

# **Sparse Principal Component Analysis using Semidefinite Programming**

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Support from NSF, DHS and Google.

# Introduction

## Principal Component Analysis (PCA)

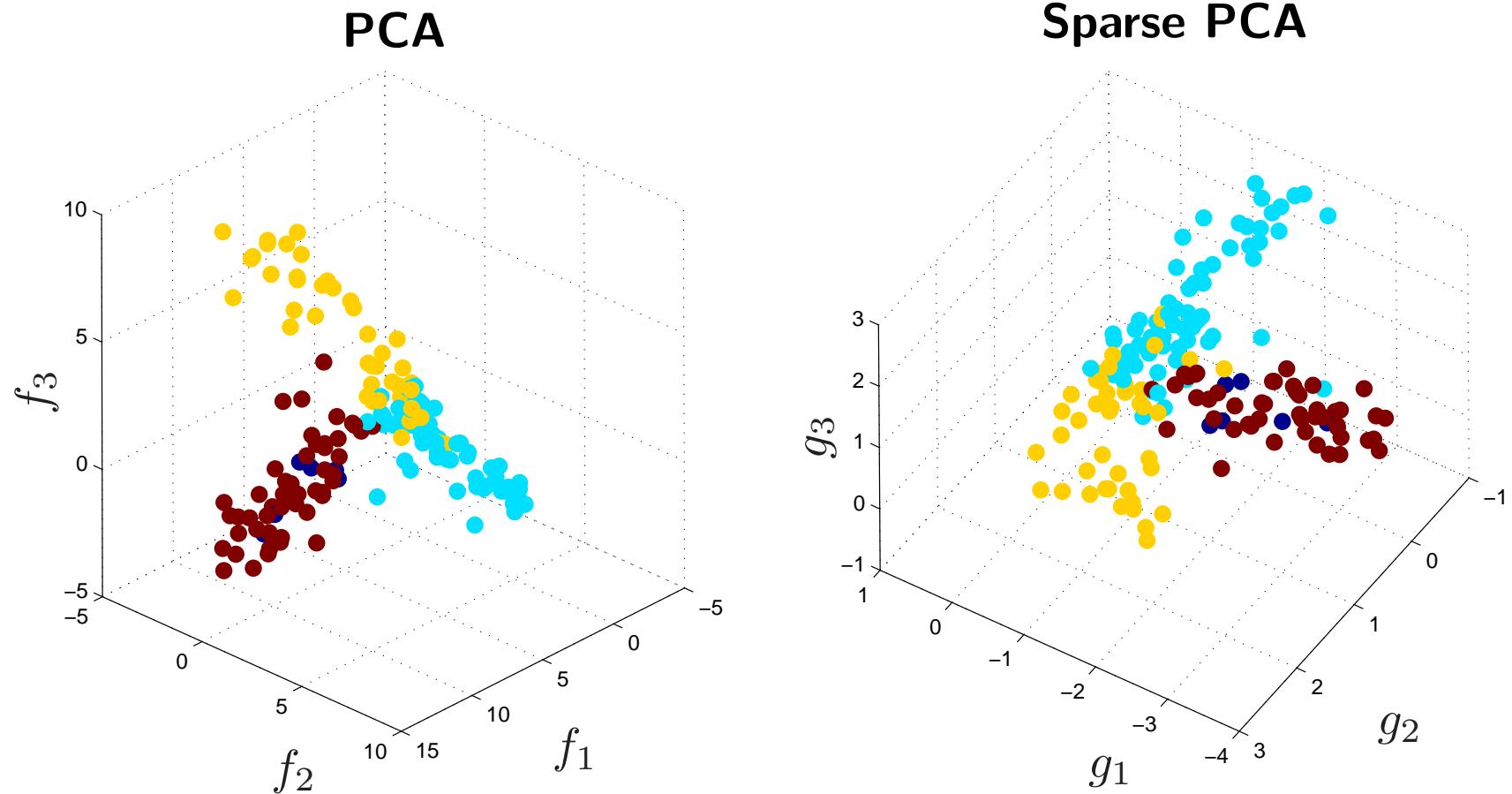
- Classical dimensionality reduction tool.
- Numerically cheap:  $O(kn^2)$ , only requires computing k leading eigenvectors.

## Sparse PCA

- Seeks factors with a few nonzero coefficients.
- **Sparse** factors capture maximum variance and improve **interpretability**.
- Numerically hard: sparsity makes it a combinatorial problem.

