

Machine Learning

Lecture 10: Dimensionality Reduction & Matrix Factorization

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Roadmap

- Chapter: Dimensionality Reduction & Matrix Factorization
 - 1. Introduction
 - 2. Principal Component Analysis (PCA)
 - 3. Singular Value Decomposition (SVD)
 - 4. Matrix Factorization
 - 5. Neighbor Graph Methods
 - 6. Autoencoders (Non-linear Dimensionality Reduction)

Introduction: Unsupervised Learning (I)

- Supervised learning aims to map inputs to targets with y = f(x), or in a probabilistic framework it models p(y|x)
- Unsupervised learning can be seen as modelling p(x)
- We are trying to find the (hidden / latent) structure in the data
 - e.g. find a latent distribution $p(\mathbf{z})$ and a generative transformation $p(\mathbf{x} \mid \mathbf{z})$ we can then obtain $p(\mathbf{x}) = \int p(\mathbf{x} \mid \mathbf{z}) \, p(\mathbf{z}) \, d\mathbf{z}$
 - latent z usually unknown and has to be estimated
- Examples:
 - Clustering: the cluster label is the latent state
 - Anomaly detection: treat instances with low p(x) as anomalies

Introduction: Unsupervised Learning (II)

- Unsupervised learning can be viewed as compression
 - compress a data point to a single label corresponding to its cluster
 - compress a data point from a higher dim. to a lower dim. latent space
- Unsupervised learning can be used ...
 - ... as a stand-alone method (e.g. to understand your data, visualization)
 - ... as a pre-processing step (e.g. use cluster label as feature for subsequent classification task; obtain small number of relevant features)
 - ... to leverage large amounts of unlabeled data for pretraining
- This lecture: Dimensionality Reduction & Matrix Factorization

Dimensionality Reduction: Motivation

- Often data has very many features, i.e. high-dimensional data
- High-dimensional data is challenging:
 - Similarity search/computation is expensive because of high complexity of distance functions
 - Highly correlated dimension could cause trouble for some algorithms
 - Curse of dimensionality: we need exponential amounts of data to characterize the density as the dimensionality goes up
 - It is hard to visualize high-dimensional data
- Often the data lies on a low-dim. manifold, embedded in a high-dim. space
- Goal: Try to reduce the dimensionality while avoiding information loss
- Benefits:
 - Computational or memory savings
 - Uncover the intrinsic dimensionality of the data
 - (more benefits later....)

Feature (Sub-)Selection

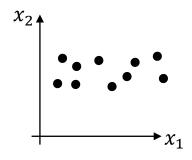
- Choose "good" dimensions using a-priori knowledge or appropriate heuristics
 - e.g. remove low-variance dimensions
 - Depending on the application only a few dimensions might be of interest
 - Example: shoe size interesting for shoe purchases, not so for car purchases

Advantages:

 No need for an intensive preprocessing or training phase to determine relevant dimensions

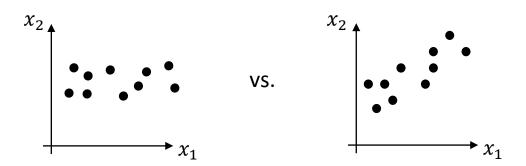


- Expert knowledge required; misjudgment possible
- Univariate feature selection ignores correlations

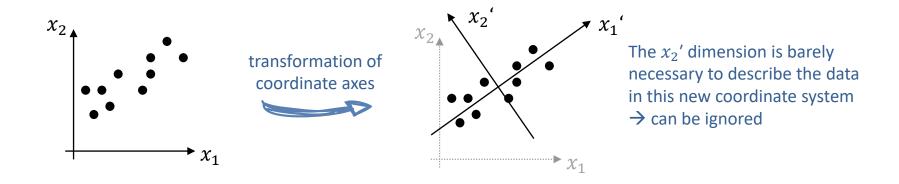


Beyond Feature (Sub-)Selection

- Can we do
 - better?
 - automatic?

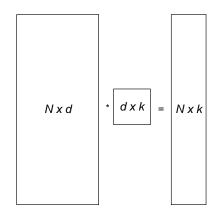


- Obviously: Simply discarding whole features not a good idea
 - Features are often correlated



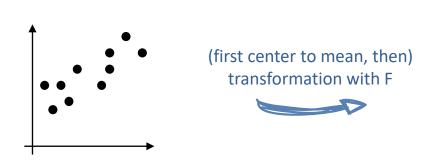
Dim. Reduction via Linear Transformations

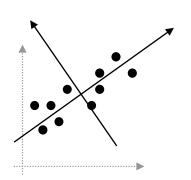
- Represent data in a different coordinate system via linear transformations
 - change of basis (orthogonal basis transformations)
 potentially discarding dimensions



Technical:

- use orthonormal transformation matrix $\mathbf{F} \in \mathbb{R}^{d \times k}$
- $-(x')^T = x^T \cdot F$ is the transformation of (column) vector x into the new coordinate system defined by F
- $X'=X\cdot F$ is the matrix containing all the transformed points x_i'





Discussion: Linear Transformations

- Feature selection is a linear transformation
 - What is the matrix F?
- Let \overline{x} be the mean vector (here: row vector) in the original data space, the mean vector in the transformed space is given by $\overline{x'} = \overline{x} \cdot F$

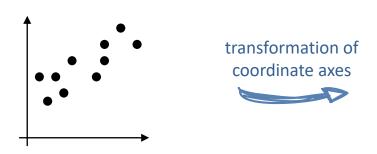
• Let Σ_X be the covariance matrix in the original data space, the covariance matrix in the transformed space is then $\Sigma_{X'} = F^T \cdot \Sigma_X \cdot F$

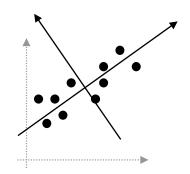
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Principal Component Analysis: Motivation

- Question: Which transformation matrix F to use?
 - Is there an optimal orthogonal transformation (depending on the data)?
 - Optimality: Approximate the data with few coefficients as well as possible

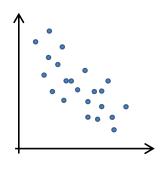




- Approach: Principal Component Analysis (PCA)
 - Find a coordinate system in which the (possibly originally correlated) points are linearly uncorrelated
 - The dimensions with no or low variance can then be ignored

Goal:

- Transform the data, such that the covariance
 between the new dimensions is 0
- The transformed data points are not linearly correlated any more



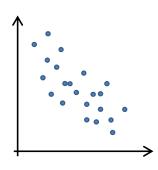
- Given: N d-dimensional data points: $\{m{x}_i\}_{i=1}^N$, $\mbox{m{x}}_i \in \mathbb{R}^d \ orall i \in \{1,\dots,N\}$
- We represent this set of points by a matrix $X \in \mathbb{R}^{N \times d}$:

$$\mathbf{X} = \begin{bmatrix} x_{11} & \cdots & x_{1d} \\ \vdots & \ddots & \vdots \\ x_{N1} & \cdots & x_{Nd} \end{bmatrix}$$

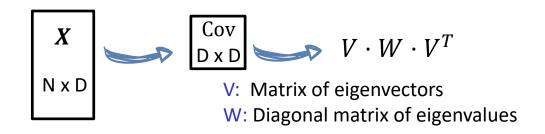
- The row $x_i = \{x_{i1}, ..., x_{id}\} \in \mathbb{R}^d$ denotes the i-th point and the column $X_{:,j}$ denotes the vector containing all values from the j-th dimension

Goal:

- Transform the data, such that the covariance
 between the new dimensions is 0
- The transformed data points are not linearly correlated any more



- General approach
 - Center the data
 - 2. Compute the covariance matrix
 - 3. Use the Eigenvector decomposition to transform the coordinate system



• Given:
$$\mathbf{X} \in \mathbb{R}^{N \times d}$$
: $\mathbf{X} = \begin{bmatrix} x_{11} & \cdots & x_{1d} \\ \vdots & \ddots & \vdots \\ x_{N1} & \cdots & x_{Nd} \end{bmatrix}$

• Shift the points by their mean $\overline{x} \in \mathbb{R}^d$ (centralized data): $\widetilde{x}_i = x_i - \overline{x}$

Statistics:

Zero order statistic : number of points N

First order statistic: the mean of the N points, the vector $\overline{x} \in \mathbb{R}^d$:

$$\overline{\boldsymbol{x}} = \begin{bmatrix} \bar{x}_1 \\ \vdots \\ \bar{x}_d \end{bmatrix} = \frac{1}{N} \cdot \boldsymbol{X}^T \cdot \boldsymbol{1}_N$$

where $\mathbf{1}_N$ is an N-dimensional vector of ones

• Determine the variances $Var(\widetilde{X}_j)$ for each dimension $j \in \{1, ... d\}$

 \widetilde{X}

• Determine the covariance $Cov(\widetilde{X}_{j_1}, \widetilde{X}_{j_2})$ between dimensions j_1 and j_2 , $\forall j_1 \neq j_2 \in \{1, ... d\}$

 $\widetilde{X}^t \mid \int Cov$

ightharpoonup Leads to the covariance matrix $\mathbf{\Sigma}_{\widetilde{\mathbf{X}}} \in \mathbb{R}^{d imes d}$

Statistics:

Second order statistic: variance and covariance

The variance within the j-th dimension in \boldsymbol{X} is:

$$\operatorname{Var}(X_j) = \frac{1}{N} \sum_{i=1}^{N} (x_{ij} - \overline{x}_j)^2 = \frac{1}{N} \cdot X_j^T X_j - \overline{x}_j^2$$

The covariance between dimension j_1 and j_2 is:

$$Cov(X_{j_1}, X_{j_2}) = \frac{1}{N} \sum_{i=1}^{N} (x_{ij_1} - \overline{x}_{j_1}) \cdot (x_{ij_2} - \overline{x}_{j_2}) = \frac{1}{N} \cdot X_{j_1}^T X_{j_2} - \overline{x}_{j_1} \overline{x}_{j_2}$$

Statistics (continued):

For the set of points contained in **X** the corresponding covariance matrix is defined as:

$$\Sigma_{\boldsymbol{X}} = \begin{bmatrix} \operatorname{Var}(\boldsymbol{X}_{1}) & \operatorname{Cov}(\boldsymbol{X}_{1}, \boldsymbol{X}_{2}) & \dots & \operatorname{Cov}(\boldsymbol{X}_{1}, \boldsymbol{X}_{d}) \\ \operatorname{Cov}(\boldsymbol{X}_{2}, \boldsymbol{X}_{1}) & \operatorname{Var}(\boldsymbol{X}_{2}) & & & \\ \vdots & & \ddots & & \vdots \\ \operatorname{Cov}(\boldsymbol{X}_{d}, \boldsymbol{X}_{1}) & \dots & \operatorname{Var}(\boldsymbol{X}_{d}) \end{bmatrix} = \frac{1}{N} \boldsymbol{X}^{T} \boldsymbol{X} - \overline{\boldsymbol{x}} \, \overline{\boldsymbol{x}}^{T}$$

Remark: Covariance matrices are symmetric

$$Cov' = \begin{pmatrix} Var(1)' & 0 & \cdots & 0 \\ 0 & Var(2)' & \cdots & 0 \\ \vdots & \vdots & & \vdots \\ 0 & 0 & \cdots & Var(D)' \end{pmatrix}$$
Goal of PCA: Transformation of the coordinate system

- such that the covariances between the new axes are 0
- Approach:
 - Diagonalization by changing the basis (= adapt the coordinate system)
 - According to the spectral theorem, the eigenvectors of a symmetric matrix form an orthogonal basis
- Eigendecomposition of the covariance matrix: $\Sigma_{\widetilde{X}} = \Gamma \cdot \Lambda \cdot \Gamma^T$

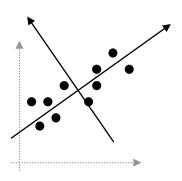
Eigendecomposition (spectral decomposition) is the factorization of $A \in \mathbb{R}^{d \times d}$:

$$A = \mathbf{\Gamma} \cdot \mathbf{\Lambda} \cdot \mathbf{\Gamma}^T$$

- \rightarrow matrices Γ , $\Lambda \in \mathbb{R}^{d \times d}$ with columns of Γ being the normalized eigenvectors γ_i
- $\rightarrow \Gamma$ is an orthonormal matrix: $\Gamma \cdot \Gamma^T = \Gamma^T \cdot \Gamma = \mathrm{Id} \ (\Gamma^T = \Gamma^{-1})$
- $\rightarrow \Lambda$ is a diagonal matrix with eigenvalues λ_i as the diagonal elements

ightharpoonup Eigendecomposition of the covariance matrix: $\Sigma_{\widetilde{X}} = \Gamma \cdot \Lambda \cdot \Gamma^T$

- The **new coordinate system** is defined by the eigenvectors γ_i :
 - Transformed data: $Y = \widetilde{X} \cdot \Gamma$
 - Λ is the covariance matrix in this new coordinate system
 - \triangleright New system has variance λ_i in dimension i
 - $\forall i_1 \neq i_2 : Cov(Y_{i_1}, Y_{i_2}) = 0$



Dimensionality Reduction with PCA

- Approach
 - The coordinates with low variance (hence low λ_i) can be ignored
 - − W.l.o.g. let us assume $\lambda_1 \ge \lambda_2 \ge \cdots \ge \lambda_d$
- \triangleright Truncation of Γ
 - Keep only columns (i.e. eigenvectors) of Γ corresponding to the largest k eigenvalues $\lambda_1,\ldots,\lambda_k$
 - $Y_{\text{reduced}} = \widetilde{X} \cdot \Gamma_{\text{truncated}}$
- How to pick k?
 - Frequently used: 90% rule; the k variances should explain 90% of the energy
 - k = smallest value ensuring $\sum_{i=1}^k \lambda_i \geq 0.9 \cdot \sum_{i=1}^d \lambda_i$
- The modified points (transformed and truncated) contain most of the information of the original points and are low dimensional

Complexity

Complexity of PCA:

$$O(N \cdot d^2) + O(d^3) + O(N \cdot d \cdot k) = O(N \cdot d^2 + d^3)$$
Compute Eigenvalue Project data onto decomposition the k-dimensional space

- Remarks on eigenvalue decomposition:
 - Usually we are interested in the reduced data only
 - > Only the k largest eigenvectors required (i.e. not all of them)
 - Use iterative approaches (next slide) for finding eigenvectors

How to Compute Eigenvectors?

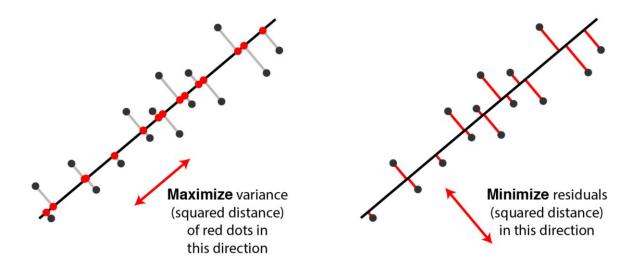
- Eigenvalues are important for many machine learning/data mining tasks
 - PCA, Ranking of Websites, Community Detection, ...
 // see our other lecture!
 - How to compute them efficiently?
- Power iteration (a.k.a. Von Mises iteration)
 - Iterative approach to compute a single eigenvector
- ullet Let $oldsymbol{A}$ be a matrix and $oldsymbol{v}$ be an arbitrary (normalized) vector
 - Iteratively compute $v \leftarrow \frac{A \cdot v}{\|A \cdot v\|}$ until convergence
 - $-\,$ in each step, $oldsymbol{v}$ is simply multiplied with $oldsymbol{A}$ and normalized
 - $-\ v$ converges to the eigenvector of A with largest absolute value
 - Highly efficient for sparse data

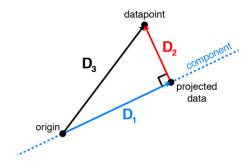
How to Compute Eigenvectors?

Convergence:

- Linear convergence with rate $|\lambda_2/\lambda_1|$
- Fast convergence if first and second eigenvalue are dissimilar
- How to find multiple (the k largest) eigenvectors?
 - Let us focus on symmetric matrices A
 - Eigenvalue decomposition leads to: $\mathbf{A} = \mathbf{\Gamma} \cdot \mathbf{\Lambda} \cdot \mathbf{\Gamma}^T = \sum_{i=1}^d \lambda_i \cdot \mathbf{\gamma}_i \cdot \mathbf{\gamma}_i^T$
 - Define deflated matrix: $\widehat{A} = A \lambda_1 \cdot \gamma_1 \cdot \gamma_1^T$
 - $-\widehat{A}$ has the same eigenvectors as A except the first one has become zero
 - ightharpoonup Apply power iteration on \widehat{A} to find the second largest eigenvector of A

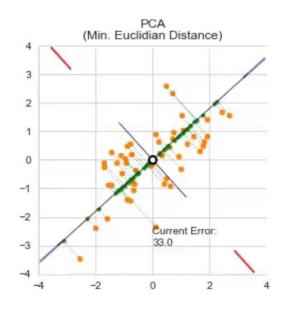
Alternative views of PCA

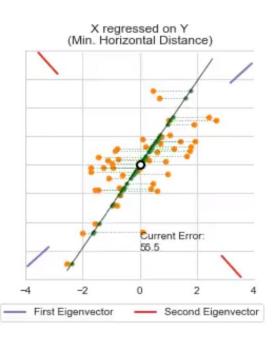




Images adapted from Alexh Williams

PCA vs. Regression





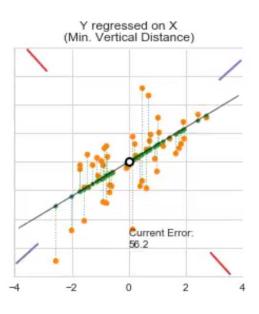


Image adapted from Quentin André

PCA: Summary

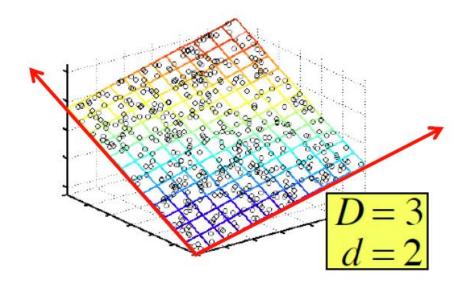
- PCA finds the optimal transformation by deriving uncorrelated dimensions
 - Exploits eigendecomposition
- Dimensionality reduction
 - After transformation simply remove dimensions with lowest variance (or use only the k largest eigenvectors for transformation)
- Limitations
 - Only captures linear relationships (one solution: Kernel PCA)

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 - Idea: Low Rank Approximation
 - SVD & Latent Factors
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Low-Dimensional Manifold

Data often lies on a low-dimensional manifold embedded in higher dimensional space



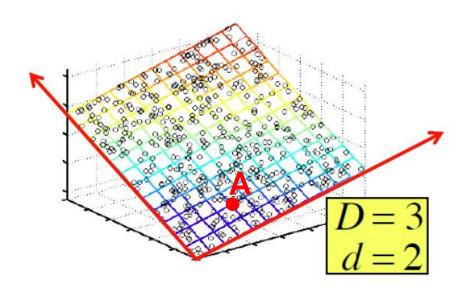
- How can me measure the dimensionality of this manifold?
 - put differently how to measure the intrinsic dimensionality of the data
- How can we find this manifold?

Rank of a Matrix

- Q: What is the rank of a matrix A?
- A: Number of linearly independent columns/rows of A
- Example: $\text{ Matrix A} = \begin{bmatrix} 1 & 2 & 1 \\ -2 & -3 & 1 \\ 3 & 5 & 0 \end{bmatrix}$ has rank r=2
 - Why? The first two rows are linearly independent, so the rank is at least 2, but all three rows are linearly dependent (the first is equal to the sum of the second and third) so the rank must be less than 3.
- Why do we care about low rank?
 - We can write A as two "basis" vectors: [1 2 1] [-2 -3 1]
 - And new coordinates of: [1 0] [0 1] [1 -1]

Rank is "Dimensionality"

- Cloud of points in 3D space:
 - Think of point positions as a matrix:



- We can rewrite coordinates more efficiently!
 - Old basis vectors: [1 0 0] [0 1 0] [0 0 1]
 - New basis vectors: [1 2 1] [-2 -3 1]
 - Then A has new coordinates: [1 0], B: [0 1], C: [1 -1]
 - Notice: We reduced the number of coordinates!

Low Rank Approximation

Idea: approximate original data A by a low rank matrix B

$$\mathbf{A} = \begin{bmatrix} 1.01 & 2.05 & 0.9 \\ -2.1 & -3.05 & 1.1 \\ 2.99 & 5.01 & 0.3 \end{bmatrix} \approx \begin{bmatrix} 1 & 2 & 1 \\ -2 & -3 & 1 \\ 3 & 5 & 0 \end{bmatrix} = \mathbf{B}$$

$$rank(A) = 3$$

we need 3 coordinates
to describe each point

$$rank(\mathbf{B}) = 2$$

we need only 2 coordinates
per point

Low Rank Approximation

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- Important: Even though both A and B are $\in R^{n \times d}$ we need only two coordinates per point to describe B
 - rank(A)=3 vs. rank(B)=2 (3 vs. 2 coordinates per point)
- Goal: Find the best low rank approximation
 - best = minimize the sum of reconstruction error
 - Given matrix $A \in \mathbb{R}^{n \times d}$, find $B \in \mathbb{R}^{n \times d}$ with rank(B) = k that minimizes

$$\|\mathbf{A} - \mathbf{B}\|_F^2 = \sum_{i=1}^N \sum_{j=1}^D (a_{ij} - b_{ij})^2$$

Roadmap

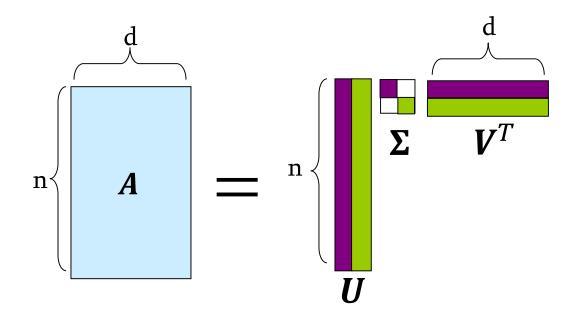
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Singular Value Decomposition (SVD): Definition

- Each real matrix $A \in \mathbb{R}^{n \times d}$ can be decomposed into $A = U \cdot \Sigma \cdot V^{T}$ (note: **exact representation**, no approximation), where
 - $\mathbf{U} \in \mathbb{R}^{n \times r}$, $\mathbf{\Sigma} \in \mathbb{R}^{r \times r}$, $\mathbf{V} \in \mathbb{R}^{d \times r}$
 - *U*, *V*: column orthonormal
 - i.e. $U^T U = I$; $V^T V = I$ (I: identity matrix)
 - \boldsymbol{U} are called the left singular vectors , \boldsymbol{V} the right singular vectors
 - Σ: diagonal
 - $-r \times r$ diagonal matrix (r: rank of matrix A)
 - entries (called singular values) are positive, and sorted in decreasing order ($\sigma_1 \ge \sigma_2 \ge ... \ge \sigma_r > 0$)
- Remark: The decomposition is (almost) unique
 - see e.g. multiplication by -1

Singular Value Decomposition

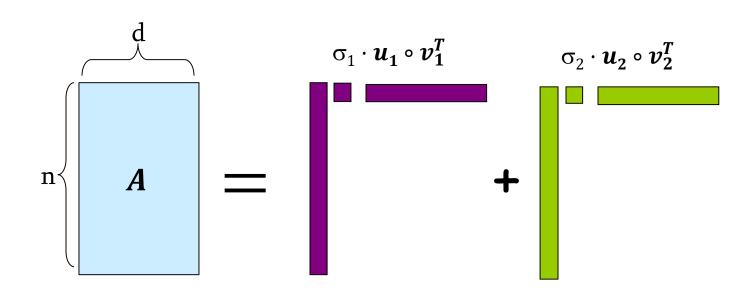
$$A = U\Sigma V^T$$



here: r = 2

Singular Value Decomposition

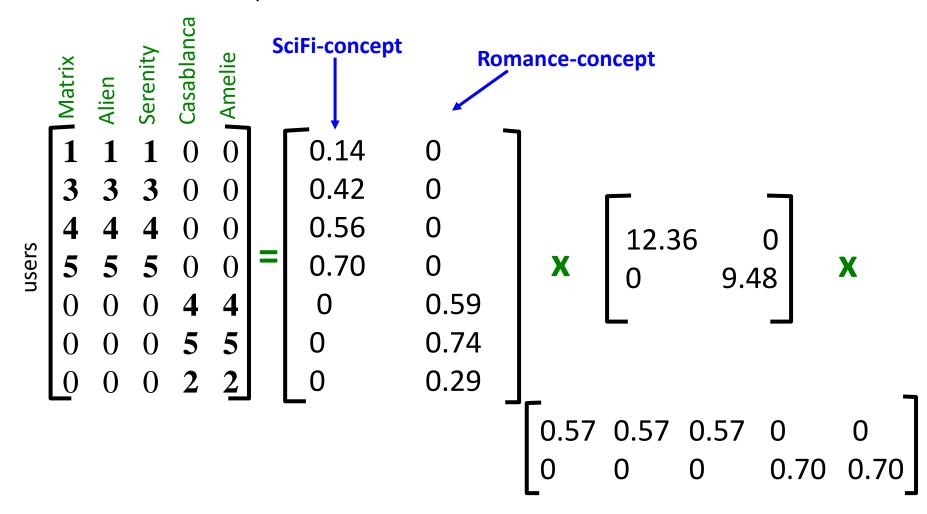
$$m{A} = m{U} m{\Sigma} m{V}^T = \sum_{i=1}^r \sigma_i \cdot m{u}_i \circ m{v}_i^T$$



here: r = 2

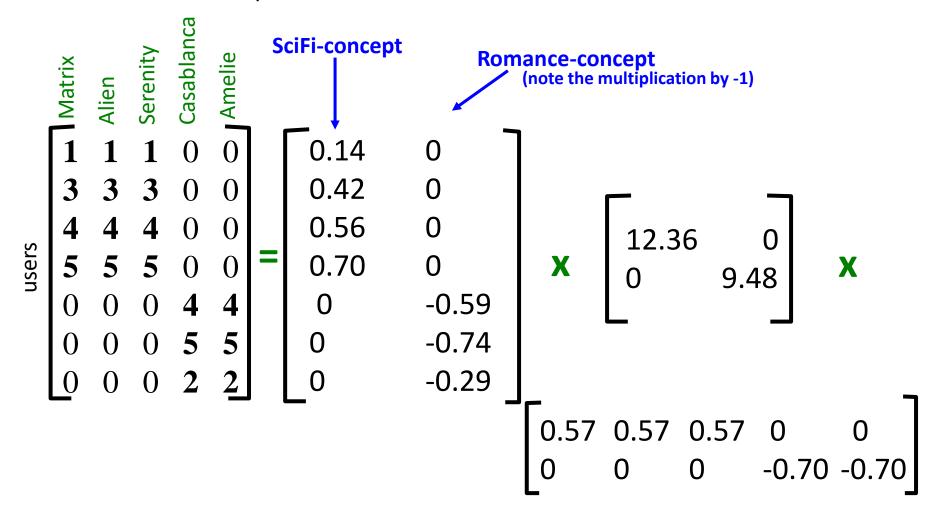
SVD Example: Users-to-Movies

• $A = U\Sigma V^T$ - example: Users to Movies



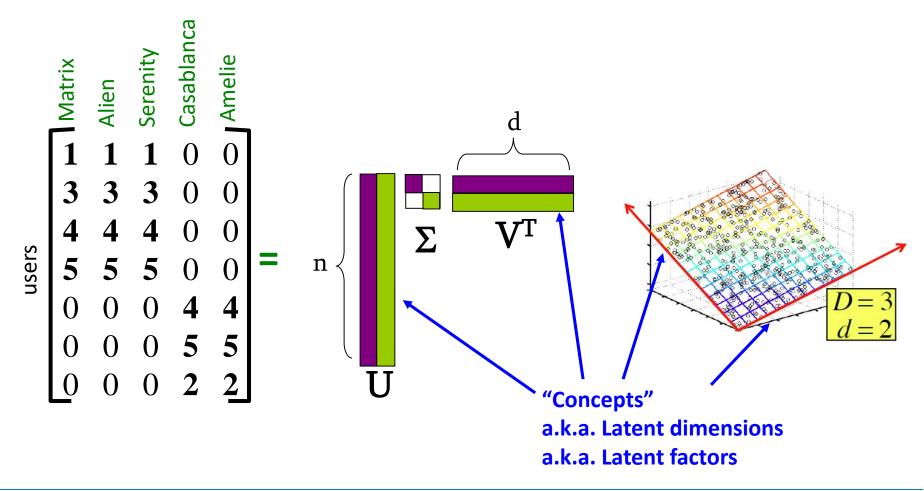
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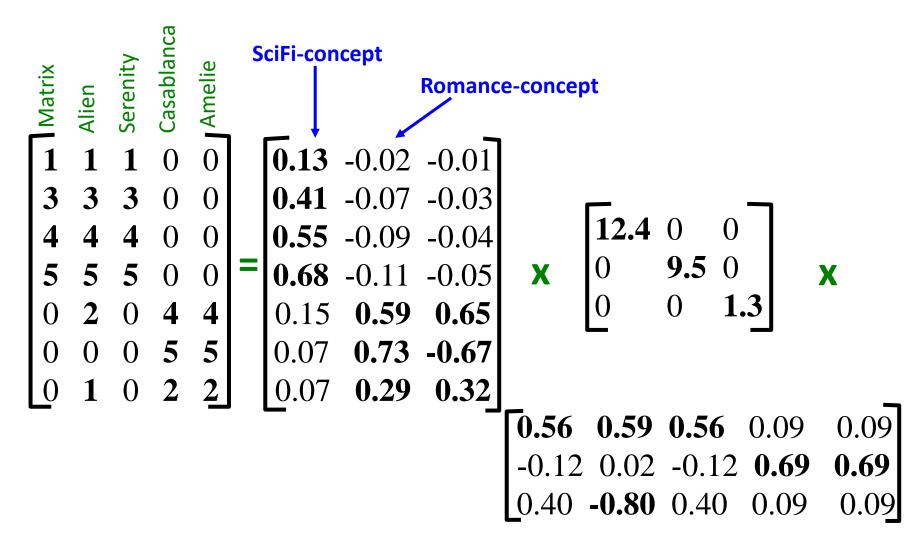


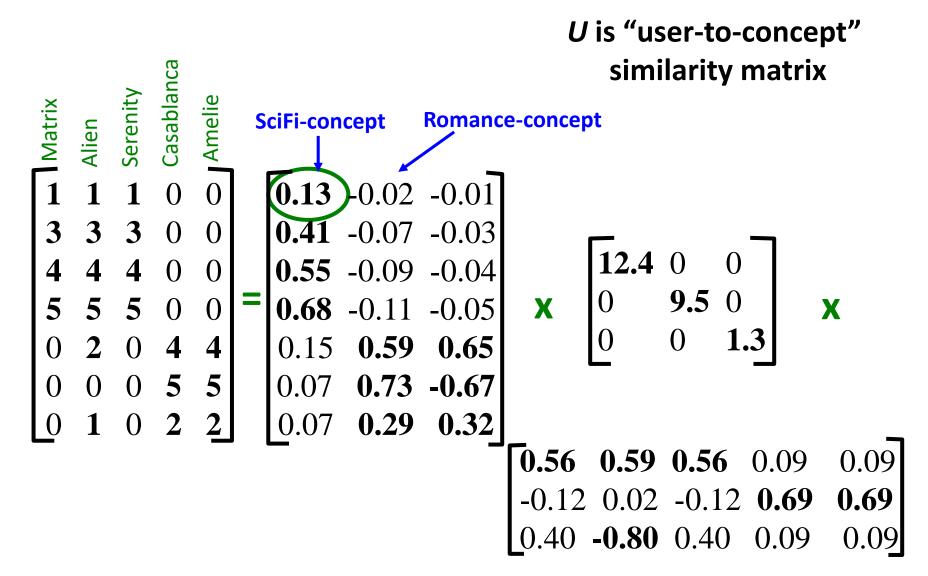
SVD Example: Latent Factors

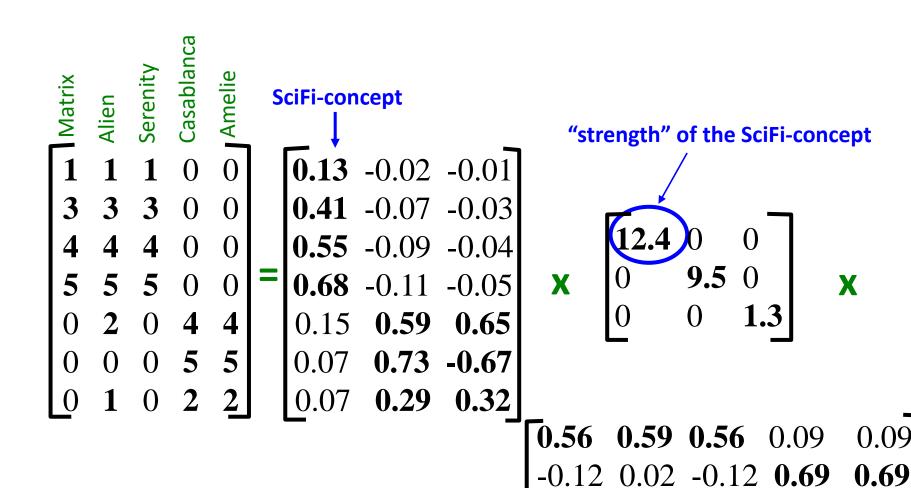
• $A = U\Sigma V^T$ - example: Users to Movies

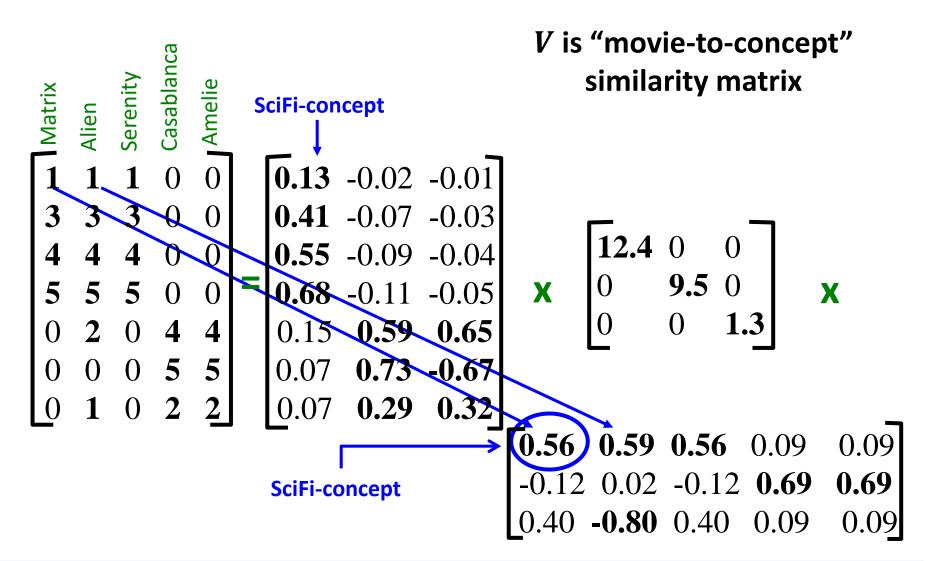


0.69









SVD: Interpretation

- $A = U\Sigma V^T$
- 'movies', 'users' and 'concepts':
 - A original data: movies-to-users
 - U: user-to-concept similarity matrix
 - V: movie-to-concept similarity matrix
 - Σ: its diagonal elements: 'strength' of each concept

Benefits of SVD (or in general matrix decomposition):

- Discover hidden correlations/topics
 - Words that occur commonly together; movies of the same genre; ...
- Interpretation and visualization

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Recap: Dim. Reduction by Low Rank Approx.

• Idea: approximate original data A by a low rank matrix B

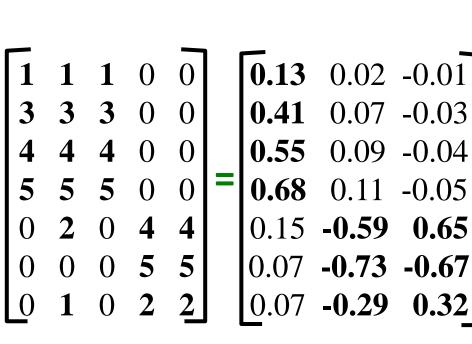
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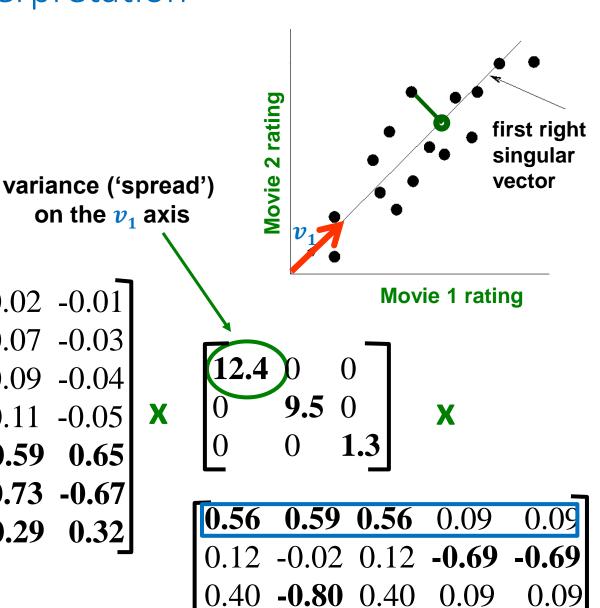
- Important: Even though both A and B are $\in \mathbb{R}^{n \times d}$ we need only two coordinates per point to describe B
 - $-\operatorname{rank}(\mathbf{A}) = 3 \text{ vs. } \operatorname{rank}(\mathbf{B}) = 2$ (3 vs. 2 coordinates per point)
- Goal: Find the best low rank approximation
 - best = minimize the sum of reconstruction error
 - Given matrix $\mathbf{A} \in \mathbb{R}^{n \times d}$, find $\mathbf{B} \in \mathbb{R}^{n \times d}$ with rank $(\mathbf{B}) = k$ that minimizes

$$\|\mathbf{A} - \mathbf{B}\|_F^2 = \sum_{i=1}^N \sum_{j=1}^D (a_{ij} - b_{ij})^2$$

SVD: Alternative Interpretation

• A = $U\Sigma V^T$ example:





on the v_1 axis

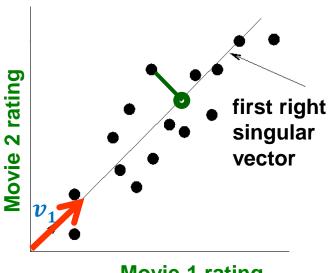
SVD: Alternative Interpretation

- A = $U\Sigma V^T$ example:
- U Σ: Gives the coordinates of the points in the projection axis

Projection of users on the "Sci-Fi" axis U Σ :

1	1	1	0	0
3	3	3	0	0
4	4	4	0	0
5	5	5	0	0
0	2	0	4	4
0	0	0	5	5
0	1	0	2	2

1.61	0.19	-0.01
5.08	0.66	-0.03
6.82	0.85	-0.05
8.43	1.04	-0.06
1.86	-5.60	0.84
0.86	-6.93	-0.87
0.86	-2.75	0.41 _



Movie 1 rating

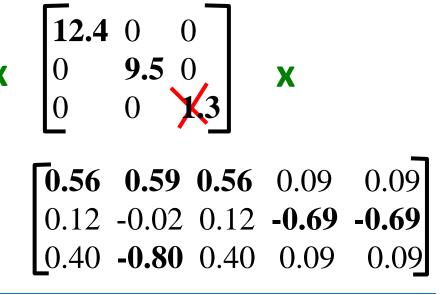
 0.56
 0.59
 0.56
 0.09
 0.09

 0.12
 -0.02
 0.12
 -0.69
 -0.69

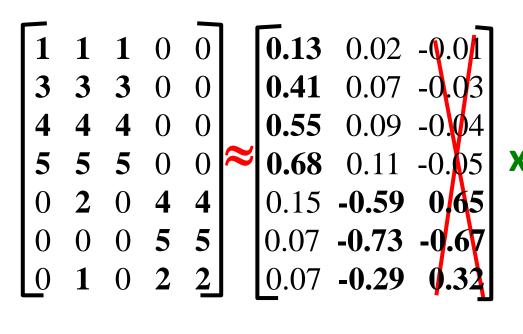
 0.40
 -0.80
 0.40
 0.09
 0.09

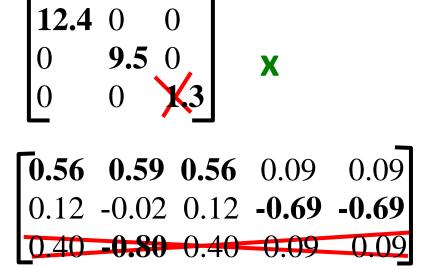
• How to find the best approximation?

- How to find the best approximation?
- Set smallest singular values to zero!

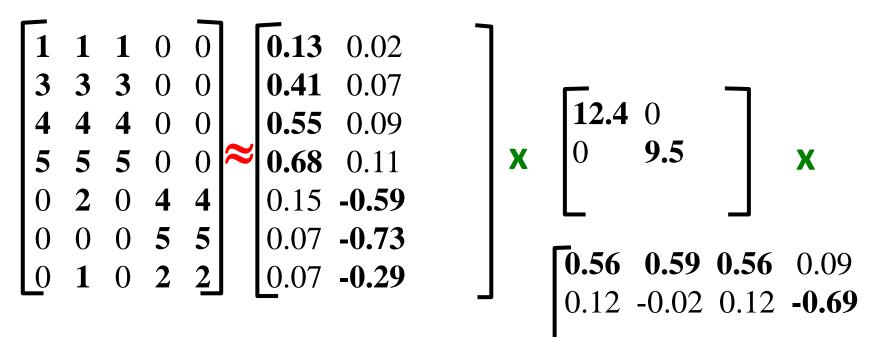


- How to find the best approximation?
- Set smallest singular values to zero!





- How to find the best approximation?
- Set smallest singular values to zero!



- How to find the best approximation?
- Set smallest singular values to zero!

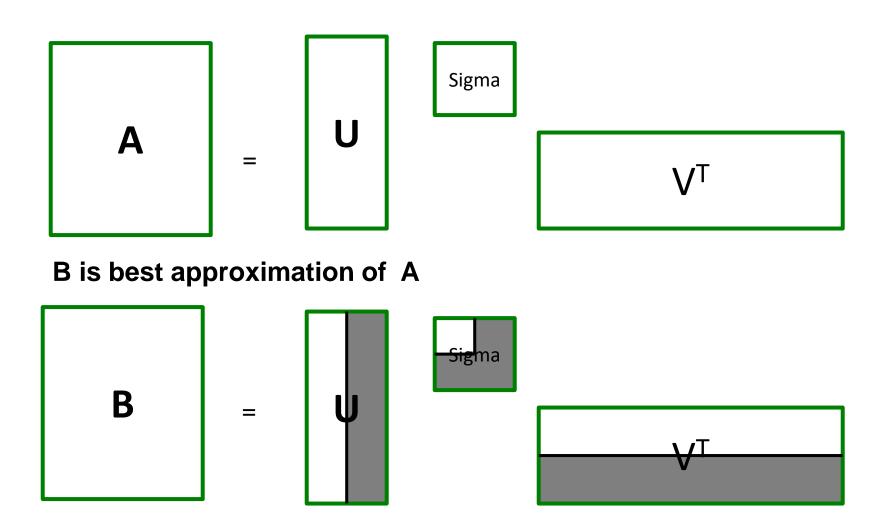
by construction, the rank of the new matrix is 2

1	1	1	0	0
3	3	3	0	0
4	4	4	0	0
5	5	5	0	0
0	2	0	4	4
0	0	0	5	5
0	1	0	2	2



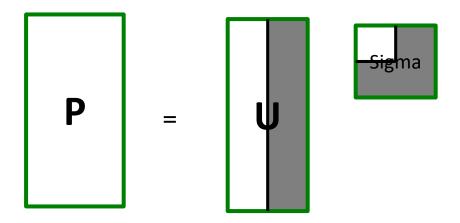
0.92	0.95	0.92	0.01	0.01
2.91	3.01	2.91	-0.01	-0.01
3.90	4.04	3.90	0.01	0.01
4.82	5.00	4.82	0.03	0.03
0.70	0.53	0.70	4.11	4.11
-0.69				
0.32	0.23	0.32	2.01	2.01

