

Dimension Reduction

2018 Statistics Summer School

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- ① An Intuition Introduction for PCA
- ② PCA: A Dimension Reduction Method for vector data
- ③ SVD: A Dimension Reduction Method for a Matrix
- ④ MPCA: A Dimension Reduction Method for Matrix Data

The 'Learning Evolution' of Statistics

- Mean (medium,...) \rightarrow Mean Vector
- Variance \rightarrow Covariance Matrix
- What can you do about the covariance matrix?
- Ans: Eigenvalue Decomposition!

Review of Covariance Matrix

- Let x_1, \dots, x_n be length- p observation vectors
- WLOG, let their mean be length- p 0-vector.
- Let the data matrix $X = (x_1, \dots, x_n)$ be a p by n matrix.
- The sample covariance matrix

$$\Sigma = XX^T / (n - 1) = \sum_{i=1}^n x_i x_i^T / (n - 1).$$

A Proof of $XX^T = \sum_{i=1}^n x_i x_i^T$

- By $X = (x_1, \dots, x_n)$, we have $X_{ij} = (x_j)_i$.

$$\begin{aligned}
 (XX^T)_{jk} &= \sum_{i=1}^n X_{ji} X_{ki} \\
 &= \sum_{i=1}^n (x_i)_j (x_i)_k \\
 &= \sum_{i=1}^n (x_i x_i^T)_{jk} \\
 &= \left(\sum_{i=1}^n x_i x_i^T \right)_{jk}
 \end{aligned}$$

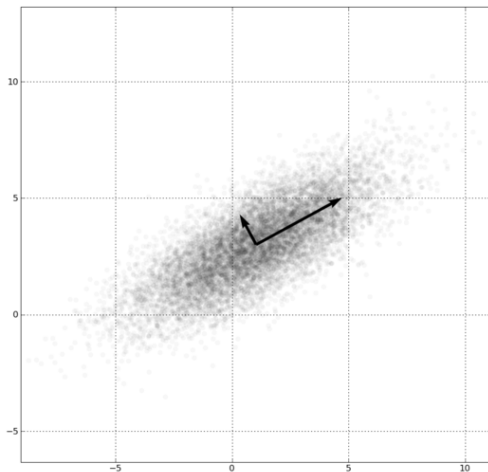
An Intuition Definition for PCA

- PCA: Principal Component Analysis
- The underlying statistical philosophy is "Larger variance captures more information."

An Intuition Definition for PCA

- Find a direction vector ($\in R^p$) p_1 , such that the variance of the data $\{x_i\}_{i \leq n}$ projected to this direction $\{x_i^T p_1\}_{i \leq n}$ has maximum variance.
- Then find p_2 orthonormal to p_1 , such that the variance of $\{x_i^T p_2\}_{i \leq n}$ has maximum variance.
- \dots find p_k orthonormal to p_1, \dots, p_{k-1} , such that the variance of $\{x_i^T p_k\}_{i \leq n}$ has maximum variance.

An Example with $p = 2$



Eigenvalue Decomposition

- Given a symmetric p.d. matrix $\Sigma_{p \times p}$, Σ can be decomposed as

$$\Sigma = \sum_{i=1}^p \lambda_i u_i u_i^t,$$

where u_1, \dots, u_p are orthonormal and $\lambda_1 \geq \dots \geq \lambda_p$. Furthermore, u_i can be uniquely decided up to $+/-$ sign if eigenvalue λ_i is distinct.

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$$\Sigma u_i = \lambda_i u_i$$

-

$$\Sigma U = U \Lambda$$

The Intuition Definition vs Eigenvalue Decomposition

- $p_1 = \operatorname{argmax}_q \sum_{i=1}^n (x_i^T q)^2$
- Will $p_1 = u_1$?

The Intuition Definition vs Eigenvalue Decomposition

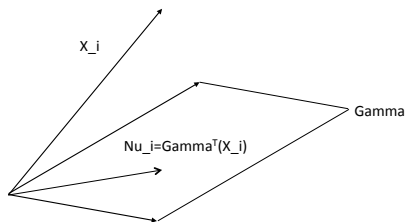
$$\begin{aligned}
 p_1 &= \operatorname{argmax}_{|q|=1} \sum_{i=1}^n (x_i^T q)^2 = \operatorname{argmax}_{|q|=1} \sum_{i=1}^n (q^T x_i)(x_i^T q) \\
 &= \operatorname{argmax}_{|q|=1} \sum_{i=1}^n q^T (x_i x_i^T) q = \operatorname{argmax}_{|q|=1} q^T \left(\sum_{i=1}^n x_i x_i^T \right) q \\
 &= \operatorname{argmax}_{|q|=1} q^T (X X^T) q = \operatorname{argmax}_{|q|=1} q^T \left(\sum_{i=1}^p \lambda_i u_i u_i^T \right) q \\
 &= \operatorname{argmax}_{|q|=1} \sum_{i=1}^p \lambda_i (u_i^T q)^2 \\
 &= u_1
 \end{aligned}$$

PCA Model

- $X \in \mathbb{R}^p$ is a random vector.
- $X = \Gamma\nu + \varepsilon$.
- $\min_{\Gamma \in \mathcal{O}_{p \times p'}, \nu \in \mathbb{R}^{p'}} E[\|X - \Gamma\nu\|_F^2]$.
- Sample version
 - Assume the mean \bar{X} has been taken off.
 - $\min_{\Gamma \in \mathcal{O}_{p \times p'}, \nu_i \in \mathbb{R}^{p'}, i \leq n} \frac{1}{n} \sum_{i=1}^n \|X_i - \Gamma\nu_i\|^2$.

PCA vs Linear Regression: A Geometric Point of View, after Γ is chosen.

Geometric interpretation



Ordinary Least-Squares

PCA Demo: Data Matrix ($p \times n$)

PCA Demo:

Data Matrix vs the first 5 Eigenvectors

Data Matrix



Eigenvectors



PCA Demo: Data Matrix ($p \times n$)

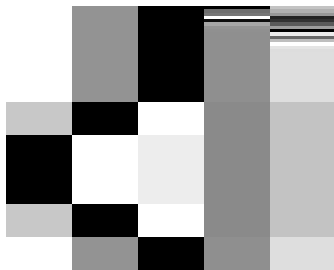
PCA Demo:

Data Matrix vs the first 5 Eigenvectors

Data Matrix



Eigenvectors



What's SVD

- Given a $p \times n$ real value matrix X , $X = UDV^T$,
- U : a $p \times p$ orthonormal matrix,
- V : a $p \times p$ orthonormal matrix,
- D : $p \times q$ and D 's nonzero elements only appear at diagonal with size $\min(p, q)$.

SVD: An Optimization Formula

- Given rank k , SVD can be formulated as an optimization problem.

$$(U_{1:k,\cdot}, D_{1:k,1:k}, V_{1:k,\cdot}) = \underset{A \in \mathcal{O}_{p \times k}, B \in \mathcal{O}_{q \times k}, D \in R^{k \times k}}{\operatorname{argmin}} \|X - ADB^T\|_F^2$$

A Neat Formula of SVD

- Let the column vectors of U and V be u_1, \dots, u_k and v_1, \dots, v_k .
- Let the diagonal elements of D be d_1, \dots, d_k .
- Then, $\mathbf{X} = \mathbf{U}\mathbf{D}\mathbf{V}^T = \sum_{i=1}^k d_i u_i v_i^T$.
- Each $u_i v_i^T$ represents a rank one $p \times q$ matrix.