# Introduction

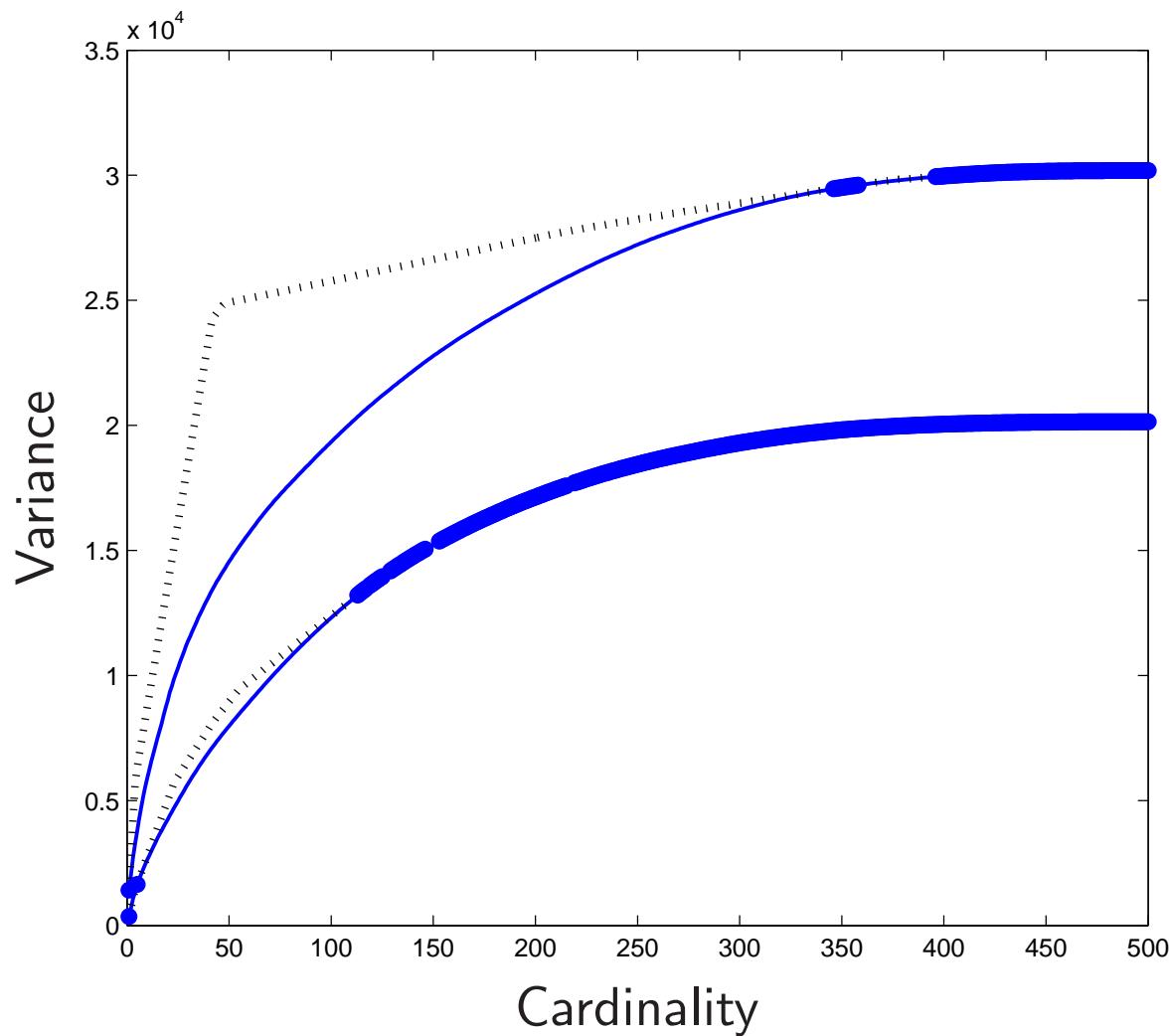
Clustering of gene expression data in PCA versus sparse PCA, on 500 genes.



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.

# Introduction



Variance (solid lines) versus cardinality tradeoff curve for two gene expression data sets, lymphoma (top) and colon cancer (bottom).

# Introduction

Given a (centered) data set  $A \in \mathbf{R}^{n \times m}$  composed of  $m$  observations on  $n$  variables, we form the covariance matrix  $C = AA^T/(m - 1)$ .

**Principal Component Analysis.** To get the first factor, we solve:

$$\begin{aligned} & \text{maximize} && x^T C x \\ & \text{subject to} && \|x\| = 1, \end{aligned}$$

in the variable  $x \in \mathbf{R}^n$ , i.e. we maximize the **variance** explained by the **factor**  $x$ .

