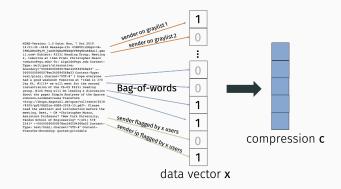
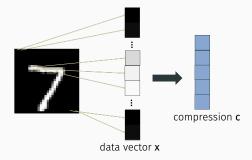
CS-GY 9223 I: Lecture 9 Low-rank approximation and singular value decomposition

NYU Tandon School of Engineering, Prof. Christopher Musco

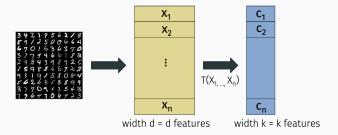
Return to data compression:



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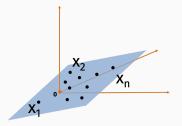
Main difference from randomized methods:



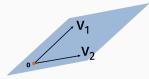
In this section, we will discuss <u>data dependent</u> data transformations. Johnson-Lindenstrauss, MinHash, SimHash were all <u>data oblivious</u>. Advantages of data independent methods:

Advantages of data dependent methods:

Suppose $\mathbf{x}_1, \ldots, \mathbf{x}_n \in \mathbb{R}^d$ lie on a <u>low-dimensional</u> subspace *S* through the origin. I.e. our data set is rank *k* for k < d.



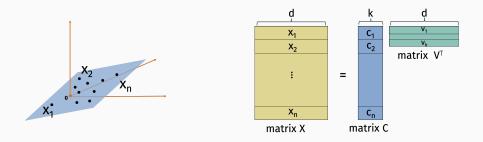
Let $\mathbf{v}_1, \ldots, \mathbf{v}_k$ be orthogonal unit vectors spanning S.



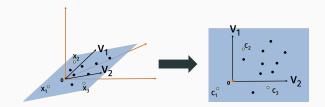
For all *i*, we can write:

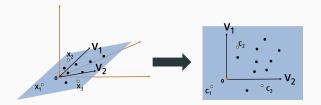
$$\mathbf{X}_i = C_{i,1}\mathbf{V}_1 + \ldots + C_{i,k}\mathbf{V}_k.$$

LOW-RANK DATA



What are $\mathbf{c}_1, \ldots, \mathbf{c}_n$?

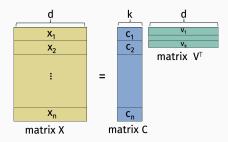




Lots of information preserved:

- $\|\mathbf{x}_{i} \mathbf{x}_{j}\|_{2} = \|\mathbf{c}_{i} \mathbf{c}_{j}\|_{2}$ for all *i*, *j*.
- $\mathbf{x}_i^T \mathbf{x}_j = \mathbf{c}_i^T \mathbf{c}_j$ for all i, j.
- Norms preserved, linear separability preserved, min ||Xy - b|| = min ||Cz - b||, etc., etc.

LOW-RANK DATA



Formally, $\mathbf{C} = \mathbf{X}\mathbf{V}^{\mathsf{T}}$:

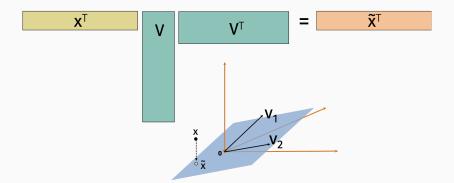
$$X = CV^T \Rightarrow XV = CV^T V$$

Since V's columns are an orthonormal basis, $V^T V = I$.

So $\mathbf{X} = \mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}}$.

PROJECTION MATRICES

VV^{T} is a symmetric projection matrix.



When all data points already lie in the subspace spanned by V's columns, projection doesn't do anything. So $\mathbf{X} = \mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}}$.