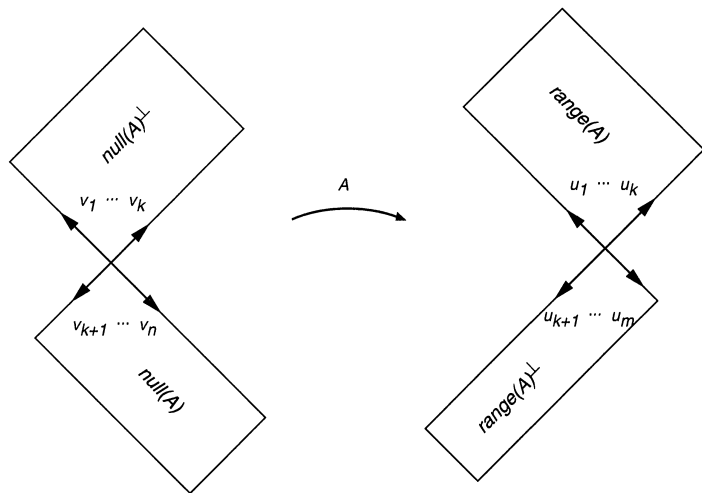
A Transformation Point of View: $\mathbf{X} = \mathbf{UDV}^T$ 

Figure 1. Strang's diagram.

What does SVD say?

Any real value transformation matrix from R^q to R^p can be decomposed to 3 steps:

- A rotation matrix V in R^q .
- A scale matrix D .
- A rotation matrix U in R^p .

A Neat Formula of SVD

- Let the column vectors of U and V be u_1, \dots, u_k and v_1, \dots, v_k .
- Let the diagonal elements of D be d_1, \dots, d_k .
- Then, $\mathbf{X} = \mathbf{U}\mathbf{D}\mathbf{V}^T = \sum_{i=1}^k d_i u_i v_i^T$.
- SVD finds the k vector-pairs with one to one mapping from R^q to R^p .
- Each $u_i v_i^T$ represents a rank one matrix with dimension $p \times q$.

SVD vs PCA

- If by SVD, $X = UDV^T$, where U is an orthonormal matrix such that $UU^T = I_p$
- By PCA, U is the eigenmatrix of $XX^T = UD^2U^T$.
- Furthermore, V is the eigenmatrix of $X^TX = VD^2V^T$.

Application of SVD: Dimension Reduction

- $\mathbf{X}_{p \times q} = \mathbf{U}_{p \times p} \mathbf{D}_{p \times q} \mathbf{V}_{q \times q} \approx \tilde{\mathbf{U}}_{p \times k} \tilde{\mathbf{D}}_{k \times k} \tilde{\mathbf{V}}_{k \times q}^T$
- k : selected rank.

196 X 257



Rank = 100



Another Point of View

$$\mathbf{X} = \mathbf{U}\mathbf{D}\mathbf{V}^T = \sum_{i=1}^k d_i u_i v_i^T$$

- A linear combination of bases: $\{u_i v_i^T, 1 \leq i \leq k\}$.
- Each $u_i v_i^T$ is a rank one matrix.

