Lecture 19 – Matrix rank minimization

# Semidefinite programming

A reminder

# Primal Semidefinite Programming Problem

min trace(CX),  
s.t. trace(A<sub>i</sub>X) = b<sub>i</sub>, i = 1,..., m  

$$X \in \mathbf{S}^n \ X \succeq 0$$
  
 $C, A_i \in \mathbf{S}^n, b \in \mathbf{R}^m$ .

SDP cone 
$$K = \{x \in \mathbf{S}^n : X \succeq 0\}$$
 - self dual.

# **Dual Semidefinite Programming Problem**

max 
$$b^T y$$
,
s.t.  $\sum_{i=1}^{m} y_i A_i + S = C$ 
 $S \succeq 0$ .



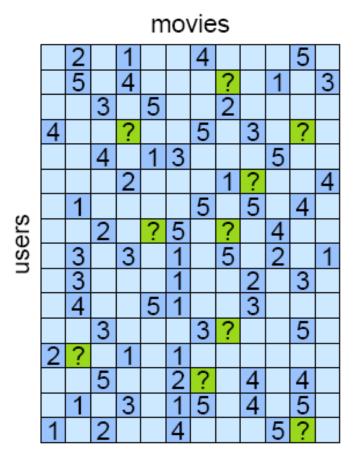
- Some users rate some movies they watched (or didn't!)
- Predict the rating (1..5) for each user/ movie pair.
- Use this prediction to recommend users the movies that they would like

#### Matrix completion problem, collaborative filtering

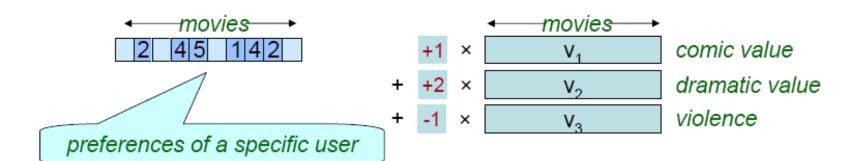
Collaborative filtering: famous Netflix challenge

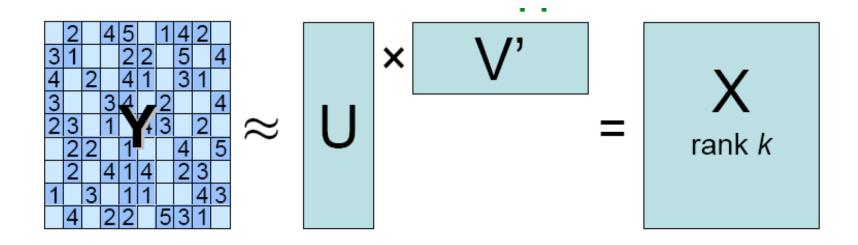
Will user i like movie j?

Complete the matrix based on partially filled information.



#### Linear factor model





#### Convex relaxation via nuclear norm

 Given the values for a subset of entries, find the matrix with these entries and the smallest (or given) rank.

$$\min_{X \in \mathbf{R}^{m \times n}} \quad \mathbf{rank}(X)$$
s.t.  $X_{ij} = M_{ij}, \ (i, j) \in I$ 

NP-hard problem.

$$\operatorname{rank}(X) = \|\sigma(X)\|_0$$
, where  $\sigma(X)$  is the vector of the singular values.

 $\|\cdot\|_0$ ,  $\Rightarrow \|\cdot\|_1$  - the tightest convex relaxation.

Convex relaxation: 
$$\|\sigma(X)\|_1 = \sum_{i=1}^n \sigma_i(X)$$

#### Under suitable randomness hypothesis

#### **ACMRM**

Recht, Fazel and Parrilo, 2007:
 For fixed 0 < δ < 1, when p = O((m + n)r log(mn)), with high probability, A satisfies the Restricted Isometry Property (RIP):</li>

$$(1 - \delta_r(A)) \|X\|_F \le \|AX\|_2 \le (1 + \delta_r(A)) \|X\|_F$$

with  $\delta_r(A) \leq \delta$  for all matrices X of rank r.