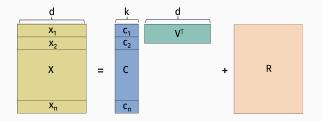
When **X**'s rows lie <u>close</u> to a *k* dimensional subspace, we can still approximate

 $\mathbf{X} \approx \mathbf{X} \mathbf{V} \mathbf{V}^{\mathsf{T}}.$

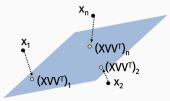
 XVV^{T} is a <u>low-rank approximation</u> for X.

For a given subspace ${\mathcal V}$ spanned by the columns in V,

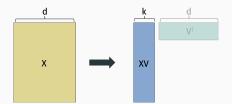
$$\mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}} = \underset{\mathbf{C}}{\arg\min} \|\mathbf{X} - \mathbf{C}\mathbf{V}^{\mathsf{T}}\|_{F}^{2} = \sum_{i,j} (\mathbf{X}_{i,j} - (\mathbf{C}\mathbf{V}^{\mathsf{T}})_{i,j})^{2}.$$



LOW-RANK APPROXIMATION



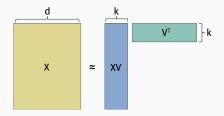
 $\|\mathbf{x}_{i} - \mathbf{x}_{j}\|_{2} \approx \|(\mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}})_{i} - (\mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}})_{j}\|_{2} = \|(\mathbf{X}\mathbf{V}^{\mathsf{T}})_{i} - (\mathbf{X}\mathbf{V}^{\mathsf{T}})_{j}\|_{2}$



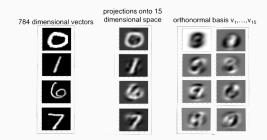
XV can be used as a compressed version of data matrix X.

WHY IS DATA APPROXIMATELY LOW-RANK?

Rows of **X** (data points) are approximately spanned by *k* vectors. Columns of **X** (data features) are approximately spanned by *k* vectors.



If a data set only had *k* unique data points, it would be exactly rank *k*. If it has *k* "clusters" of data points (e.g. the 10 digits) it's often very close to rank *k*.



Colinearity/correlation of data features leads to a low-rank data matrix.

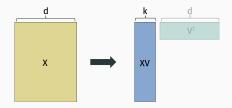
| | bedrooms | bathrooms | sq.ft. | floors | list price | sale price |
|--------|----------|-----------|--------|--------|------------|------------|
| home 1 | 2 | 2 | 1800 | 2 | 200,000 | 195,000 |
| home 2 | 4 | 2.5 | 2700 | 1 | 300,000 | 310,000 |
| | | | | | | |
| | | | | • | • | |
| | • | • | • | • | • | • |
| | • | • | • | • | • | • |
| | | | | | | |
| | | | | | | |
| home n | 5 | 3.5 | 3600 | 3 | 450,000 | 450,000 |

When encoded as a matrix, which image has lower approximate rank?





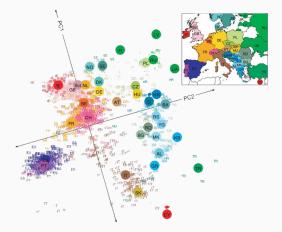
APPLICATIONS OF LOW-RANK APPROXIMATION



- $XV \cdot V^T$ takes O(k(n + d)) space to store instead of O(nd).
- Regression problems involving $XV \cdot V^T$ can be solved in $O(nk^2)$ instead of $O(nd^2)$ time.
- XV can be used for visualization when k = 2, 3.
- We will discuss many more next class.

APPLICATIONS OF LOW-RANK APPROXIMATION

"Genes Mirror Geography Within Europe" – Nature, 2008.



Each data vector \mathbf{x}_i contains genetic information for one person in Europe. Set k = 2 and plot (XV)_i for each *i* on a 2-d plane. Color points by what country they are from.

Given a subspace \mathcal{V} spanned by the k columns in V,

$$\|\mathbf{X} - \mathbf{X}\mathbf{V}\mathbf{V}^T\|_F^2 = \min_{\mathbf{C}} \|\mathbf{X} - \mathbf{C}\mathbf{V}^T\|_F^2$$

We want to find the best $\mathbf{V} \in \mathbb{R}^{d \times k}$:

$$\min_{\substack{\text{orthonormal } \mathbf{V} \in \mathbb{R}^{d \times h}} \|\mathbf{X} - \mathbf{X} \mathbf{V} \mathbf{V}^{\mathsf{T}}\|_{F}^{2}$$
(1)

Note that $\|\mathbf{X} - \mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}}\|_{F}^{2} = \|\mathbf{X}\|_{F}^{2} - \|\mathbf{X}\mathbf{V}\mathbf{V}^{\mathsf{T}}\|_{F}^{2}$ for all orthonormal **V** (since $\mathbf{V}\mathbf{V}^{\mathsf{T}}$ is a projection). Equivalent form:

$$\max_{\text{orthonormal } \mathbf{V} \in \mathbb{R}^{d \times k}} \| \mathbf{X} \mathbf{V} \mathbf{V}^{\mathsf{T}} \|_{F}^{2} = \| \mathbf{X} \mathbf{V} \|_{F}^{2}$$
(2)

If k = 1, want to find a single vector \mathbf{v}_1 which maximizes:

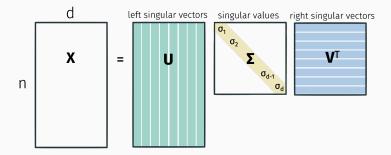
$$\|\mathbf{X}\mathbf{v}_{1}\mathbf{v}_{1}^{T}\|_{F}^{2} = \|\mathbf{X}\mathbf{v}_{1}\|_{F}^{2} = \|\mathbf{X}\mathbf{v}_{1}\|_{2}^{2} = \mathbf{v}_{1}^{T}\mathbf{X}^{T}\mathbf{X}\mathbf{v}_{1}.$$

Choose v_1 to be the top eigenvector of $X^T X$.

What about higher k?

One-stop shop for computing optimal low-rank approximations.

Any matrix **X** can be written:



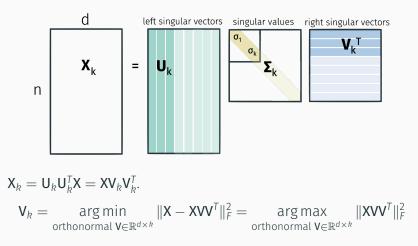
Where $\mathbf{U}^T \mathbf{U} = \mathbf{I}$, $\mathbf{V}^T \mathbf{V} = \mathbf{I}$, and $\sigma_1 \ge \sigma_2 \ge \dots \sigma_d \ge 0$.

CONNECTION TO EIGENDECOMPOSITION

- U contains the orthonormal eigenvectors of XX^{T} .
- V contains the orthonormal eigenvectors of $X^T X$.
- $\sigma_i^2 = \lambda_i(\mathbf{X}\mathbf{X}^T) = \lambda_i(\mathbf{X}^T\mathbf{X})$

This can be checked directly:

Can read off optimal low-rank approximations from the SVD:



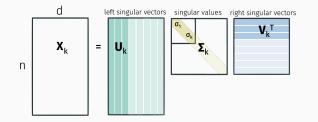
SINGULAR VALUE DECOMPOSITION

- V_k 's columns are called the "top right singular vectors of X"
- $\cdot ~ \mathbf{U}_k$'s columns are called the "top left singular vectors of \mathbf{X} "
- $\sigma_1, \ldots, \sigma_k$ are the "top singular values". $\sigma_1, \ldots, \sigma_d$ are sometimes called the "spectrum of X" (although this is more typically used to refer to eigenvalues).