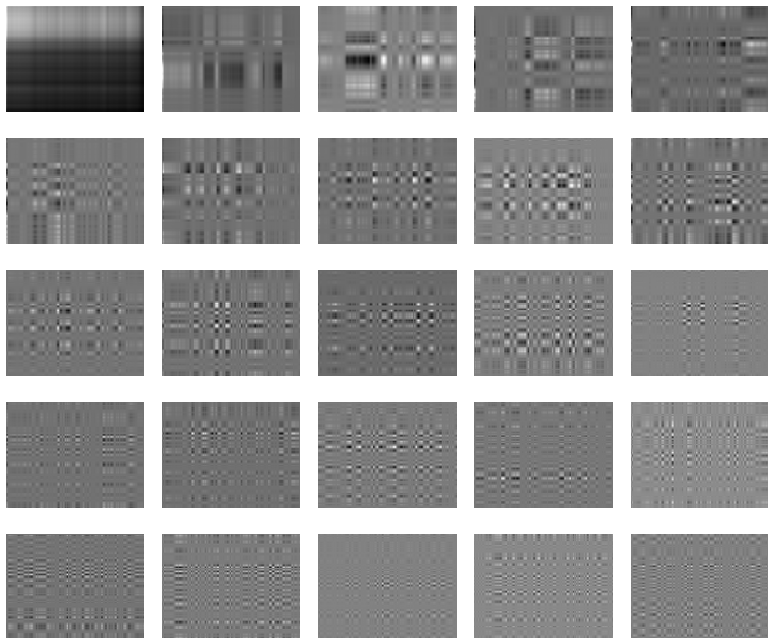


Figure: The first 25 rank-one bases.

SVD Rank Demo: The more, the better.

196 X 257



Rank =25



Rank =50

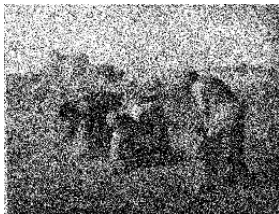


Rank =100

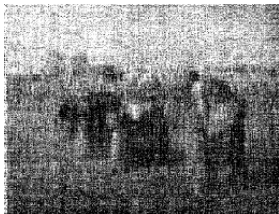


SVD Rank Demo: The more, the better?

196 X 257



Rank =25



Rank =50



Rank =100



Gleaners and Angelus

Angelus



Gleaner



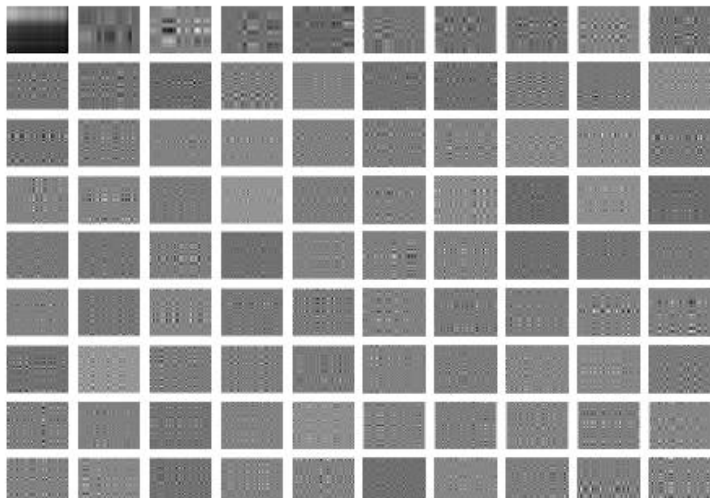
Angelus Reconstructed by Gleaners

Gleaner



Angelus Reconstructed by Gleaner



Gleaners Basis 10×10 

MPCA, HOSVD: multi-matrix versions of SVD

- We have matrix A_1, \dots, A_n , all from R^q to R^p .
- We want to stick on one set of U and V by sacrificing some bias.
- We relax the condition on the diagonal property of D .

Multilinear Principal Component Analysis (MPCA)

- We define **column** basis $A \in \mathcal{O}_{p \times \tilde{p}}$, **row** basis $B \in \mathcal{O}_{q \times \tilde{q}}$.
- SVD: Find A, B (with $\tilde{p} = \tilde{q}$) and a diagonal matrix D that

$$\min_{A, B, D} \|X - ADB^T\|_F^2$$

- MPCA: Find simultaneous A, B , and U_i by

$$\min_{A, B, U_i, 1 \leq i \leq n} \frac{1}{n} \sum_{i=1}^n \|(X_i - \bar{X}) - AU_i B^T\|_F^2$$

- D diagonal and U_i usually not.