#### Matrix Completion

- Candès and Recht, 2008:  $O(n^{1.2}r \log n)$
- Candès and Tao, 2009: O(nrpoly log n)

#### Convex relaxation via nuclear norm

 Given the values for a subset of entries, find the matrix with these entries and the smallest "nuclear norm".

$$\min_{X \in \mathbf{R}^{m \times n}} ||X||_*$$
s.t. 
$$X_{ij} = M_{ij}, \ (i, j) \in I$$

Convex problem

$$||X||_* = ||\sigma(X)||_1 = \sum_{i=1}^n \sigma_i(X)$$

Convex Cone

$$||X||_* \leq t$$

## Trace norm properties

**Definition 1.** The trace norm<sup>1</sup>  $||X||_{\Sigma}$  is the sum of the singular values of X.

$$\mathbf{Lemma} \ \mathbf{1.} \ \|X\|_{\Sigma} = \min_{X = UV'} \|U\|_{Fro} \, \|V\|_{Fro} = \min_{X = UV'} \tfrac{1}{2} (\|U\|_{Fro}^2 + \|V\|_{Fro}^2)$$

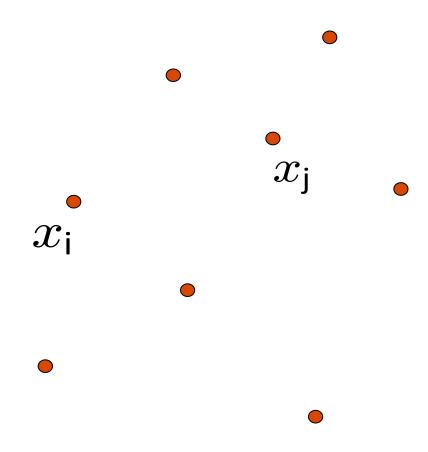
**Lemma 3 ([7, Lemma 1]).** For any  $X \in \mathbb{R}^{n \times m}$  and  $t \in \mathbb{R}$ :  $||X||_{\Sigma} \leq t$  iff there exists  $A \in \mathbb{R}^{n \times n}$  and  $B \in \mathbb{R}^{m \times m}$  such that  $2 \begin{bmatrix} A & X \\ X' & B \end{bmatrix} \succcurlyeq 0$  and  $\operatorname{tr} A + \operatorname{tr} B \leq 2t$ .

*Proof.* Note that for any matrix W,  $\|W\|_{\operatorname{Fro}} = \operatorname{tr} WW'$ . If  $\begin{bmatrix} A & X \\ X' & B \end{bmatrix} \succcurlyeq 0$ , we can write it as a product  $\begin{bmatrix} U \\ V \end{bmatrix} \begin{bmatrix} U' & V' \end{bmatrix}$ . We have X = UV' and  $\frac{1}{2}(\|U\|_{\operatorname{Fro}}^2 + \|V\|_{\operatorname{Fro}}^2) = \frac{1}{2}(\operatorname{tr} A + \operatorname{tr} B) \le t$ , establishing  $\|X\|_{\Sigma} \le t$ . Conversely, if  $\|X\|_{\Sigma} \le t$  we can write it as X = UV' with  $\operatorname{tr} UU' + \operatorname{tr} VV' \le 2t$  and consider the p.s.d. matrix  $\begin{bmatrix} UU' & X \\ X' & VV' \end{bmatrix}$ .

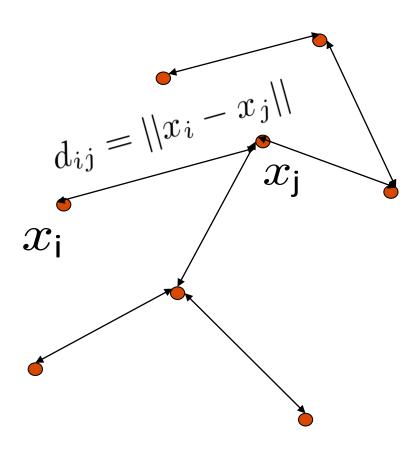
#### Matrix Completion formulation

min trace 
$$\begin{pmatrix} W_1 & X \\ X^{\top} & W_2 \end{pmatrix}$$
  
s.t.  $X_{ij} = M_{ij}, \ (i,j) \in I$   
 $\begin{pmatrix} W_1 & X \\ X^{\top} & W_2 \end{pmatrix} \succeq 0$   
 $X \in \mathbf{R}^{m \times n},$   
 $W_1 \in \mathbf{R}^{m \times m}, \ W_2 \in \mathbf{R}^{n \times n}$ 