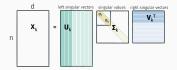
Connection to Principal Component Analysis:

- Let $\bar{\mathbf{X}} = \mathbf{X} \mathbf{1}\boldsymbol{\mu}^{\mathsf{T}}$ where $\boldsymbol{\mu} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_{i}$. I.e. $\bar{\mathbf{X}}$ is obtained by mean centering X's rows.
- Let ŪΣ̄V^T be the SVD of X̄. Ū's first columns are the "top principal components" of X. V's first columns are the "weight vectors" for these principal components.

USEFUL OBSERVATIONS



Observation 1: The optimal compression XV_k has orthogonal columns.



Observation 2: The optimal low-rank approximation error $E_k = \|\mathbf{X} - \mathbf{U}_k \mathbf{U}_k^T \mathbf{X}\|_F^2 = \|\mathbf{X} - \mathbf{X} \mathbf{V}_k \mathbf{V}_k^T\|_F^2$ can be written:

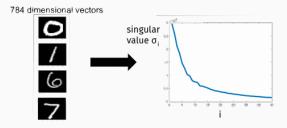
$$E_k = \sum_{i=k+1}^d \sigma_i^2$$

SPECTRAL PLOTS

Observation 2: The optimal low-rank approximation error $E_k = \|\mathbf{X} - \mathbf{U}_k \mathbf{U}_k^T \mathbf{X}\|_F^2 = \|\mathbf{X} - \mathbf{X} \mathbf{V}_k \mathbf{V}_k^T\|_F^2$ can be written:

$$E_k = \sum_{i=k+1}^d \sigma_i^2$$

Can immediately get a sense of "how low-rank" a matrix is from it's spectrum:

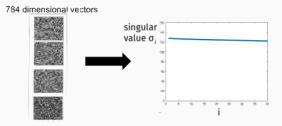


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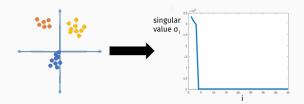


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$$E_k = \sum_{i=k+1}^d \sigma_i^2.$$

Can immediately get a sense of "how low-rank" a matrix is from it's spectrum:



Suffices to compute V. Then $U\Sigma = XV$.

- Compute $\mathbf{X}^{\mathsf{T}}\mathbf{X}$.
- Find eigendecomposition $V \Lambda V^T = X^T X$.
- Compute $\mathbf{L} = \mathbf{XV}$. Set $\sigma_i = \|\mathbf{L}_i\|_2$ and $\mathbf{U}_i = \mathbf{L}_i / \|\mathbf{L}_i\|_2$.

Total runtime \approx

COMPUTING THE SVD (FASTER)

- Use an iterative algorithm.
- Compute <u>approximate</u> solution.
- Only compute top k singular vectors/values. Runtime will depend on k. When k = d we can't do any better than classical algorithms based on eigendecomposition.

What we won't discuss today: sketching methods and stochastic methods (which are faster in some settings).

Today: What about when k = 1? Goal: Find some $\mathbf{z} \approx \mathbf{v}_1$. Input: $\mathbf{X} \in \mathbb{R}^{n \times d}$ with SVD $\mathbf{U} \mathbf{\Sigma} \mathbf{V}$.

Power method:

- Choose $\mathbf{z}^{(0)}$ randomly. E.g. $\mathbf{z}_0 \sim \mathcal{N}(0, 1)$.
- For i = 1, ..., T• $\mathbf{z}^{(i)} = \mathbf{X}^T \cdot (\mathbf{X} \mathbf{z}^{(i-1)})$ • $n_i = \|\mathbf{z}^{(i)}\|_2$ • $\mathbf{z}^{(i)} = \mathbf{z}^{(i)}/n_i$

Return **z**_T

Write $\mathbf{z}^{(0)}$ in the right singular vector basis:

$$\mathbf{z}^{(0)} = c_1 \mathbf{v}_1 + \mathbf{c}_2 \mathbf{v}_2 + \ldots + c_d \mathbf{v}_d$$

Update step: $\mathbf{z}^{(i)} = \mathbf{X}^T \cdot (\mathbf{X}\mathbf{z}^{(i-1)}) = \mathbf{V}\mathbf{\Sigma}^2 \mathbf{V}^T \mathbf{z}^{(i-1)}$ (then normalize) Claim:

$$\mathbf{z}^{(1)} = \frac{1}{n_1} \left[c_1 \cdot \sigma_1^2 \mathbf{v}_1 + \mathbf{c}_2 \cdot \sigma_2^2 \mathbf{v}_2 + \ldots + c_d \cdot \sigma_d^2 \mathbf{v}_d \right]$$

POWER METHOD INTUITION

Claim:

$$\mathbf{z}^{(T)} = \frac{1}{\prod_{i=1}^{T} n_i} \left[c_1 \cdot \sigma_1^{2T} \mathbf{v}_1 + \mathbf{c}_2 \cdot \sigma_2^{2T} \mathbf{v}_2 + \ldots + c_d \cdot \sigma_d^{2T} \mathbf{v}_d \right]$$

Theorem (Basis Power Method Convergence)

Let $\gamma = \frac{\sigma_1 - \sigma_2}{\sigma_1}$ be parameter capturing the "gap" between the first and second largest singular values. If Power Method is initialized with a random Gaussian vector then, with high probability, after $T = O\left(\frac{\log d/\epsilon}{\gamma}\right)$ steps, we have:

$$\|\mathbf{v}_1 - \mathbf{z}^{(T)}\|_2 \le \epsilon.$$

Total runtime:

First observation: For all i

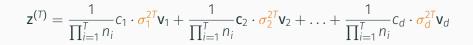
$$O(1/d^2) \le c_i \le O(d)$$

with probability $\frac{1}{d}$. This is a very loose bound, but it's all that we will need. **Prove at home.**

Corollary:

$$\max_j \frac{c_j}{c_1} \leq O(d^3).$$

POWER METHOD FORMAL CONVERGENCE



Since $\mathbf{z}^{(T)}$ is a unit vector, $\sum_{i=1}^{d} \alpha_i^2 = 1$.

• $\alpha_1 \leq 1$.

•
$$\alpha_j^2 \leq (\epsilon/2d)^2$$
 for $j \geq 2$.

• $\alpha_1^2 \ge 1 - d \cdot (\epsilon/2d)^2 \Longrightarrow \alpha_1 \ge 1 - \epsilon/2.$

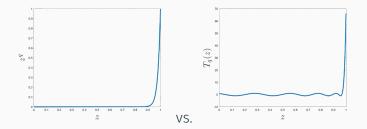
$$\|v_1 - z^{(7)}\|_2 \le$$

Theorem (Basis Power Method Convergence)

If Power Method is initialized with a random Gaussian vector then, with high probability, after $T = O\left(\frac{\log d/\epsilon}{\epsilon}\right)$ steps, we obtain a **z** satisfying:

$$\|\mathbf{X} - \mathbf{X}\mathbf{z}\mathbf{z}^{\mathsf{T}}\|_{F}^{2} \leq (1 + \epsilon)\|\mathbf{X} - \mathbf{X}\mathbf{v}_{1}\mathbf{v}_{1}^{\mathsf{T}}\|_{F}^{2}$$

KRYLOV SUBSPACE METHODS



Lanczos method, Arnoldi method, etc. require $T = O\left(\frac{\log d/\epsilon}{\sqrt{\epsilon}}\right)$ steps for the same guarantee.

GENERALIZATIONS TO LARGE k

- Block Power Method aka Simultaneous Iteration aka Subspace Iteration aka Orthogonal Iteration
- Block Krylov methods

Runtime:
$$O\left(ndk \cdot \frac{\log d/\epsilon}{\sqrt{\epsilon}}\right)$$

to obtain a nearly optimal low-rank approximation.