Implementation: iterative alternating eigenvalue decompositions(Ye, 2005)

- \hat{A} : leading \tilde{p} eigenvectors of $\hat{\Sigma}_{\hat{B}} = \frac{1}{n} \sum_{i=1}^n (X_i - \bar{X}) P_{\hat{B}} (X_i - \bar{X})^T$.
solve a small eigenvalue problem: a $p \times p$ matrix, where $P_{\hat{B}} = \hat{B} \hat{B}^T$
- \hat{B} : leading \tilde{q} eigenvectors of $\hat{\Sigma}_{\hat{A}} = \frac{1}{n} \sum_{i=1}^n (X_i - \bar{X})^T P_{\hat{A}} (X_i - \bar{X})$.
solve a small eigenvalue problem: a $q \times q$ matrix, where $P_{\hat{A}} = \hat{A} \hat{A}^T$.
- **Iterative alternating until convergence.**

Experimental Setting

- 400 face images of 64×64 : partition them to 100-300 training-test sets.
- Both MPCA and PCA are applied on the training images to produce basis to [reconstruct the test images](#).

Basis from the training set

- MPCA: 24 row and 24 column eigenvectors are used to generate 576 basis (24 is selected by hypothesis test for 95% explained-variation)
- PCA: 576 ($= 24 \times 24$) eigenvectors
- 500 replicates, for random partition into training-test subsets, are performed to compare the mean test error.

	MPCA	PCA
Mean	452	2870
SD	4	43

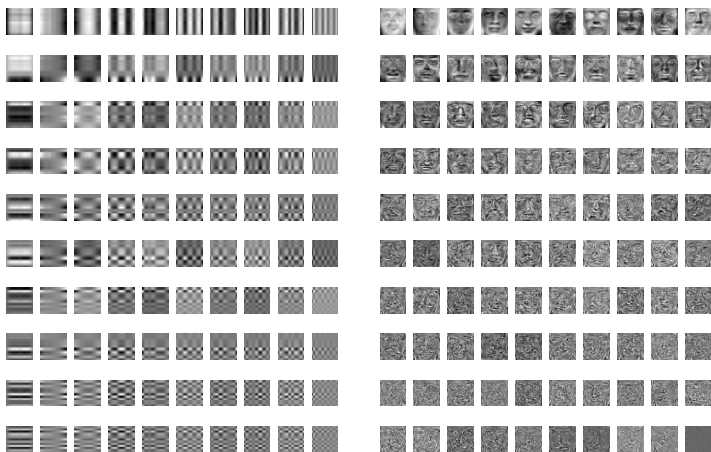
The error is defined as the Frobenius norm for two images.

Visual Comparisons

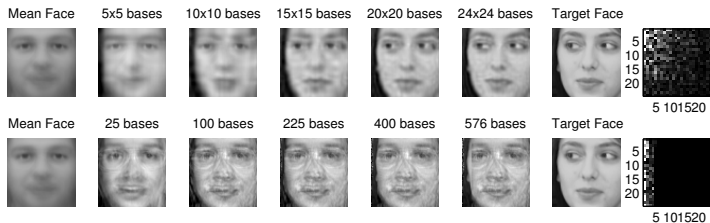


20 test faces randomly drawn (rows 1-2), reconstructions by MPCA (rows 3-4) and PCA (rows 5-6).

Basis Comparisons: MPCA vs PCA

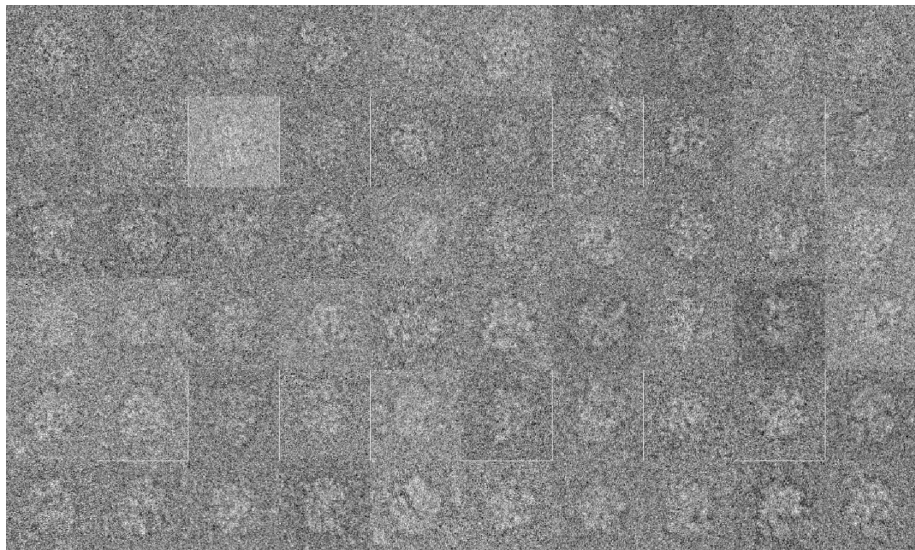


Stepwise Comparison

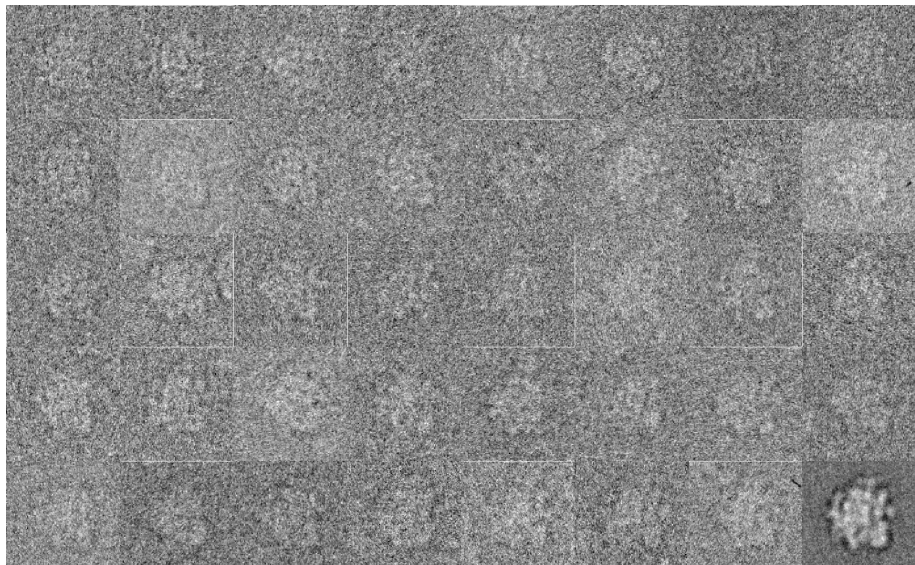


Test image reconstruction, MPCA (top) and PCA (bottom).

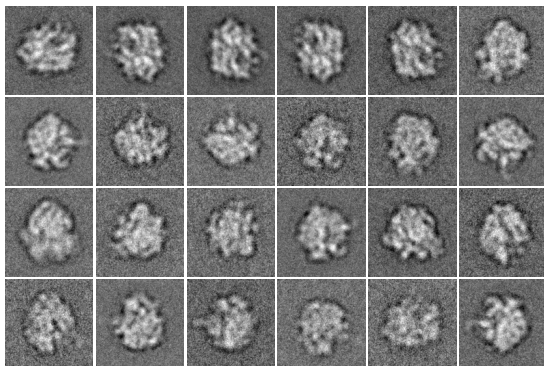
5000 Ribosome cryoEM images



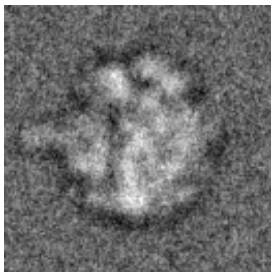
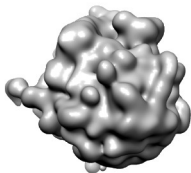
A Cluster Example



24 Cluster Averages for Ribosome Data



One Interesting Example



Summary: Properties and Applications of

- PCA
- SVD
- MPCA