## Sensor network localization



## Sensor network localization



#### SDP relaxation of sensor network localization problem

 Given partial information on pair-wise distances find all distances and the exact locations of sensors.

$$d_{ij} = ||x_i - x_j||^2 = (x_i - x_j)^\top (x_i - x_j)$$
Looking for matrix  $X \in \mathbf{R}^{n \times 3}$ 
such that for all pairs  $(i, j)$ 
for which  $d_{ij}$  is known

Find 
$$Y = XX^{\top}$$
:  $Y_{ij} = x_i^{\top} x_j$   
linear constaints  $Y_{ii} + Y_{jj} - 2Y_{ij} = d_{ij}$ , Y-low rank

 $||x_i - x_j||^2 = ||x_i||^2 + ||x_j||^2 - 2x_i^\top x_j$ 

#### Convex relaxation via trace

 Given the distances between some elements, find the matrix with a given rank (2 or 3)

rank
$$(Y) = 2 (3)$$
  
s.t.  $Y_{ii} + Y_{jj} - 2Y_{ij} = d_{ij}, (i, j) \in I$   
 $Y \succeq 0$ 

NP-hard problem.

$$\operatorname{rank}(Y) = \|\lambda(Y)\|_0$$
, where  $\lambda(Y)$  is the vector of eignevalues values of Y.

Convex relaxation: 
$$\|\lambda(Y)\|_1 = \sum_{i=1}^n \lambda_i(Y) = \operatorname{trace}(Y)$$

#### SDP relaxation for sensor network localization

min trace(Y)  
s.t. 
$$Y_{ii} + Y_{jj} - 2Y_{ij} = d_{ij}, (i, j) \in I$$
  
 $Y \succeq 0$   
 $Y \in \mathbf{R}^{n \times n}$ 

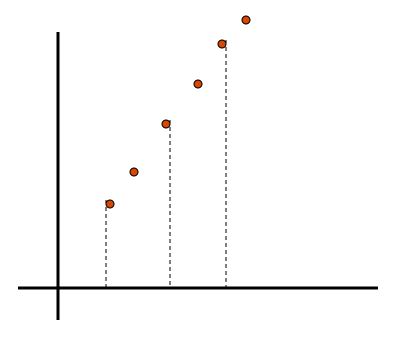
## Metric multidimensional scaling

Let us take three points in  $\mathbb{R}^2$ :

$$x_1 = (2, 1)$$
 $x_2 = (4, 2)$ 
 $x_3 = (6, 2)$ 

$$x_2 = (4,2)$$

$$x_3 = (6, 2)$$



$$Y = XX^{\top} = \begin{bmatrix} 5 & 10 & 15 \\ 10 & 20 & 30 \\ 15 & 30 & 45 \end{bmatrix} = \begin{bmatrix} \sqrt{5} \\ 2\sqrt{5} \\ 3\sqrt{5} \end{bmatrix} \begin{bmatrix} \sqrt{5} & 2\sqrt{5} & 3\sqrt{5} \end{bmatrix}$$

Preserving the inner products  $x_i^T x_i$  we preserve distances  $||x_i - x_i||^2$ 

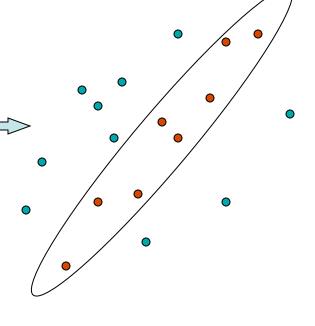
## Dimensionality reduction

# **Principal Component Analysis**

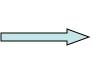
Select few largest eigenvectors of Y=X<sup>T</sup>X/n

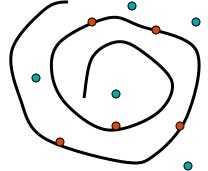
## **Multidimensional Metric Scaling**

Select few largest eigenvectors of Y=XX<sup>T</sup>









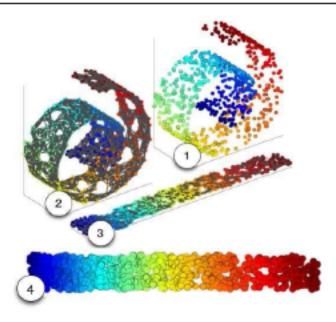


Figure 1. The problem of manifold learning, illustrated for N=800 data points sampled from a "Swiss roll". (1). A discretized manifold is revealed by connecting each data point and its k=6 nearest neighbors (2). An unsupervised learning algorithm unfolds the Swiss roll while preserving the local geometry of nearby data points (3). Finally, the data points are projected onto the two dimensional subspace that maximizes their variance, yielding a faithful embedding of the original manifold (4).

## Nonlinear dimensionality reduction

Preserve pair-wise distances between neighboring points.

$$d_{ij} = ||x_i - x_j||^2 = (x_i - x_j)^{\top} (x_i - x_j), \text{ for } (i, j) \in I$$

 $x_{\mathsf{i}}$ 

Looking for matrix  $Z \in \mathbf{R}^{n \times k}$  (for some k) such that for all pairs  $(i, j) \in I$   $||z_i - z_j||^2 = ||z_i||^2 + ||z_j||^2 - 2z_i^{\mathsf{T}} z_j$ 

Find 
$$Y = ZZ^{\top}$$
:  $Y_{ij} = z_i^{\top} z_j$   
constaints  $Y_{ii} + Y_{jj} - 2Y_{ij} = d_{ij}, Y \succeq 0$ 

## SDP formulation for nonlinear dimensionality reduction

max 
$$\operatorname{trace}(Y)$$

$$\sum_{ij} Y_{ij} = 0$$
s.t.  $Y_{ii} + Y_{jj} - 2Y_{ij} = d_{ij}, \ (i, j) \in I$ 

$$Y \succeq 0$$

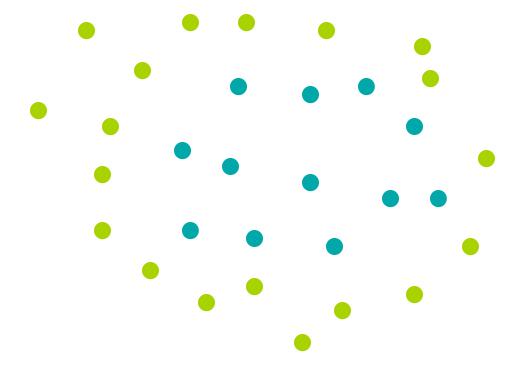
$$Y \in \mathbf{R}^{n \times n}$$

Distance metric learning

# Gaussian Kernel

$$K(x_i, x_j) = e^{-\frac{\|x_i - x_j\|^2}{2\sigma}}$$

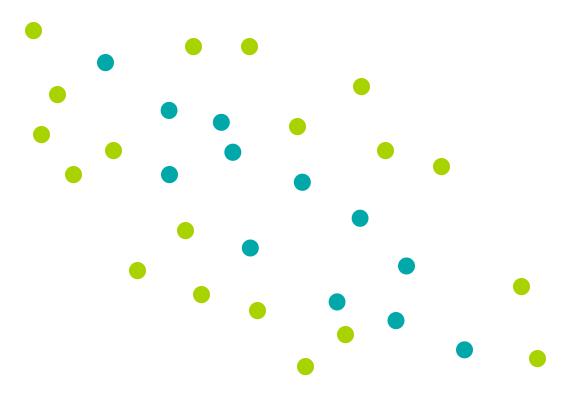
Euclidean distance 
$$||x_i - x_j||^2 = (x_i - x_j)^\top (x_i - x_j)$$



# Gaussian Kernel

$$K(x_i, x_j) = e^{-\frac{\|x_i - x_j\|^2}{2\sigma}}$$

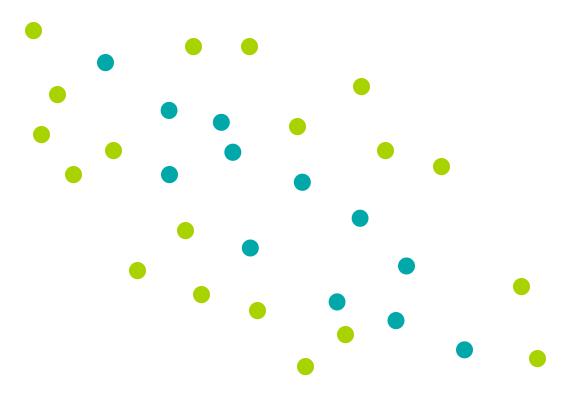
Euclidean distance 
$$||x_i - x_j||^2 = (x_i - x_j)^\top (x_i - x_j)$$



# Gaussian Kernel

$$K(x_i, x_j) = e^{-\frac{\|x_i - x_j\|_M^2}{2\sigma}}$$

Mahalanobis distance 
$$||x_i - x_j||_M^2 = (x_i - x_j)^\top M(x_i - x_j)$$



# Distance Metric Learning

$$\max_{M} \sum_{\substack{(i,j) \in D}} (x_i - x_j)^{\top} M(x_i - x_j)$$
s.t. 
$$\sum_{\substack{(i,j) \in S \\ M \succeq 0}} (x_i - x_j)^{\top} M(x_i - x_j) \leqslant 1$$

S – the set of similarly labeled examples,  $card(S) \sim O(n^2)$ 

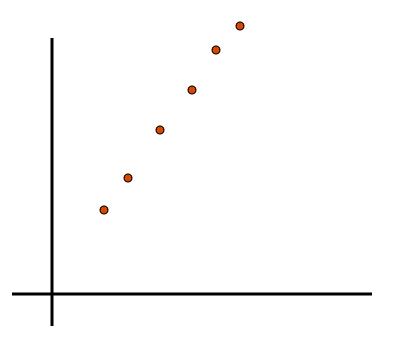
D – the set of differently labeled examples,  $card(D)\sim O(n^2)$ 

IPM is too expensive, need a first order method approach

## Principal component analysis

Let us take three points in  $\mathbb{R}^2$ :

$$x_1 = (2,1)$$
 $x_2 = (4,2)$ 
 $x_3 = (6,2)$ 



$$Y = \frac{1}{3} \sum_{i=1}^{3} x_i x_i^{\top} = \frac{1}{3} \left( \begin{bmatrix} 4 & 2 \\ 2 & 1 \end{bmatrix} + \begin{bmatrix} 16 & 8 \\ 8 & 4 \end{bmatrix} + \begin{bmatrix} 36 & 18 \\ 18 & 9 \end{bmatrix} \right)$$

$$Y = \frac{1}{3} \sum_{i=1}^{3} x_i x_i^{\top} = \frac{1}{3} \begin{bmatrix} 56 & 28 \\ 28 & 14 \end{bmatrix} = \frac{14}{3} \begin{bmatrix} 2 \\ 1 \end{bmatrix} \begin{bmatrix} 2 & 1 \end{bmatrix}$$

## Principal component analysis

#### Find direction of largest variance

Let us take three points in  $\mathbb{R}^2$ :

$$y_1 = (2,1) + (O(\epsilon), O(\epsilon))$$
  
 $y_2 = (4,2) + (O(\epsilon), O(\epsilon))$ 

$$y_3 = (6,3) + (O(\epsilon), O(\epsilon))$$

$$A = \frac{14}{\sqrt{3}} \begin{bmatrix} 2/\sqrt{3} \\ 1/\sqrt{3} \end{bmatrix} \begin{bmatrix} 2/\sqrt{3} & 1/\sqrt{3} \end{bmatrix} + \begin{bmatrix} O(\epsilon) & O(\epsilon) \\ O(\epsilon) & O(\epsilon) \end{bmatrix}$$

$$\begin{bmatrix} 2/\sqrt{3} \\ 1/\sqrt{3} \end{bmatrix} = \operatorname{argmax}_{x \in \mathbf{R}^2, \|x\| = 1} x^{\top} A x$$

## Sparse principal component analysis

#### Find a sparse direction of largest variance

Let us take three points in  $\mathbb{R}^2$ :

