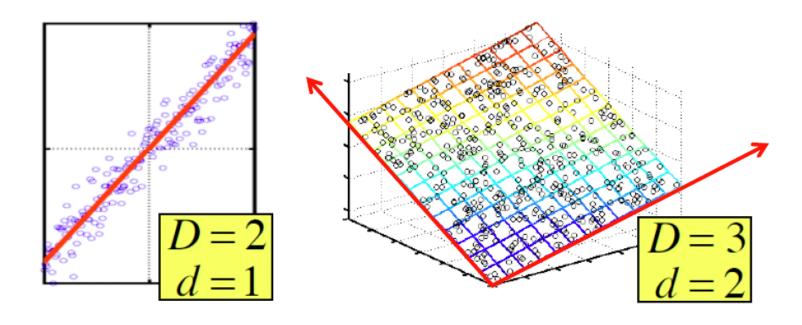
# Dimensionality Reduction: SVD & CUR

CS246: Mining Massive Datasets
Jure Leskovec, Stanford University
http://cs246.stanford.edu



# **Dimensionality Reduction**



- Assumption: Data lies on or near a low d-dimensional subspace
- Axes of this subspace are effective representation of the data

# **Dimensionality Reduction**

- Compress / reduce dimensionality:
  - 10<sup>6</sup> rows; 10<sup>3</sup> columns; no updates
  - Random access to any cell(s); small error: OK

$\mathbf{day}$	We	${ m Th}$	$\mathbf{F}$ r	$\mathbf{S}\mathbf{a}$	$\mathbf{S}\mathbf{u}$
customer	7/10/96	7/11/96	7/12/96	7/13/96	7/14/96
ABC Inc.	1	1	1	0	0
DEF Ltd.	2	2	2	0	0
GHI Inc.	1	1	1	0	0
KLM Co.	5	5	5	0	0
$\mathbf{Smith}$	0	0	0	2	2
$_{ m Johnson}$	0	0	0	3	3
Thompson	0	0	0	1	1

The above matrix is really "2-dimensional." All rows can be reconstructed by scaling [1 1 1 0 0] or [0 0 0 1 1]

## Rank of a Matrix

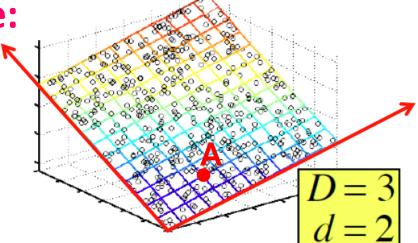
- Q: What is rank of a matrix A?
- A: Number of linearly independent columns of A
- For example:
  - Matrix  $\mathbf{A} = \begin{bmatrix} 1 & 2 & 1 \\ -2 & -3 & 1 \\ 3 & 5 & 0 \end{bmatrix}$  has rank  $\mathbf{r} = \mathbf{2}$ 
    - Why? The first two rows are linearly independent, so the rank is 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 3D space:

■ Think of point positions

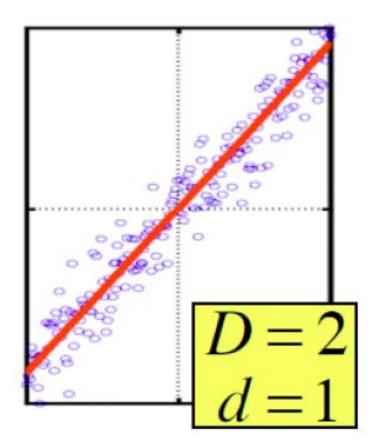
as a matrix:  $\begin{bmatrix} 1 & 2 & 1 \\ -2 & -3 & 1 \\ 3 & 5 & 0 \end{bmatrix}$  A B C



- We can rewrite coordinates more efficiently!
  - Old coordinate system: [1 0 0] [0 1 0] [0 0 1]
  - New coordinate system: [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!

# **Dimensionality Reduction**

Goal of dimensionality reduction is to discover the axis of data!



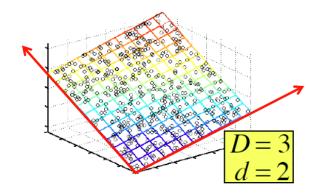
Rather than representing every point with 2 coordinates we represent each point with 1 coordinate (corresponding to the position of the point on the red line).

By doing this we incur a bit of **error** as the points do not exactly lie on the line

# Why Reduce Dimensions?

#### Why reduce dimensions?

- Discover hidden correlations/topics
  - Words that occur commonly together
- Remove redundant and noisy features
  - Not all words are useful
- Interpretation and visualization
- Easier storage and processing of the data



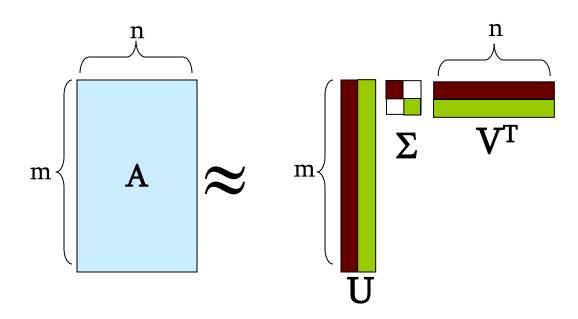
## **SVD** - Definition

$$\mathbf{A}_{[m \times n]} = \mathbf{U}_{[m \times r]} \boldsymbol{\Sigma}_{[r \times r]} (\mathbf{V}_{[n \times r]})^{\mathsf{T}}$$

- A: Input data matrix
  - m x n matrix (e.g., m documents, n terms)
- U: Left singular vectors
  - m x r matrix (m documents, r concepts)
- Σ: Singular values
  - r x r diagonal matrix (strength of each 'concept')
     (r: rank of the matrix A)
- V: Right singular vectors
  - n x r matrix (n terms, r concepts)

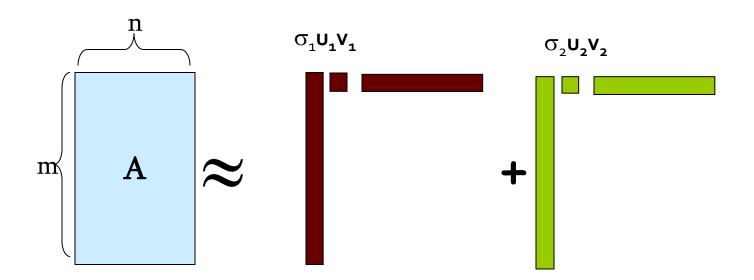
### SVD

$$\mathbf{A} pprox \mathbf{U} \mathbf{\Sigma} \mathbf{V}^T = \sum_i \sigma_i \mathbf{u}_i \circ \mathbf{v}_i^{\mathsf{T}}$$



### SVD

$$\mathbf{A} pprox \mathbf{U} \mathbf{\Sigma} \mathbf{V}^T = \sum_i \sigma_i \mathbf{u}_i \circ \mathbf{v}_i^{\mathsf{T}}$$



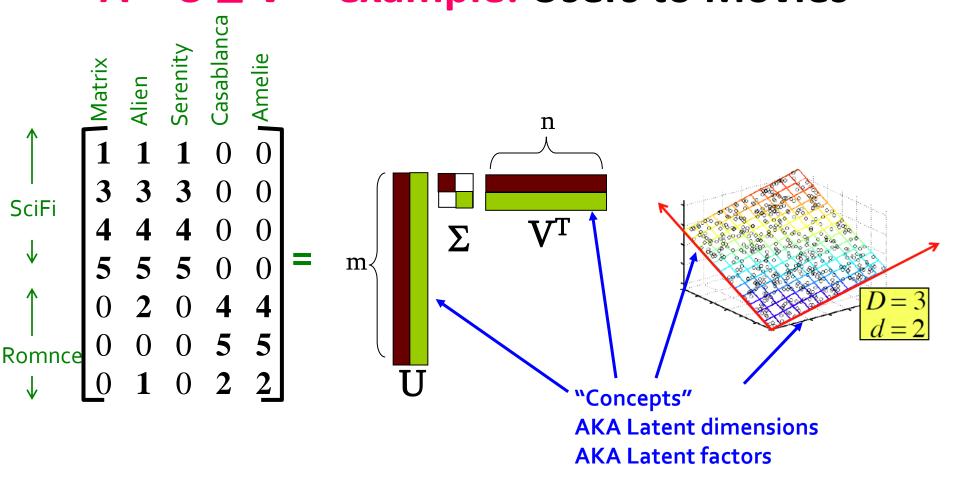
 $\sigma_i$  ... scalar  $u_i$  ... vector  $v_i$  ... vector

## **SVD** - Properties

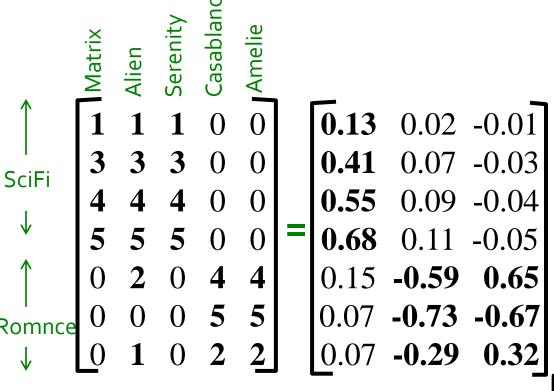
It is **always** possible to decompose a real matrix  $\boldsymbol{A}$  into  $\boldsymbol{A} = \boldsymbol{U} \boldsymbol{\Sigma} \boldsymbol{V}^{\mathsf{T}}$ , where

- **U**, Σ, *V*: unique
- U, V: column orthonormal
  - $U^T U = I$ ;  $V^T V = I$  (I: identity matrix)
  - (Columns are orthogonal unit vectors)
- Σ: diagonal
  - Entries (singular values) are positive, and sorted in decreasing order  $(\sigma_1 \ge \sigma_2 \ge ... \ge 0)$

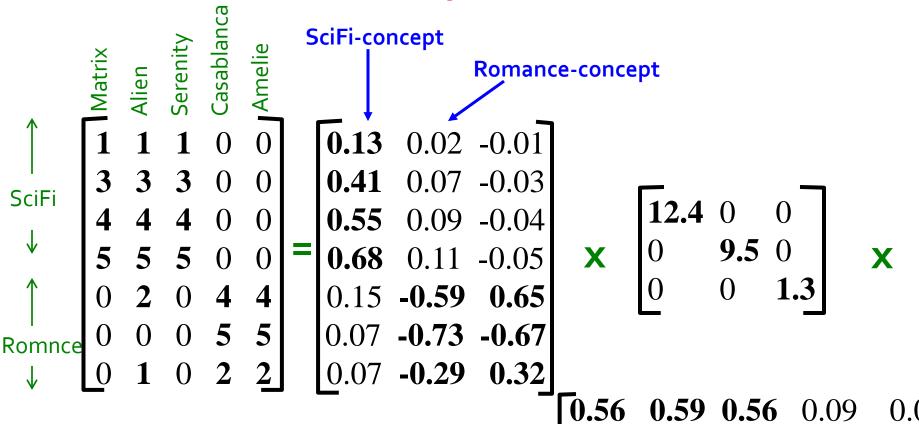
## ■ A = U $\Sigma$ V<sup>T</sup> - example: Users to Movies



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



## ■ A = U $\Sigma$ V<sup>T</sup> - example: Users to Movies



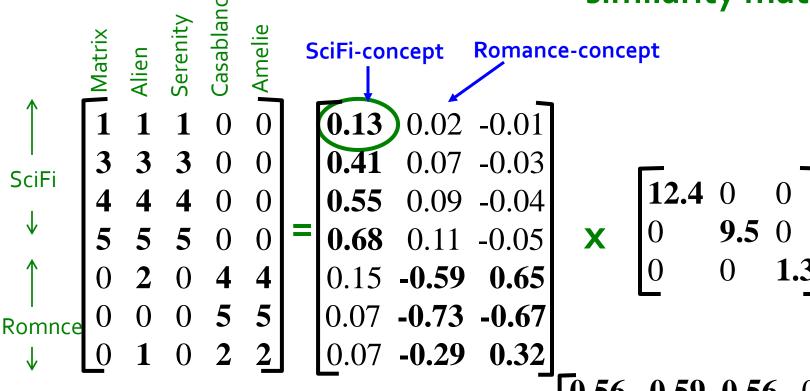
**-0.02 0.12 -0.69 -0.69** 

0.09

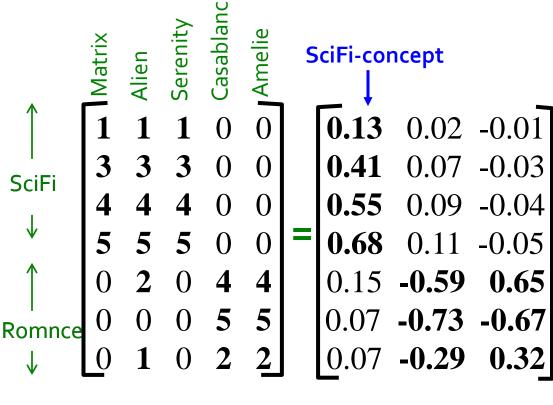
**-0.80** 0.40

## ■ $A = U \Sigma V^T$ - example:

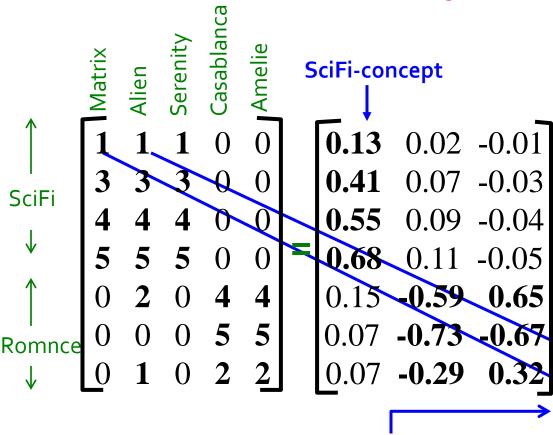
*U* is "user-to-concept" similarity matrix



## • $A = U \Sigma V^T$ - example:



## • $A = U \Sigma V^T$ - example:



V is "movie-to-concept" similarity matrix

$$\begin{array}{c|cccc}
\mathbf{X} & \begin{bmatrix}
\mathbf{12.4} & 0 & 0 \\
0 & \mathbf{9.5} & 0 \\
0 & 0 & \mathbf{1.3}
\end{bmatrix} \quad \mathbf{X}$$

SciFi-concept

**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

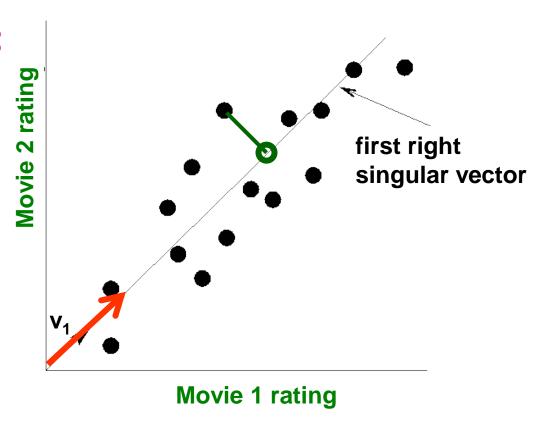
## 'movies', 'users' and 'concepts':

- U: user-to-concept similarity matrix
- V: movie-to-concept similarity matrix
- Σ: its diagonal elements: 'strength' of each concept

# Dimensionality Reduction with SVD

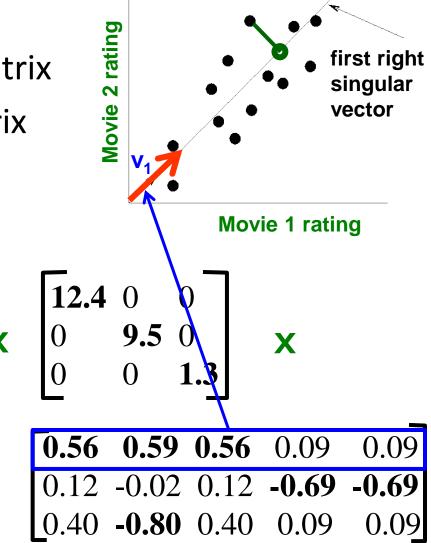
# SVD – Dimensionality Reduction

- SVD gives 'best' axis to project on:
  - 'best' = min sum of squares of projection errors
- In other words,
   minimum
   reconstruction
   error



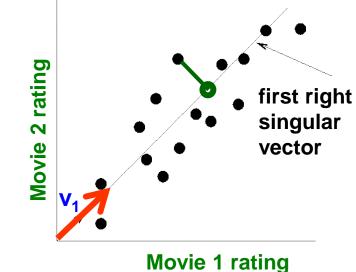
## • $A = U \Sigma V^T$ - example:

- V: "movie-to-concept" matrix
- U: "user-to-concept" matrix

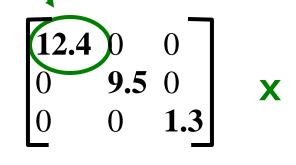




variance ('spread') on the v<sub>1</sub> axis

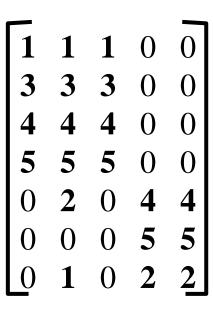


1	1	1	0	0	
3	3	3	0	0	
4	4	4	0	0	
5	5	5	0	0	:
0	2	0	4	4	
0	0	0	5	5	
<b>0</b>	1	<b>0</b>	2	2	

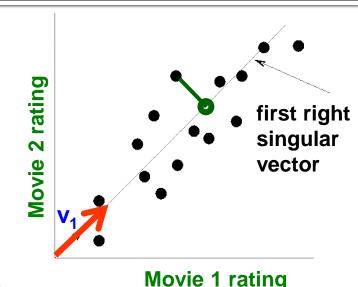


## $A = U \Sigma V^{T}$ - example:

 U Σ: Gives the coordinates of the points in the projection axis



Projection of users on the "Sci-Fi" axis  $((U \Sigma)^T)$ :



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

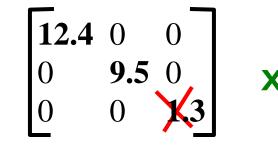
#### **More details**

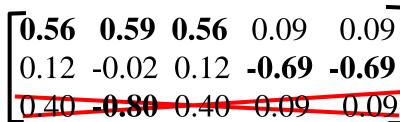
Q: How exactly is dim. reduction done?

- Q: How exactly is dim. reduction done?
- A: Set smallest singular values to zero

- Q: How exactly is dim. reduction done?
- A: Set smallest singular values to zero

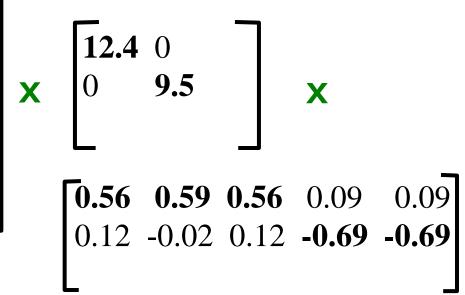
- Q: How exactly is dim. reduction done?
- A: Set smallest singular values to zero





- Q: How exactly is dim. reduction done?
- A: Set smallest singular values to zero

$$\begin{bmatrix} \mathbf{1} & \mathbf{1} & \mathbf{1} & \mathbf{0} & \mathbf{0} \\ \mathbf{3} & \mathbf{3} & \mathbf{3} & \mathbf{0} & \mathbf{0} \\ \mathbf{4} & \mathbf{4} & \mathbf{4} & \mathbf{0} & \mathbf{0} \\ \mathbf{5} & \mathbf{5} & \mathbf{5} & \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \mathbf{2} & \mathbf{0} & \mathbf{4} & \mathbf{4} \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{5} & \mathbf{5} \\ \mathbf{0} & \mathbf{1} & \mathbf{0} & \mathbf{2} & \mathbf{2} \end{bmatrix}$$
 
$$\begin{bmatrix} \mathbf{0.13} & 0.02 \\ \mathbf{0.41} & 0.07 \\ \mathbf{0.55} & 0.09 \\ \mathbf{0.68} & 0.11 \\ \mathbf{0.15} & \mathbf{-0.59} \\ \mathbf{0.07} & \mathbf{-0.73} \\ \mathbf{0.07} & \mathbf{-0.73} \\ \mathbf{0.07} & \mathbf{-0.29}$$



#### **More details**

- Q: How exactly is dim. reduction done?
- A: Set smallest singular values to zero

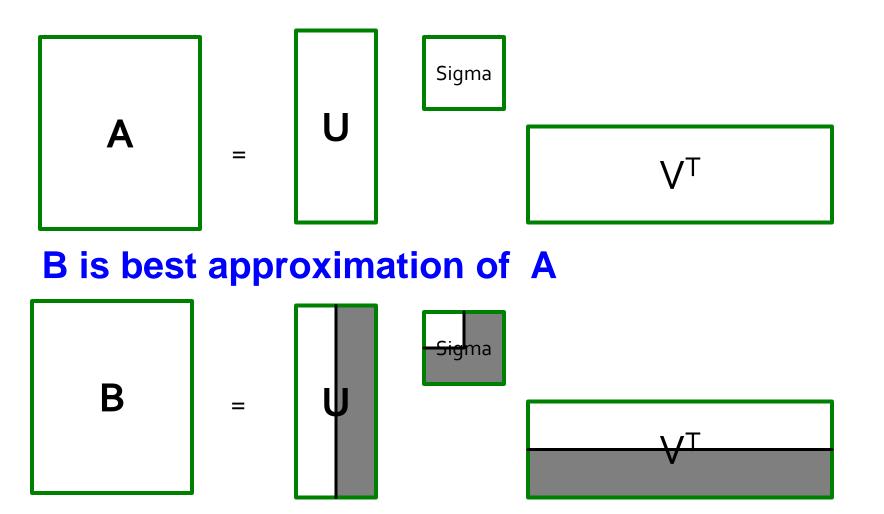


#### Frobenius norm:

$$\|\mathbf{M}\|_{\mathrm{F}} = \sqrt{\sum_{ij} \mathbf{M}_{ij}}^2$$

 $\|\mathbf{A} - \mathbf{B}\|_{\mathbf{F}} = \sqrt{\Sigma_{ii} (\mathbf{A}_{ii} - \mathbf{B}_{ii})^2}$ 

## SVD – Best Low Rank Approx.



## SVD – Best Low Rank Approx.

#### Theorem:

Let  $A = U \sum V^T$  where  $\sum : \sigma_1 \ge \sigma_2 \ge ...$ , and rank(A) = rthen  $B = U \sum V^T$  is a **best** rank(B) = k approx. to A

• S = diagonal  $n \times n$  matrix where  $s_i = \sigma_i$  (i = 1...k) else  $s_i = 0$ 

#### What do we mean by "best":

■ B is a solution to  $\min_{B} ||A-B||_{F}$  where  $\operatorname{rank}(B)=k$ 

$$\begin{pmatrix} x_{11} & x_{12} & \dots & x_{1n} \\ x_{21} & x_{22} & \dots & \\ \vdots & \vdots & \ddots & \\ x_{m1} & & & x_{mn} \end{pmatrix} = \begin{pmatrix} u_{11} & \dots & \\ \vdots & \ddots & \\ u_{m1} & & & \\ m \times n \end{pmatrix} \begin{pmatrix} \sigma_{11} & 0 & \dots \\ 0 & \ddots & \\ \vdots & \ddots & \\ r \times r \end{pmatrix} \begin{pmatrix} v_{11} & \dots & v_{1n} \\ \vdots & \ddots & \\ r \times n \end{pmatrix}$$

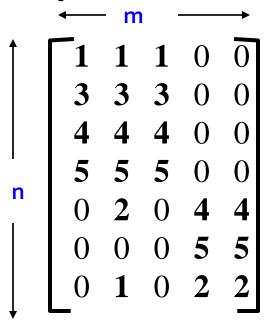
$$\|A-B\|_{F} = \sqrt{\sum_{ij} (A_{ij}-B_{ij})^2}$$

#### **Equivalent:**

'spectral decomposition' of the matrix:

#### **Equivalent:**

#### 'spectral decomposition' of the matrix



#### Why is setting small $\sigma_i$ to 0 the right thing to do?

Vectors  $\mathbf{u}_{i}$  and  $\mathbf{v}_{i}$  are unit length, so  $\mathbf{\sigma}_{i}$ scales them.

So, zeroing small  $\sigma_i$  introduces less error.

Q: How many  $\sigma_s$  to keep?

A: Rule-of-a thumb:

keep 80-90% of 'energy' (= $\sum \sigma_i^2$ )

# **SVD - Complexity**

- To compute SVD:
  - O(nm²) or O(n²m) (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 - Conclusions so far

- SVD:  $A = U \Sigma V^T$ : unique
  - U: user-to-concept similarities
  - V: movie-to-concept similarities
  - lacksquare  $\Sigma$  : strength of each concept
- Dimensionality reduction:
  - keep the few largest singular values (80-90% of 'energy')
  - SVD: picks up linear correlations

# Relation to Eigen-decomposition

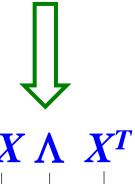
- SVD gives us:
  - $\bullet A = U \Sigma V^T$
- Eigen-decomposition:
  - $A = X \Lambda X^T$ 
    - A is symmetric
    - U, V, X are orthonormal (U<sup>T</sup>U=I),
    - $\Lambda, \Sigma$  are diagonal
- What is:
  - AA<sup>T</sup>=

# Relation to Eigen-decomposition

- SVD gives us:
  - $\bullet A = U \Sigma V^T$
- Eigen-decomposition:
  - $A = X \wedge X^T$ 
    - A is symmetric
    - U, V, X are orthonormal (U<sup>T</sup>U=I),
    - $\Lambda, \Sigma$  are diagonal
- What is:
  - $AA^T = U\Sigma V^T(U\Sigma V^T)^T = U\Sigma V^T(V\Sigma^TU^T) = U\Sigma\Sigma^T U^T$



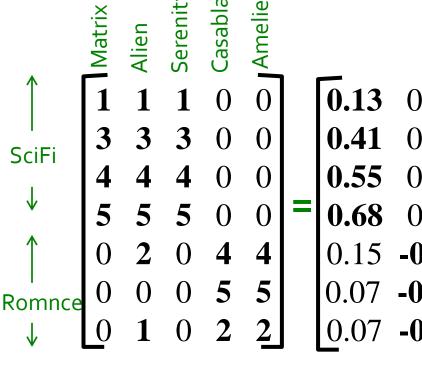
Shows how to compute SVD using eigenvalue decomposition!



So, 
$$\lambda_i = \sigma_i^2$$

# Example of SVD & Conclusion

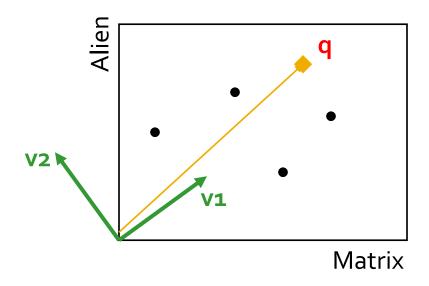
- Q: Find users that like 'Matrix'
- A: Map query into a 'concept space' how?



- Q: Find users that like 'Matrix'
- A: Map query into a 'concept space' how?

#### **Project into concept space:**

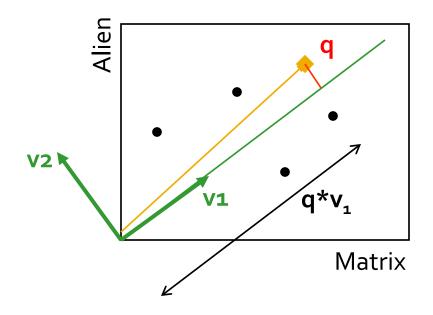
Inner product with each 'concept' vector **v**<sub>i</sub>



- Q: Find users that like 'Matrix'
- A: Map query into a 'concept space' how?

#### **Project into concept space:**

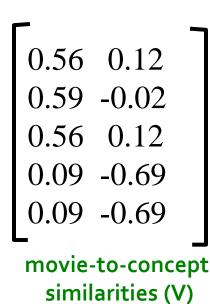
Inner product with each 'concept' vector **v**<sub>i</sub>

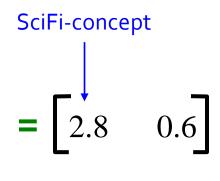


## Compactly, we have:

$$q_{concept} = q V$$

## **E.g.:**



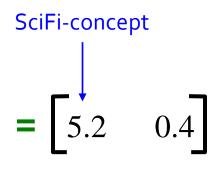


How would the user d that rated ('Alien', 'Serenity') be handled?

$$d_{concept} = d V$$

## **E.g.:**

movie-to-concept similarities (V)



Observation: User d that rated ('Alien', 'Serenity') will be similar to user q that rated ('Matrix'), although d and q have zero ratings in common!

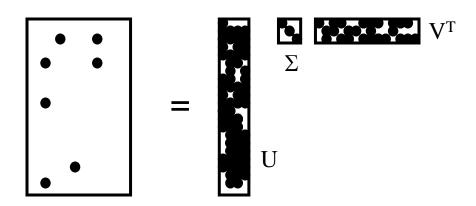
$$\mathbf{d} = \begin{bmatrix} 0 & 4 & 5 & 0 & 0 \end{bmatrix} \qquad \qquad \mathbf{SciFi\text{-}concept}$$

$$\mathbf{q} = \begin{bmatrix} 5 & 0 & 0 & 0 & 0 \end{bmatrix} \qquad \qquad \mathbf{5.2} \qquad \mathbf{5.2} \qquad \mathbf{0.4}$$

$$\mathbf{Zero \ ratings \ in \ common} \qquad \qquad \mathbf{Similarity} \neq \mathbf{0}$$

## **SVD: Drawbacks**

- Optimal low-rank approximation in terms of Frobenius norm
- Interpretability problem:
  - A singular vector specifies a linear combination of all input columns or rows
- Lack of sparsity:
  - Singular vectors are dense!



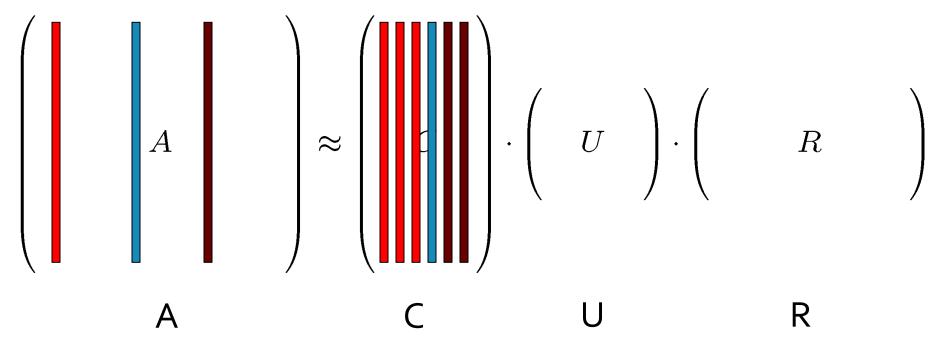
## **Announcements:**

HW2 has been posted

# **CUR Decomposition**

# **CUR Decomposition**

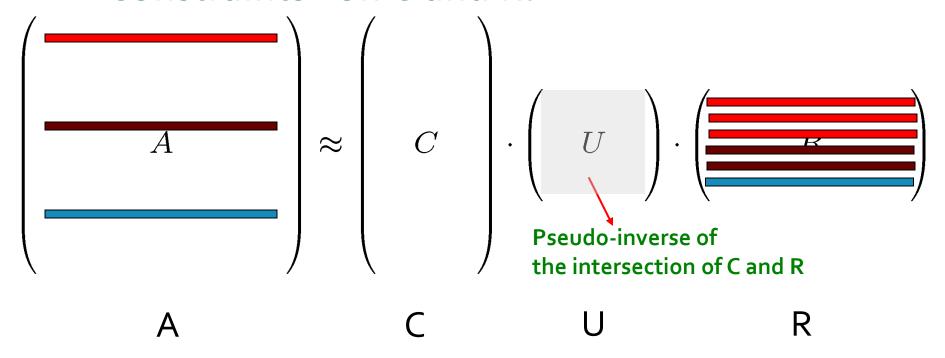
- Goal: Express A as a product of matrices C,U,R
   Make ||A-C·U·R||<sub>F</sub> small
- "Constraints" on C and R:



## **CUR Decomposition**

Frobenius norm: 
$$\|X\|_F = \sqrt{\sum_{ij} X_{ij}^2}$$

- Goal: Express A as a product of matrices C,U,R
   Make ||A-C·U·R||<sub>F</sub> small
- "Constraints" on C and R:



# CUR: Provably good approx. to SVD

#### Let:

 $A_k$  be the "best" rank k approximation to A (that is,  $A_k$  is SVD of A)

## Theorem [Drineas et al.]

**CUR** in  $O(\mathbf{m} \cdot \mathbf{n})$  time achieves

- $\|\mathbf{A}\text{-CUR}\|_{F} \leq \|\mathbf{A}\text{-}\mathbf{A}_{k}\|_{F} + \epsilon \|\mathbf{A}\|_{F}$  with probability at least 1- $\delta$ , by picking
- O(k log(1/ $\delta$ )/ $\epsilon^2$ ) columns, and
- $O(k^2 \log^3(1/\delta)/\epsilon^6)$  rows

In practice:
Pick 4k cols/rows

## **CUR: How it Works**

## Sampling columns (similarly for rows):

**Input**: matrix  $\mathbf{A} \in \mathbb{R}^{m \times n}$ , sample size c

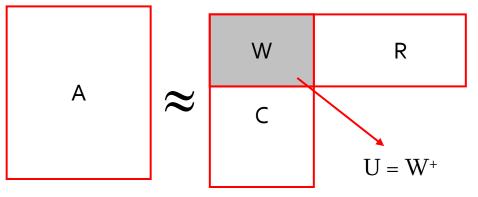
Output:  $\mathbf{C}_d \in \mathbb{R}^{m \times c}$ 

- 1. for x = 1 : n [column distribution]
- 2.  $P(x) = \sum_{i} \mathbf{A}(i, x)^{2} / \sum_{i,j} \mathbf{A}(i, j)^{2}$
- 3. for i = 1 : c [sample columns]
- 4. Pick  $j \in 1 : n$  based on distribution P(x)
- 5. Compute  $\mathbf{C}_d(:,i) = \mathbf{A}(:,j)/\sqrt{cP(j)}$

Note this is a randomized algorithm, same column can be sampled more than once

# Computing U

- Let W be the "intersection" of sampled columns C and rows R
  - Let SVD of W = X Z Y<sup>T</sup>
- Then: U = W<sup>+</sup> = Y Z<sup>+</sup> X<sup>T</sup>
  - $Z^+$ : reciprocals of non-zero singular values:  $Z^+_{ii} = 1/Z_{ii}$
  - W<sup>+</sup> is the "pseudoinverse"



#### Why pseudoinverse works?

W = X Z Y then W<sup>-1</sup> = X<sup>-1</sup> Z<sup>-1</sup> Y<sup>-1</sup>
Due to orthonomality  $X^{-1}=X^{T}$  and  $Y^{-1}=Y^{T}$ Since Z is diagonal  $Z^{-1}=1/Z_{ii}$ Thus, if W is nonsingular,
pseudoinverse is the true inverse

# CUR: Provably good approx. to SVD

## For example:

- Select  $c = O\left(\frac{k \log k}{\varepsilon^2}\right)$  columns of A using ColumnSelect algorithm
- Select  $r = O\left(\frac{k \log k}{\varepsilon^2}\right)$  rows of A using ColumnSelect algorithm
- Set  $U = W^+$
- Then:  $||A CUR||_F \le (2 + \epsilon) ||A A_k||_F$  with probability 98%

## **CUR: Pros & Cons**

## + Easy interpretation

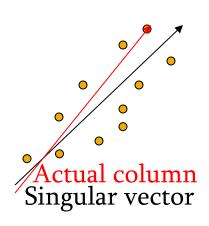
Since the basis vectors are actual columns and rows

## Sparse basis

Since the basis vectors are actual columns and rows

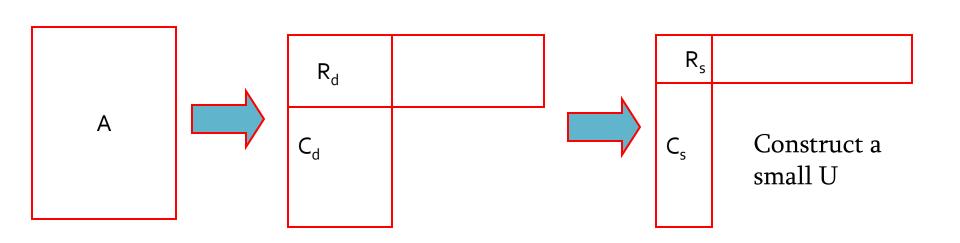


 Columns of large norms will be sampled many times

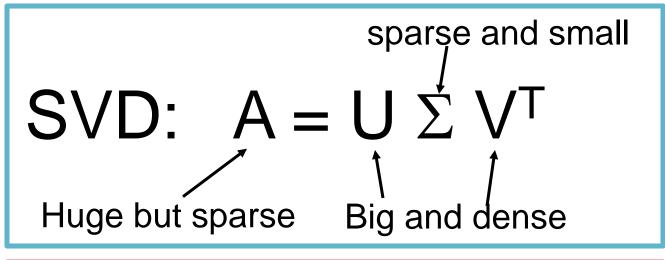


## Solution

- If we want to get rid of the duplicates:
  - Throw them away
  - Scale (multiply) the columns/rows by the square root of the number of duplicates



## SVD vs. CUR

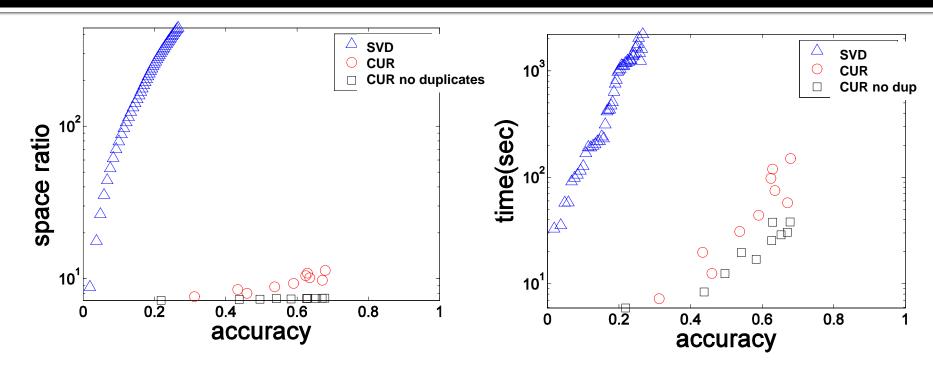


# Simple Experiment

## DBLP bibliographic data

- Author-to-conference big sparse matrix
- A<sub>ij</sub>: Number of papers published by author *i* at conference *j*
- 428K authors (rows), 3659 conferences (columns)
  - Very sparse
- Want to reduce dimensionality
  - How much time does it take?
  - What is the reconstruction error?
  - How much space do we need?

# Results: DBLP- big sparse matrix



#### Accuracy:

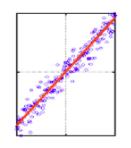
- 1 relative sum squared errors
- Space ratio:
  - #output matrix entries / #input matrix entries
- CPU time

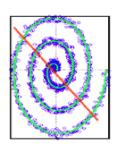
Sun, Faloutsos: Less is More: Compact Matrix Decomposition for Large Sparse Graphs, SDM '07.

# What about linearity assumption?

## SVD is limited to linear projections:

 Lower-dimensional linear projection that preserves Euclidean distances

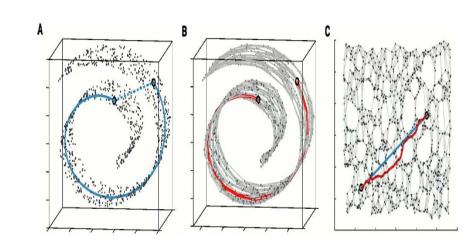




- Non-linear methods: Isomap
  - Data lies on a nonlinear low-dim curve aka manifold
    - Use the distance as measured along the manifold

#### How?

- Build adjacency graph
- Geodesic distance is graph distance
- SVD/PCA the graph pairwise distance matrix



# Further Reading: CUR

- Drineas et al., Fast Monte Carlo Algorithms for Matrices III: Computing a Compressed Approximate Matrix Decomposition, SIAM Journal on Computing, 2006.
- J. Sun, Y. Xie, H. Zhang, C. Faloutsos: Less is More: Compact Matrix Decomposition for Large Sparse Graphs, SDM 2007
- Intra- and interpopulation genotype reconstruction from tagging SNPs, P. Paschou, M. W. Mahoney, A. Javed, J. R. Kidd, A. J. Pakstis, S. Gu, K. K. Kidd, and P. Drineas, Genome Research, 17(1), 96-107 (2007)
- Tensor-CUR Decompositions For Tensor-Based Data, M. W. Mahoney, M. Maggioni, and P. Drineas, Proc. 12-th Annual SIGKDD, 327-336 (2006)