**Sparse Principal Component Analysis.** We constrain the cardinality of the factor  $x$  and solve instead:

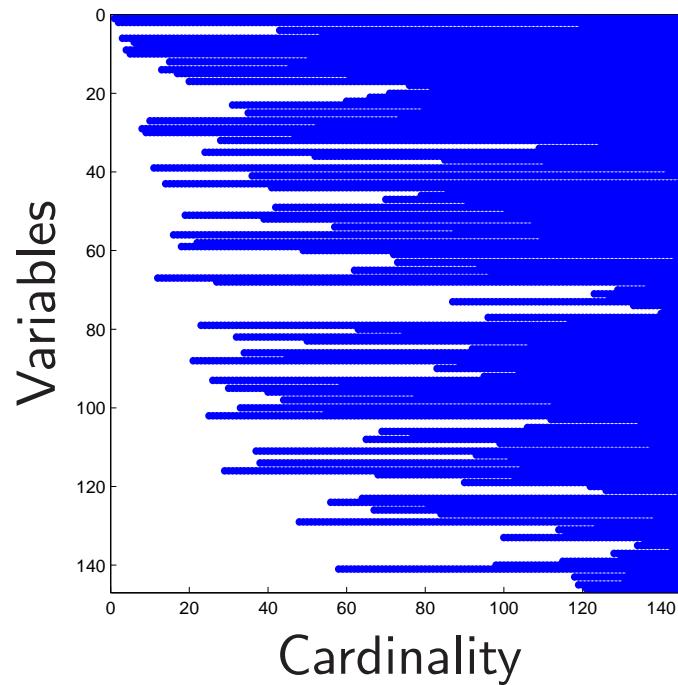
$$\begin{aligned} & \text{maximize} && x^T C x \\ & \text{subject to} && \mathbf{Card}(x) = k \\ & && \|x\| = 1, \end{aligned}$$

in the variable  $x \in \mathbf{R}^n$ , where **Card**( $x$ ) is the number of nonzero coefficients in the vector  $x$  and  $k > 0$  is a parameter controlling **sparsity**.

# Sorting

Simplest solution: just sort variables according to variance.

**Schur-Horn theorem:** the diagonal of a matrix majorizes its eigenvalues so the diagonal of a matrix is a diffused vector of eigenvalues.



In this example, we selected variables according to their variance, but we ordered them according to their true ranking (computed from the optimal solution).

## Related Work

- Cadima & Jolliffe (1995): the loadings with small absolute value are thresholded to zero.
- Johnstone & Lu (2004) apply this to ECG data and show consistency.
- Zou, Hastie & Tibshirani (2006), non-convex algo. (SPCA) based on a  $l_1$  penalized representation of PCA as a regression problem.
- Non-convex methods (SCoTLASS) by Jolliffe, Trendafilov & Uddin (2003).
- A greedy search algorithm by Moghaddam, Weiss & Avidan (2006).

This talk is mostly about the results in d'Aspremont, El Ghaoui, Jordan & Lanckriet (2007). New results in:

- d'Aspremont, Bach & El Ghaoui (2007) compute a full approximate regularization path in  $O(n^3)$ .
- Sriperumbudur, Torres & Lanckriet (2007) apply D.C. algorithms to the penalized eigenvalue problem.

# Outline

- Introduction
- Sparse PCA
  - **Semidefinite Relaxation**
  - Smooth Optimization
- Sparse Eigenvalues
  - Variable Selection
  - Compressed Sensing
- Numerical Experiments

## Related Work

- Non-convex methods produce approximate solution with unpredictable complexity.
- Here, we produce approximate solutions with **predictable complexity**, together with bounds on suboptimality.

Combine two classic relaxation techniques:

- The lifting procedure à la MAXCUT by Goemans & Williamson (1995).
- A  $\ell_1$  norm relaxation of the cardinality constraint. Used in basis pursuit by Chen, Donoho & Saunders (2001), LASSO by Tibshirani (1996), etc.

# Semidefinite relaxation

Start from:

$$\begin{aligned} & \text{maximize} && x^T A x \\ & \text{subject to} && \|x\|_2 = 1 \\ & && \mathbf{Card}(x) \leq k, \end{aligned}$$

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

$$\begin{aligned} & \text{maximize} && \mathbf{Tr}(AX) \\ & \text{subject to} && \mathbf{Tr}(X) = 1 \\ & && \mathbf{Card}(X) \leq k^2 \\ & && X = xx^T, \end{aligned}$$

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

$$\begin{aligned} & \text{maximize} && \mathbf{Tr}(AX) \\ & \text{subject to} && \mathbf{Tr}(X) = 1 \\ & && \mathbf{Card}(X) \leq k^2 \\ & && X \succeq 0, \quad \mathbf{Rank}(X) = 1, \end{aligned}$$

again, this is the **same problem**.

# Semidefinite relaxation

We have made **some progress**:

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

But this is still a hard problem:

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

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

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

# Semidefinite Programming

Semidefinite relaxation:

$$\begin{aligned} & \text{maximize} && x^T Ax \\ & \text{subject to} && \|x\|_2 = 1 \\ & && \text{Card}(x) \leq k, \end{aligned}$$

**becomes**

$$\begin{aligned} & \text{maximize} && \text{Tr}(AX) \\ & \text{subject to} && \text{Tr}(X) = 1 \\ & && \mathbf{1}^T |X| \mathbf{1} \leq k \\ & && X \succeq 0, \end{aligned}$$

- 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. . .