$$y_1 = (2,1) + (O(\epsilon), O(\epsilon))$$
  
 $y_2 = (4,2) + (O(\epsilon), O(\epsilon))$ 

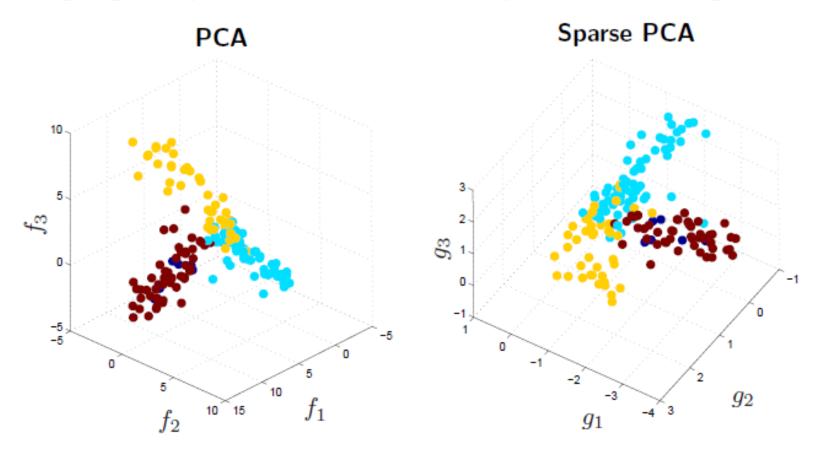
$$y_3 = (6,3) + (O(\epsilon), O(\epsilon))$$

$$A = \frac{1}{3} \begin{bmatrix} 56 + O(\epsilon) & 28 + O(\epsilon) \\ 28 + O(\epsilon) & 14 + O(\epsilon) \end{bmatrix} \approx \frac{56}{3} \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 1 & 0 \end{bmatrix} + \begin{bmatrix} O(\epsilon) & O(\epsilon) \\ O(\epsilon) & O(\epsilon) \end{bmatrix}$$

$$\begin{bmatrix} 1 \\ 0 \end{bmatrix} = \operatorname{argmax}_{x \in \mathbf{R}^2, \|x\| = 1, \operatorname{card}(x) = 1} x^{\top} A x$$

#### Introduction

Clustering of gene expression data in PCA versus sparse PCA, on 500 genes.



The PCA factors  $f_i$  on the left are dense and each use all 500 genes. The sparse factors  $g_1$ ,  $g_2$  and  $g_3$  on the right involve 6, 4 and 4 genes respectively.

## Sparse PCA

Given a set  $Y \in \mathbf{R}^m \times n$  compute empirical covariance matrix  $A = \frac{1}{m} Y^{\top} Y$ 

#### Principal component analysis

Maximize the variance explained by factor x

$$\max_{x \in \mathbf{R}^n} \quad x^{\top} A x$$

s.t. 
$$||x||_2 = 1$$

# **Sparse** principal component analysis

Maximize the variance explained by a factor x with bounded cardinality

$$\max_{x \in \mathbf{R}^n} \quad x^{\top} A x$$
  
s.t. 
$$card(x) = k$$
  
$$||x||_2 = 1$$

#### Semidefinite relaxation

Start from:

maximize 
$$x^T A x$$
  
subject to  $||x||_2 = 1$   
 $\operatorname{Card}(x) \le k$ ,

where  $x \in \mathbb{R}^n$ . Let  $X = xx^T$  and write everything in terms of the matrix X:

Replace  $X = xx^T$  by the equivalent  $X \succeq 0$ ,  $\operatorname{Rank}(X) = 1$ :

again, this is the same problem.

#### Semidefinite relaxation

We have made some progress:

- The objective Tr(AX) is now linear in X
- The (non-convex) constraint  $||x||_2 = 1$  became a linear constraint Tr(X) = 1.

But this is still a hard problem:

- The  $Card(X) \leq k^2$  is still non-convex.
- So is the constraint  $\operatorname{Rank}(X) = 1$ .

We still need to relax the two non-convex constraints above:

- If  $u \in \mathbb{R}^p$ , Card(u) = q implies  $||u||_1 \le \sqrt{q}||u||_2$ . So we can replace  $\operatorname{Card}(X) \leq k^2$  by the weaker (but convex):  $\mathbf{1}^T |X| \mathbf{1} \leq k$ .
- We simply drop the rank constraint

#### Semidefinite Programming

Semidefinite relaxation:

$$\begin{array}{lll} \text{maximize} & x^TAx \\ \text{subject to} & \|x\|_2 = 1 \\ & \operatorname{Card}(x) \leq k, \end{array} \qquad \begin{array}{ll} \text{becomes} & \operatorname{maximize} & \operatorname{Tr}(AX) \\ \text{subject to} & \operatorname{Tr}(X) = 1 \\ & 1^T|X|1 \leq k \\ & X \succeq 0, \end{array}$$

- This is a semidefinite program in the variable  $X \in \mathbf{S}^n$ ...
- Solve small problems (a few hundred variables) using IP solvers, etc.
- Dimensionality reduction apps: solve very large instances.

Solution: use first order algorithm. . .