

Data Mining and Matrices

03 – Singular Value Decomposition

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The SVD is the Swiss Army knife of matrix decompositions

—Diane O'Leary, 2006

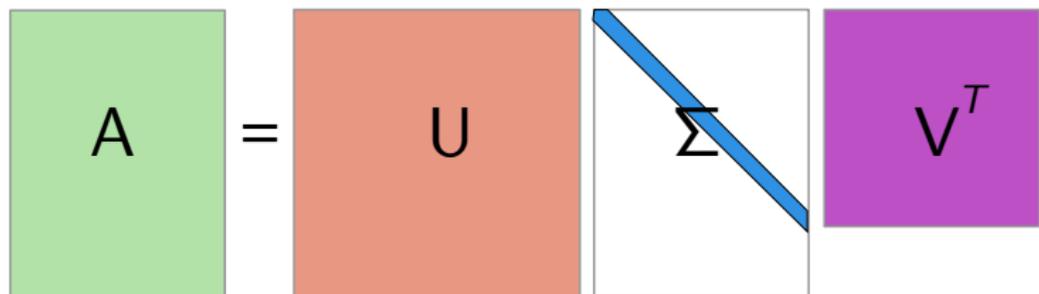
Outline

- 1 The Definition
- 2 Properties of the SVD
- 3 Interpreting SVD
- 4 SVD and Data Analysis
 - How many factors?
 - Using SVD: Data processing and visualization
- 5 Computing the SVD
- 6 Wrap-Up
- 7 About the assignments

The definition

Theorem. For every $\mathbf{A} \in \mathbb{R}^{m \times n}$ there exists $m \times m$ orthogonal matrix \mathbf{U} and $n \times n$ orthogonal matrix \mathbf{V} such that $\mathbf{U}^T \mathbf{A} \mathbf{V}$ is an $m \times n$ diagonal matrix $\mathbf{\Sigma}$ that has values $\sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_{\min\{n,m\}} \geq 0$ in its diagonal.

- I.e. every \mathbf{A} has decomposition $\mathbf{A} = \mathbf{U} \mathbf{\Sigma} \mathbf{V}^T$
 - ▶ The **singular value decomposition** (SVD)
- The values σ_i are the **singular values** of \mathbf{A}
- Columns of \mathbf{U} are the **left singular vectors** and columns of \mathbf{V} the **right singular vectors** of \mathbf{A}



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The fundamental theorem of linear algebra

The **fundamental theorem of linear algebra** states that every matrix $\mathbf{A} \in \mathbb{R}^{m \times n}$ induces four fundamental subspaces:

- The **range** of dimension $\text{rank}(\mathbf{A}) = r$
 - ▶ The set of all possible linear combinations of columns of \mathbf{A}
- The **kernel** of dimension $n - r$
 - ▶ The set of all vectors $\mathbf{x} \in \mathbb{R}^n$ for which $\mathbf{Ax} = \mathbf{0}$
- The **coimage** of dimension r
- The **cokernel** of dimension $m - r$

The bases for these subspaces can be obtained from the SVD:

- Range: the first r columns of \mathbf{U}
- Kernel: the last $(n - r)$ columns of \mathbf{V}
- Coimage: the first r columns of \mathbf{V}
- Cokernel: the last $(m - r)$ columns of \mathbf{U}

Pseudo-inverses

Problem.

Given $\mathbf{A} \in \mathbb{R}^{m \times n}$ and $\mathbf{b} \in \mathbb{R}^m$, find $\mathbf{x} \in \mathbb{R}^n$ minimizing $\|\mathbf{Ax} - \mathbf{b}\|_2$.