# Robustness & Tightness

**Robustness.** The penalized problem can be written:

$$\min_{\{ |U_{ij}| \leq \rho \}} \lambda^{\max}(A + U)$$

Natural interpretation: **robust** maximum eigenvalue problem with componentwise noise of magnitude  $\rho$  on the coefficients of the matrix  $A$ .

**Tightness.** The KKT optimality conditions are here:

$$\begin{cases} (A + U)X = \lambda^{\max}(A + U)X \\ U \circ X = \rho|X| \\ \text{Tr}(X) = 1, \quad X \succeq 0 \\ |U_{ij}| \leq \rho, \quad i, j = 1, \dots, n. \end{cases}$$

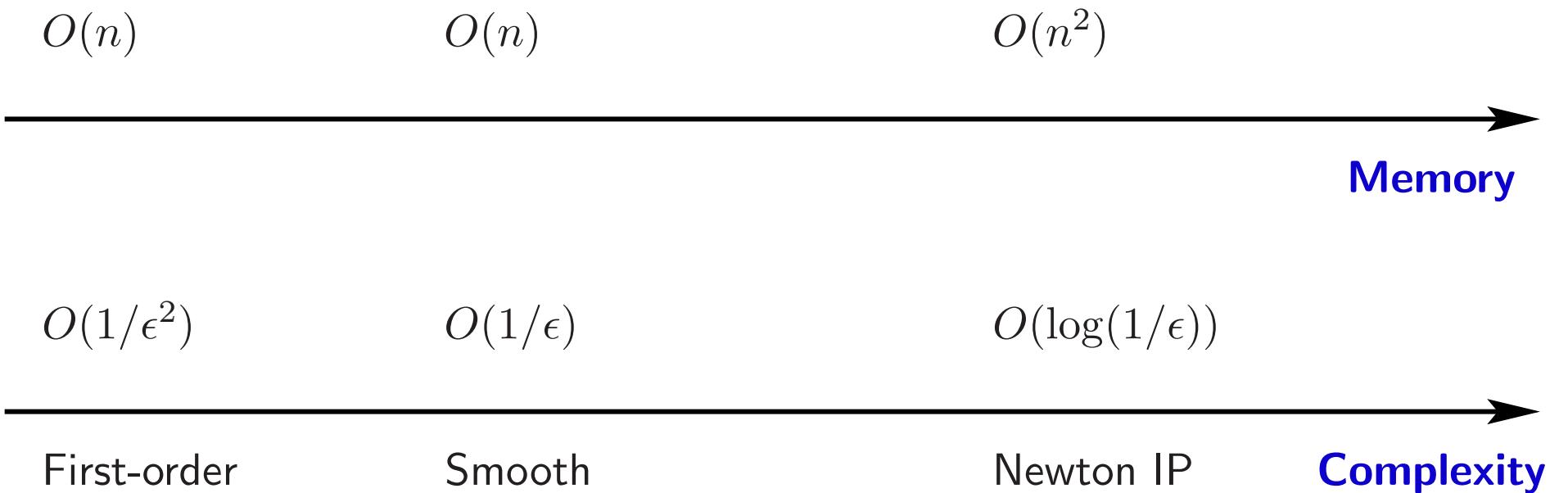
If the eigenvalue  $\lambda^{\max}(A + U)$  is simple,  $\text{Rank}(X) = 1$  and the semidefinite relaxation is **tight**.

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# First order algorithm

Complexity options. . .



# First order algorithm

Here, we can exploit problem structure

- Our problem here has a particular **min-max** structure:

$$\min_{|U_{ij}| \leq \rho} \max_{X \in \mathbf{S}^n} \text{Tr}((A + U)X)$$

- This min-max structure means that we can use prox function algorithms by Nesterov (2005) (see also Nemirovski (2004)) to solve large, dense problem instances.

# First order algorithm

If problem has min-max model, **two steps**:

- **Regularization.** Add strongly convex penalty inside the min-max representation to produce an  $\epsilon$ -approximation of  $f$  with Lipschitz continuous gradient (generalized Moreau-Yosida regularization step, see Lemaréchal & Sagastizábal (1997) for example).
- **Optimal first order minimization.** Use optimal first order scheme for Lipschitz continuous functions detailed in Nesterov (1983) to the solve the regularized problem.

**Benefits:**

- Produces an  $\epsilon$  solution is given by  $O(1/\epsilon)$  compared to  $O(1/\epsilon^2)$  for generic first-order methods.
- Low memory requirements. Change in **granularity** of the solver: larger number of cheaper iterations.

**Caveat:** