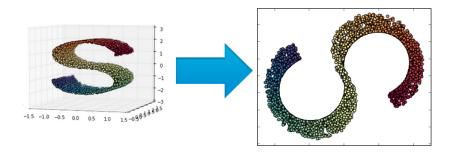
- Approaches:
 - absolute difference

$$\sum_{i,j} (\|x_i - x_j\| - \|y_i - y_j\|)^2 \to \min_{Y}$$

• relative difference (more attention to small distances)

$$\sum_{i,j} \frac{(\|x_i - x_j\| - \|y_i - y_j\|)^2}{\|x_i - x_j\|} \to \min_{Y}$$

Example



Issue: small $||x_i - x_j||$ should not always imply small $||y_i - y_j||$, such as in case of red and yellow points.

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Non-linear dimensionality reduction

Global methods

Isomap

Isomap

Map $x \to y$ preserving correspondence between distance in transformed space and "geodesic" distance along the surface in original space.

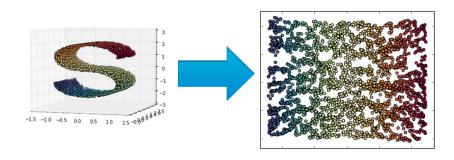
- This apprach solves the previous issue of MDS.
- Geodesic distance calculation:
 - ① for each x_n find its K nearest neighbours $x_{n_1,n_2,...n_K}$
 - Suild the pairwise distance matrix, filling distance between samples and their k-NN.
 - calculate all pairwise distances using shortest-path algorithm of Dijkstra or Floyd.
- Finally usual MDS is applied to match $||x_i x_j||_G$ and $||y_i y_j||$, where $||\cdot||_G$ is geodesic distance.

Issues of Isomap

- Noisy observations between distant parts of surfaces may make distant parts close
- Solutions:
 - remove observations with large total flows through them
 - remove nearest neighbours that violate local linearity
- Non-convexity of surface may lead to taking K-NN that are far away in the transformed space.
- Selection of K:
 - if too small, then poor approximation of geodesic distance
 - if too large, then increases chance of "short-circuiting" through noisy observations.

Global methods

Example of ISOMAP



Maximum variance unfolding

Maximum variance unfolding

Maximally unfold the transformations, preserving local geometry of data.

```
initialize neighbourhood graph G with edges being the samples x_1, x_2, ... x_N
```

for each x_n :

for
$$k = 1, 2, ...K$$
:

find k-th nearest neighbour x_{n_k} to x_n add a link to G between x_n and x_{n_k}

solve the optimization problem:

$$\sum_{i,j} ||y_i - y_j|| \to \min \text{ subject to: } ||y_i - y_j||^2 = ||x_i - x_j|| \ \forall (i,j) \in G$$

Issue: noise sample may "short-circuit" leading to redundant constraint, which may prevent manifold unfolding.

Kernel PCA

- Like PCA, but input space is expanded with kernels
- Easy computation of projections of new points
- Issue: kernel selection.
 - linear (reduces to ordinary PCA)
 - Gaussian
 - polynomial

Diffusion maps

- Construct proximity graph
 - nodes: observations
 - edge weight between x_i and x_j :

$$w_{ij}=e^{-\frac{\left\|x_i-x_j\right\|^2}{2\sigma^2}}$$

2 for each x_i outgoing probabilities set to normalized weights:

$$p_{ij}^{(1)} = \frac{w_{ij}}{\sum_k w_{ik}}$$

- **3** random walk with probabilities $p_{ij}^{(1)}$ stored in matrix $P^{(1)}$ is assumed.
- based on random walk assumption, the probability of walking from x_i to x_j after T steps is:

$$p_{ij}^{(T)} = \left(P^{(1)}\right)^T$$

Diffusion maps

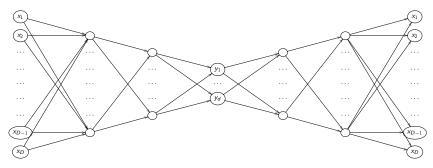
• Finally MDS is applied to match $||y_i - y_j||$ to diffusion distance:

$$D^{T}(x_{i}, x_{j}) = \sqrt{\sum_{k} \frac{(p_{ik}^{(T)} - p_{jk}^{(T)})}{\psi(x_{k})}}$$

where $\psi(x_k)$ is normalization term, that attributes more weight to high density regions.

• Benefit: distance between points is based on multiple paths through the graph - more robust to noise.

Autoencoders



- feed-forward neural network, tranined to reproduce input with MSE loss.
- \bullet D input and D output nodes
- d nodes in the central layer
- User-defined number of layers and nodes

Autoencoders

- Benefits: can map new points to reduced space
- Issues:
 - optimization may get stuck in local optima
 - slow convergence (can be cured with specific starting weights)
 - infeasible to apply to high d (too many connections).

Kitov Victor - Dimensionality reduction Non-linear dimensionality reduction Local methods

- 4 Non-linear dimensionality reduction
 - Global methods
 - Local methods

Local linear embedding

Local linear embedding

Method preserves reconstruction weights of objects through their nearest neighbors.

INPUT:

Local methods

training sample $x_1, x_2, ... x_N$ number of neighbours K

ALGORITHM:

for each x_i :

find its K nearest neighbours: $x_{i(1)}, x_{i(2)}, ... x_{i(K)}$ find weights to reconstruct x_i using its neighbours: $x_i \approx \sum_{k=1}^K w_{ik} x_{i(k)}$

solve optimization problem: $\sum_{n=1}^{N}(y_i-\sum_{k=1}^{K}w_{ik}x_{ik})^2 o \min_Y$

<u>OUTPUT</u>: reduced space representation: $y_1, y_2, ... y_N$.

Laplacian eigenmaps

Laplacian eigenmaps

Method preserves distances of points with nearest neighbors.

INPUT:

training sample $x_1, x_2, ... x_N$ number of neighbours K

ALGORITHM:

for each x_i :

find its K nearest neighbours: $x_{i(1)}, x_{i(2)}, ...x_{i(K)}$ for each nearest neighbbour j=i(1), i(2), ...i(K):

calculate distance-based weights: $w_{ii} = e^{-\frac{\left|\left|x_{i}-x_{j}\right|\right|^{2}}{2\sigma^{2}}}$

solve optimization problem:

$$\sum_{n=1}^{N} \sum_{i=1}^{N} \sum_{j \in \{i(1), \dots, i(K)\}} w_{ij} (y_i - y_j)^2 \to \min_{Y}$$

<u>OUTPUT</u>: reduced space representation: $y_1, y_2, ... y_N$.

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Comments on local methods

- short-circuiting affects only local points in space
- local method, relying on K-NN => prone to curse of dimensionality
- prone to overfitting on outliers (when they become nearest neighbors)

Properties

Technique	Convex	Parameters	Computational	Memory
PCA	yes	none	$O(D^3)$	$O(D^2)$
MDS	yes	none	$O(N^3)$	$O(N^2)$
Isomap	yes	K	$O(N^3)$	$O(N^2)$
MVU	yes	K	$O((NK)^3)$	$O((NK)^3)$
Kernel PCA	yes	kernel	$O(N^3)$	$O(N^2)$
Diffusion maps	yes	σ , T	$O(N^3)$	$O(N^2)$
Autoencoders	no	network shape	O(INW)	O(W)
LLE	yes	K	$O(pN^2)$	$O(pN^2)$
Laplacian eigenmaps	yes	K, σ	$O(pN^2)$	$O(pN^2)$

D - input dimension, N - sample size, K - number of nearest neighbors, σ - smoothing parameter of Gaussian kernel, W number of weights in neural network, I - number of iterations (optimization passes), p - the fraction of non-zero entries in the weight matrix.

Other

- Problem of transforming new previously unobserved samples.
 - direct for PCA, Kernel PCA, autoencoders
 - only approximations possible for other methods.
- Intrinsic dimensionality estimation
 - Cross-validation of the original task (e.g. classification)
 - How many components of local PCA explain most of the variance
 - The growth rate of number of samples falling inside a growing hypersphere with center x:

$$\#\{x_i: \|x_i - x\| \le R\}$$

etc.

Evaluation

- L.J.P. van der Maaten, E.O. Postma, H.J. van den Herik.
 Dimensionality Reduction: A Comparative Review. Working paper. 2008.
 - Extensive comparison of different dimensionality reduction methods.
 - Non-linear techniques perform better than PCA on simulated data
 - PCA wins most of the time on real data
 - Problems:
 - global methods: short-circuiting
 - nearest neighbours based methods: curse of dimensionality, overfitting to outliers
 - unstable optimization for local methods: they reduce to eigenproblems, frequently $\lambda_{max}/\lambda_{min}\gg 1$.
 - suboptimal local optima for autoencoders.
- Actively developing field!

References

Keith D. Copsey Andrew R. Webb. Statistical Pattern Recognition. Wiley, 3rd edition, 2011.

Ravindran Kannan John Hopcroft.
Foundations of data science.
www.cs.cornell.edu/jeh/book11April2014.pdf, 2014.
[Online; accessed 16-August-2015].

Jeff Ullman Jure Leskovec, Anand Rajaraman.
Mining of massive datasets.
www.mmds.org, 2014.
[Online; accessed 16-August-2015].

L.J.P. van der Maaten, E. O. Postma, and H. J. van den Herik. Dimensionality reduction: A comparative review, 2008.