SVD: Best Low Rank Approximation



SVD: Projection

Note: The actual projected/reduced data can be obtained by computing



• Or equivalently: $P = A \cdot V$

(since V is orthonormal)

Best Approximation – Intuitive Explanation

- Recap: Vectors $oldsymbol{u}_i$ and $oldsymbol{v}_i$ are of unit length
- W.l.o.g.: $\sigma_1 \ge \sigma_2 \ge \sigma_3 \ge ... \ge 0$

$$\begin{bmatrix} \mathbf{1} & \mathbf{1} & \mathbf{1} & 0 & 0 \\ \mathbf{3} & \mathbf{3} & \mathbf{3} & 0 & 0 \\ \mathbf{4} & \mathbf{4} & \mathbf{4} & 0 & 0 \\ \mathbf{5} & \mathbf{5} & \mathbf{5} & 0 & 0 \\ 0 & \mathbf{2} & 0 & \mathbf{4} & \mathbf{4} \\ 0 & 0 & 0 & \mathbf{5} & \mathbf{5} \\ 0 & \mathbf{1} & 0 & \mathbf{2} & \mathbf{2} \end{bmatrix} = \begin{bmatrix} \begin{vmatrix} & & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ \end{bmatrix} \times \begin{bmatrix} \sigma_1 & \swarrow \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ \end{bmatrix} \times \begin{bmatrix} \sigma_1 & \swarrow \\ & & \sigma_2 \end{bmatrix} \times \begin{bmatrix} \sigma_1 & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ \end{bmatrix}$$

Best Approximation – Intuitive Explanation

- Recap: Vectors $oldsymbol{u}_i$ and $oldsymbol{v}_i$ are of unit length
- W.l.o.g.: $\sigma_1 \ge \sigma_2 \ge \sigma_3 \ge ... \ge 0$

$$\begin{bmatrix} 1 & 1 & 1 & 0 & 0 \\ 3 & 3 & 3 & 0 & 0 \\ 4 & 4 & 4 & 0 & 0 \\ 5 & 5 & 5 & 0 & 0 \\ 0 & 2 & 0 & 4 & 4 \\ 0 & 0 & 0 & 5 & 5 \\ 0 & 1 & 0 & 2 & 2 \end{bmatrix} = \begin{matrix} & & & & & & & & & \\ & & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & \\ & & & \\ & & \\ & & & \\ & &$$

- σ_i scales the terms $\boldsymbol{u}_i \cdot \boldsymbol{v}_i^T$
- Zeroing small σ_i introduces less error

 $\sum_{i=1}^{k} \sigma_i^2 \ge 0.9 \sum_{i=1}^{r} \sigma_i^2$

SVD: Best Low Rank Approximation - Proof

- Theorem: Let A = $U\Sigma V^T$ ($\sigma_1 \geq \sigma_2 \geq \cdots$, rank(A)=r) and B = USV^T with S being a diagonal $r \times r$ matrix where
 - $s_i = \sigma_i$ for i=1...k and $s_i = 0$ else

Then B is a best rank-k approximation to A regarding Frobenius norm, i.e. B is a solution to $\min_{B} ||A - B||_{F}$ where rank(B)=k

- We have uploaded a detailed proof to the web
 - Note: Many proofs on the web and on other lecture slides are incorrect!
- Remark: B is also an optimal low-rank approximation regarding the spectral norm (operator 2-norm): $\min_{B} ||A B||_2$
 - $\|X\|_2$ = largest singular value of X

SVD: Best Low Rank Approximation - Proof

Some facts:

$$- \|X\|_F = \|X^T\|_F$$

obvious from the definition

$$- \|X\|_F^2 = \operatorname{trace}(X^T X)$$

// trace = sum of diagonal entries

- easy homework
- Frobenius norm is invariant to orthonormal transformations U
 - Note: If $U^TU = I$ then also $UU^T = I$
 - $\|UX\|_F^2 = \operatorname{trace}((UX)^T(UX)) = \operatorname{trace}(X^TU^TUX) = \operatorname{trace}(X^TX) = \|X\|_F^2$

$$- \|XU\|_F^2 = \|(XU)^T\|_F^2 = \operatorname{trace}((XU)(XU)^T)$$
$$= \operatorname{trace}(XUU^TX^T) = \operatorname{trace}(XX^T) = \|X\|_F^2$$

- Let $A = U\Sigma V^T$ then $||A||_F^2 = ||\Sigma||_F^2 = \sum_i^r \sigma_i^2$
 - follows from above results

SVD: Complexity

- To compute SVD:
 - $O(n \cdot d^2)$ or $O(n^2 \cdot d)$ (whichever is less)
- But:
 - Less work, if we just want singular values
 - or if we want first k singular vectors
 - or if the matrix is sparse
- Implemented in linear algebra packages like
 - LINPACK, Matlab, SPlus, Mathematica ...

SVD & PCA: Comparison

- Given data X (let's assume it is already centered)
- SVD gives us:
 - $X = U\Sigma V^T$
 - Projected data obtained by $X \cdot V$

(or truncated V)

- PCA computes the eigendecomposition of the covariance matrix
 - Covariance matrix: X^TX
 - Eigendecomposition leads to $X^TX = \Gamma \cdot \Lambda \cdot \Gamma^T$
 - Projected data obtained by X · Γ

(or truncated Γ)

Let us calculate:

$$- X^T X = (U \Sigma V^T)^T U \Sigma V^T = V \Sigma^T U^T (U \Sigma V^T) = V \Sigma \Sigma^T V^T = V \Sigma^2 V^T$$

- Γ=V

PCA and SVD are equivalent!

 $-\Sigma^2 = \Lambda$

squared singular values are variances in new space!

SVD & PCA: Comparison

transform the data such that dimensions of new space are uncorrelated + discard (new) dimensions with smallest variance

=

find optimal low-rank approximation (regarding Frobenius norm)

Remark: Computation of SVD

We can use the eigendecomposition to calculate the singular value decomposition

- $X^TX = (U\Sigma V^T)^TU\Sigma V^T = V\Sigma^TU^T(U\Sigma V^T) = V\Sigma\Sigma^TV^T = V\Sigma^2V^T$ - $V = \text{eigenvectors of } X^TX$
- $XX^T = U\Sigma V^T (V\Sigma^T U^T) = U\Sigma \Sigma^T U^T$
 - U = eigenvectors of XX^T
- Drawback: Numerically instable
 - better to use specialized algorithms