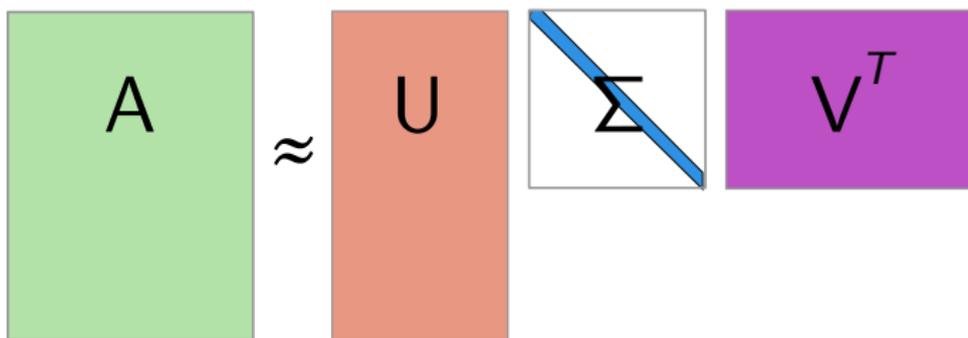
- If \mathbf{A} is invertible, the solution is $\mathbf{A}^{-1}\mathbf{Ax} = \mathbf{A}^{-1}\mathbf{b} \Leftrightarrow \mathbf{x} = \mathbf{A}^{-1}\mathbf{b}$
- A **pseudo-inverse** \mathbf{A}^+ captures some properties of the inverse \mathbf{A}^{-1}
- The **Moose–Penrose pseudo-inverse** of \mathbf{A} is a matrix \mathbf{A}^+ satisfying the following criteria
 - ▶ $\mathbf{AA}^+\mathbf{A} = \mathbf{A}$ (but it is possible that $\mathbf{AA}^+ \neq \mathbf{I}$)
 - ▶ $\mathbf{A}^+\mathbf{AA}^+ = \mathbf{A}^+$ (cf. above)
 - ▶ $(\mathbf{AA}^+)^T = \mathbf{AA}^+$ (\mathbf{AA}^+ is symmetric)
 - ▶ $(\mathbf{A}^+\mathbf{A})^T = \mathbf{A}^+\mathbf{A}$ (as is $\mathbf{A}^+\mathbf{A}$)
- If $\mathbf{A} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^T$ is the SVD of \mathbf{A} , then $\mathbf{A}^+ = \mathbf{V}\mathbf{\Sigma}^{-1}\mathbf{U}^T$
 - ▶ $\mathbf{\Sigma}^{-1}$ replaces σ_i 's with $1/\sigma_i$ and transposes the result

Theorem.

The optimum solution for the above problem can be obtained using $\mathbf{x} = \mathbf{A}^+\mathbf{b}$.

Truncated (thin) SVD

- The rank of the matrix is the number of its non-zero singular values
 - ▶ Easy to see by writing $\mathbf{A} = \sum_{i=1}^{\min\{n,m\}} \sigma_i \mathbf{u}_i \mathbf{v}_i^T$
- The truncated (or thin) SVD only takes the first k columns of \mathbf{U} and \mathbf{V} and the main $k \times k$ submatrix of Σ
 - ▶ $\mathbf{A}_k = \sum_{i=1}^k \sigma_i \mathbf{u}_i \mathbf{v}_i^T = \mathbf{U}_k \Sigma_k \mathbf{V}_k^T$
 - ▶ $\text{rank}(\mathbf{A}_k) = k$ (if $\sigma_k > 0$)
 - ▶ \mathbf{U}_k and \mathbf{V}_k are no more orthogonal, but they are **column-orthogonal**
- The truncated SVD gives a low-rank approximation of \mathbf{A}



SVD and matrix norms

Let $\mathbf{A} = \mathbf{U}\mathbf{\Sigma}\mathbf{V}^T$ be the SVD of \mathbf{A} . Then

- $\|\mathbf{A}\|_F^2 = \sum_{i=1}^{\min\{n,m\}} \sigma_i^2$
- $\|\mathbf{A}\|_2 = \sigma_1$
 - ▶ Remember: $\sigma_1 \geq \sigma_2 \geq \dots \geq \sigma_{\min\{n,m\}} \geq 0$
- Therefore $\|\mathbf{A}\|_2 \leq \|\mathbf{A}\|_F \leq \sqrt{n}\|\mathbf{A}\|_2$
- The Frobenius of the truncated SVD is $\|\mathbf{A}_k\|_F^2 = \sum_{i=1}^k \sigma_i^2$
 - ▶ And the Frobenius of the difference is $\|\mathbf{A} - \mathbf{A}_k\|_F^2 = \sum_{i=k+1}^{\min\{n,m\}} \sigma_i^2$

The Eckart–Young theorem

Let \mathbf{A}_k be the rank- k truncated SVD of \mathbf{A} . Then \mathbf{A}_k is the closest rank- k matrix of \mathbf{A} in the Frobenius sense. That is

$$\|\mathbf{A} - \mathbf{A}_k\|_F \leq \|\mathbf{A} - \mathbf{B}\|_F \quad \text{for all rank-}k \text{ matrices } \mathbf{B}.$$

Eigendecompositions

- An **eigenvector** of a square matrix \mathbf{A} is a vector \mathbf{v} such that \mathbf{A} only changes the magnitude of \mathbf{v}
 - ▶ I.e. $\mathbf{A}\mathbf{v} = \lambda\mathbf{v}$ for some $\lambda \in \mathbb{R}$
 - ▶ Such λ is an **eigenvalue** of \mathbf{A}
- The **eigendecomposition** of \mathbf{A} is $\mathbf{A} = \mathbf{Q}\mathbf{\Delta}\mathbf{Q}^{-1}$
 - ▶ The columns of \mathbf{Q} are the eigenvectors of \mathbf{A}
 - ▶ Matrix $\mathbf{\Delta}$ is a diagonal matrix with the eigenvalues
- Not every (square) matrix has eigendecomposition
 - ▶ If \mathbf{A} is of form $\mathbf{B}\mathbf{B}^T$, it always has eigendecomposition
- The SVD of \mathbf{A} is closely related to the eigendecompositions of $\mathbf{A}\mathbf{A}^T$ and $\mathbf{A}^T\mathbf{A}$
 - ▶ The left singular vectors are the eigenvectors of $\mathbf{A}\mathbf{A}^T$
 - ▶ The right singular vectors are the eigenvectors of $\mathbf{A}^T\mathbf{A}$
 - ▶ The singular values are the square roots of the eigenvalues of both $\mathbf{A}\mathbf{A}^T$ and $\mathbf{A}^T\mathbf{A}$

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Factor interpretation

- The most common way to interpret SVD is to consider the columns of \mathbf{U} (or \mathbf{V})
 - ▶ Let \mathbf{A} be objects-by-attributes and $\mathbf{U}\mathbf{\Sigma}\mathbf{V}^T$ its SVD
 - ▶ If two columns have similar values in a row of \mathbf{V}^T , these attributes are somehow similar (have strong correlation)
 - ▶ If two rows have similar values in a column of \mathbf{U} , these users are somehow similar

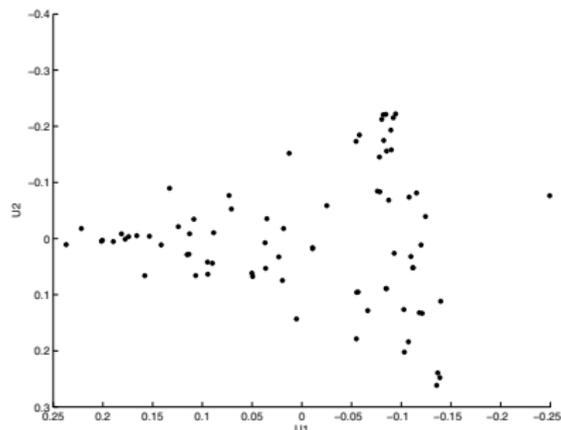
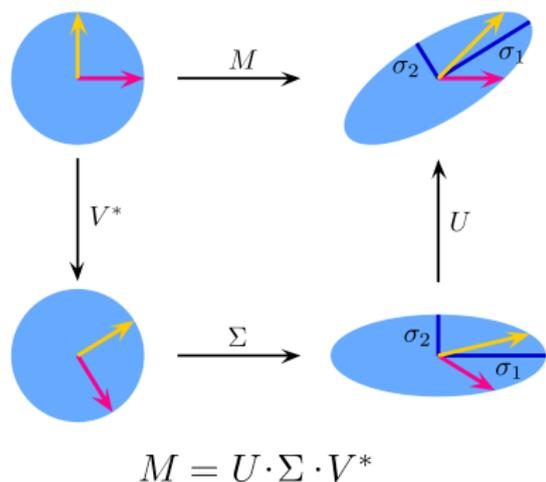


Figure 3.2. The first two factors for a dataset ranking wines.

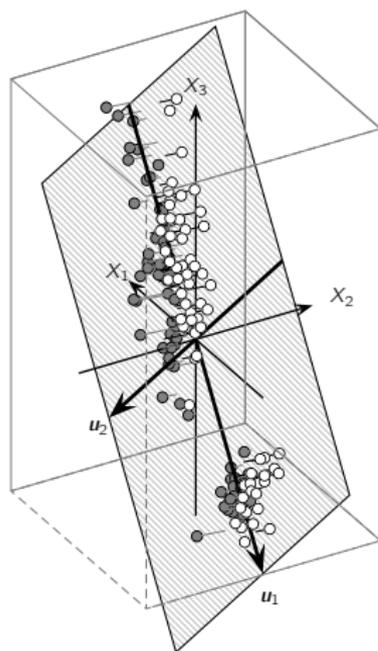
- Example: people's ratings of different wines
- Scatterplot of first and second column of \mathbf{U}
 - ▶ left: likes wine
 - ▶ right: doesn't like
 - ▶ up: likes red wine
 - ▶ bottom: likes white wine
- Conclusion: winelovers like red and white, others care more

Geometric interpretation



- Let $\mathbf{U}\Sigma\mathbf{V}^T$ be the SVD of \mathbf{M}
- SVD shows that every linear mapping $\mathbf{y} = \mathbf{M}\mathbf{x}$ can be considered as a series of rotation, stretching, and rotation operations
 - ▶ Matrix \mathbf{V}^T performs the first rotation $\mathbf{y}_1 = \mathbf{V}^T \mathbf{x}$
 - ▶ Matrix Σ performs the stretching $\mathbf{y}_2 = \Sigma \mathbf{y}_1$
 - ▶ Matrix \mathbf{U} performs the second rotation $\mathbf{y} = \mathbf{U} \mathbf{y}_2$

Dimension of largest variance



- The singular vectors give the dimensions of the variance in the data
 - ▶ The first singular vector is the dimension of the largest variance
 - ▶ The second singular vector is the orthogonal dimension of the second largest variance
 - ★ First two dimensions span a hyperplane
- From Eckart–Young we know that if we project the data to the spanned hyperplanes, the distance of the projection is minimized

Component interpretation

- Recall that we can write $\mathbf{A} = \mathbf{U}\Sigma\mathbf{V}^T = \sum_{i=1}^r \sigma_i \mathbf{u}_i \mathbf{v}_i^T = \sum_{i=1}^r \mathbf{A}_i$
 - ▶ $\mathbf{A}_i = \sigma_i \mathbf{v}_i \mathbf{u}_i^T$
- This explains the data as a sums of (rank-1) layers
 - ▶ The first layer explains the most
 - ▶ The second corrects that by adding and removing smaller values
 - ▶ The third corrects that by adding and removing even smaller values
 - ▶ ...
- The layers don't have to be very intuitive

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Problem

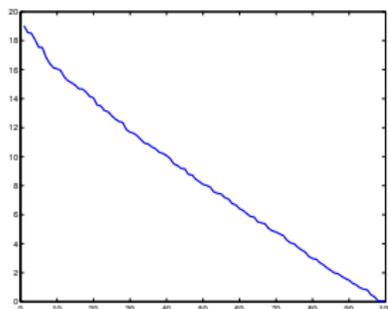
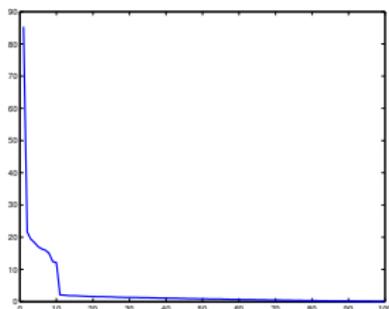
- Most data mining applications do not use full SVD, but truncated SVD
 - ▶ To concentrate on “the most important parts”
- But how to select the rank k of the truncated SVD?
 - ▶ What is important, what is unimportant?
 - ▶ What is structure, what is noise?
 - ▶ Too small rank: all subtlety is lost
 - ▶ Too big rank: all smoothing is lost
- Typical methods rely on singular values in a way or another

Guttman–Kaiser criterion and captured energy

- Perhaps the oldest method is the Guttman–Kaiser criterion:
 - ▶ Select k so that for all $i > k$, $\sigma_i < 1$
 - ▶ Motivation: all components with singular value less than unit are uninteresting
- Another common method is to select enough singular values such that the sum of their squares is 90% of the total sum of the squared singular values
 - ▶ The exact percentage can be different (80%, 95%)
 - ▶ Motivation: The resulting matrix “explains” 90% of the Frobenius norm of the matrix (a.k.a. energy)
- **Problem:** Both of these methods are based on arbitrary thresholds and do not consider the “shape” of the data

Cattell's Scree test

- The **scree plot** plots the singular values in decreasing order
 - ▶ The plot looks like a side of the hill, thence the name
- The scree test is a subjective decision on the rank based on the shape of the scree plot
- The rank should be set to a point where
 - ▶ there is a clear drop in the magnitudes of the singular values; or
 - ▶ the singular values start to even out
- **Problem:** Scree test is subjective, and many data don't have any clear shapes to use (or have many)
 - ▶ Automated methods have been developed to detect the shapes from the scree plot



Entropy-based method

- Consider the relative contribution of each singular value to the overall Frobenius norm
 - ▶ Relative contribution of σ_k is $f_k = \sigma_k^2 / \sum_i \sigma_i^2$
- We can consider these as probabilities and define the (normalized) **entropy** of the singular values as

$$E = -\frac{1}{\log(\min\{n, m\})} \sum_{i=1}^{\min\{n, m\}} f_i \log f_i$$

- ▶ The basis of the logarithm doesn't matter
- ▶ We assume that $0 \cdot \infty = 0$
- ▶ Low entropy (close to 0): the first singular value has almost all mass
- ▶ High entropy (close to 1): the singular values are almost equal
- The rank is selected to be the smallest k such that $\sum_{i=1}^k f_i \geq E$
- **Problem:** Why entropy?

Random flip of signs

- Multiply every element of the data \mathbf{A} randomly with either 1 or -1 to get $\tilde{\mathbf{A}}$
 - ▶ The Frobenius norm doesn't change ($\|\mathbf{A}\|_F = \|\tilde{\mathbf{A}}\|_F$)
 - ▶ The spectral norm does change ($\|\mathbf{A}\|_2 \neq \|\tilde{\mathbf{A}}\|_2$)
 - ★ How much this changes depends on how much "structure" \mathbf{A} has
- We try to select k such that the residual matrix contains only noise
 - ▶ The residual matrix contains the last $m - k$ columns of \mathbf{U} , $\min\{n, m\} - k$ singular values, and last $n - k$ rows of \mathbf{V}^T
 - ▶ If \mathbf{A}_{-k} is the residual matrix of \mathbf{A} after rank- k truncated SVD and $\tilde{\mathbf{A}}_{-k}$ is that for the matrix with randomly flipped signs, we select rank k to be such that $(\|\mathbf{A}_{-k}\|_2 - \|\tilde{\mathbf{A}}_{-k}\|_2) / \|\mathbf{A}_{-k}\|_F$ is small
- **Problem:** How small is small?

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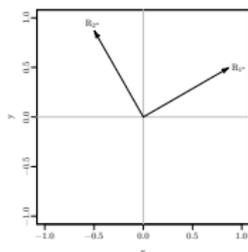
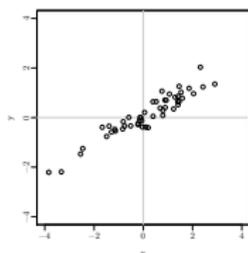
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Normalization

- Data should usually be normalized before SVD is applied
 - ▶ If one attribute is height in meters and other weights in grams, weight seems to carry much more importance in data about humans
 - ▶ If data is all positive, the first singular vector just explains where in the positive quadrant the data is
- The **z-scores** are attributes whose values are transformed by
 - ▶ centering them to 0
 - ★ Remove the mean of the attribute's values from each value
 - ▶ normalizing the magnitudes
 - ★ Divide every value with the standard deviation of the attribute
- Notice that the z-scores assume that
 - ▶ all attributes are equally important
 - ▶ attribute values are approximately normally distributed
- Values that have larger magnitude than importance can also be normalized by first taking logarithms (from positive values) or cubic roots
- The effects of normalization should always be considered

Removing noise

- Very common application of SVD is to remove the noise from the data
- This works simply by taking the truncated SVD from the (normalized) data
 - ▶ The big problem is to select the rank of the truncated SVD
- Example:



- Original data
 - ▶ Looks like 1-dimensional with some noise
- The right singular vectors show the directions
 - ▶ The first looks like the data direction
 - ▶ The second looks like the noise direction
- The singular values confirm this

$$\sigma_1 = 11.73$$

$$\sigma_2 = 1.71$$

Removing dimensions

- Truncated SVD can also be used to battle the **curse of dimensionality**
 - ▶ All points are close to each other in very high dimensional spaces
 - ▶ High dimensionality slows down the algorithms
- Typical approach is to work in a space spanned by the columns of \mathbf{V}^T
 - ▶ If $\mathbf{U}\mathbf{\Sigma}\mathbf{V}^T$ is the SVD of $\mathbf{A} \in \mathbb{R}^{m \times n}$, project \mathbf{A} to $\mathbf{A}\mathbf{V}_k \in \mathbb{R}^{m \times k}$ where \mathbf{V}_k has the first k columns of \mathbf{V}
 - ▶ This is known as the **Karhunen–Loève transform** (KLT) of the rows of \mathbf{A}
 - ★ Matrix \mathbf{A} must be normalized to z-scores in KLT

Visualization

- Truncated SVD with $k = 2, 3$ allows us to visualize the data
 - ▶ We can plot the projected data points after 2D or 3D Karhunen–Loève transform
 - ▶ Or we can plot the scatter plot of two or three (first, left/right) singular vectors

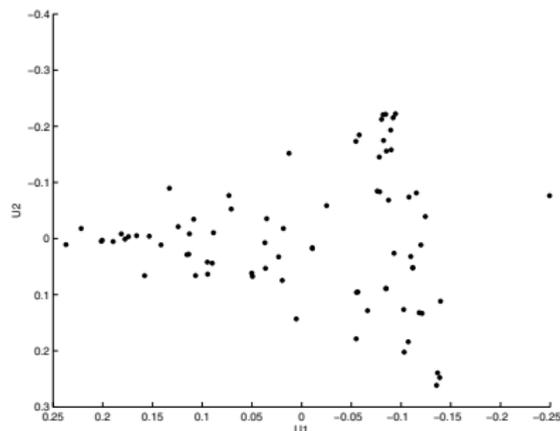
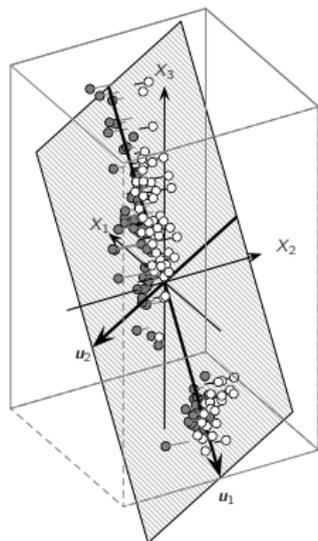


Figure 3.2. *The first two factors for a dataset ranking wines.*



Latent semantic analysis

- The **latent semantic analysis** (LSA) is an information retrieval method that uses SVD
- The data: a term–document matrix \mathbf{A}
 - ▶ the values are (weighted) term frequencies
 - ▶ typically tf/idf values (the frequency of the term in the document divided by the global frequency of the term)
- The truncated SVD $\mathbf{A}_k = \mathbf{U}_k \mathbf{\Sigma}_k \mathbf{V}_k^T$ of \mathbf{A} is computed
 - ▶ Matrix \mathbf{U}_k associates documents to topics
 - ▶ Matrix \mathbf{V}_k associates topics to terms
 - ▶ If two rows of \mathbf{U}_k are similar, the corresponding documents “talk about same things”
- A query q can be answered by considering its term vector \mathbf{q}
 - ▶ \mathbf{q} is projected to $\mathbf{q}_k = \mathbf{q} \mathbf{V} \mathbf{\Sigma}^{-1}$
 - ▶ \mathbf{q}_k is compared to rows of \mathbf{U} and most similar rows are returned

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Algorithms for SVD

- In principle, the SVD of \mathbf{A} can be computed by computing the eigendecomposition of $\mathbf{A}\mathbf{A}^T$
 - ▶ This gives us left singular vectors and squares of singular values
 - ▶ Right singular vectors can be solved: $\mathbf{V}^T = \Sigma^{-1}\mathbf{U}^T\mathbf{A}$
 - ▶ **Bad for numerical stability!**
- Full SVD can be computed in time $O(nm \min\{n, m\})$
 - ▶ Matrix \mathbf{A} is first reduced to a bidiagonal matrix
 - ▶ The SVD of the bidiagonal matrix is computed using iterative methods (similar to eigendecompositions)
- Methods that are faster in practice exist
 - ▶ Especially for truncated SVD
- Efficient implementation of an SVD algorithm requires considerable work and knowledge
 - ▶ Luckily (almost) all numerical computation packages and programs implement SVD

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Lessons learned

- SVD is the Swiss Army knife of (numerical) linear algebra
 - ranks, kernels, norms, ...
- SVD is also very useful in data analysis
 - noise removal, visualization, dimensionality reduction, ...
- Selecting the correct rank for truncated SVD is still a problem

Suggested reading

- Skillicorn, Ch. 3
- Gene H. Golub & Charles F. Van Loan: *Matrix Computations*, 3rd ed. Johns Hopkins University Press, 1996
 - ▶ Excellent source for the algorithms and theory, but very dense

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Basic information

- Assignment sheet will be made available later today/early tomorrow
 - ▶ We'll announce it in the mailing list
- DL in two weeks, delivery by e-mail
 - ▶ Details in the assignment sheet
- Hands-on assignment: data analysis using SVD
- Recommended software: R
 - ▶ Good alternatives: Matlab (commercial), GNU Octave (open source), and Python with NumPy, SciPy, and matplotlib (open source)
 - ▶ Excel is **not** a good alternative (too complicated)
- What you have to return?
 - ▶ Single document that answers to all questions (all figures, all analysis of the results, the main commands you used for the analysis if asked)
 - ▶ Supplementary material containing the transcript of all commands you issued/all source code
 - ▶ **Both files in PDF format**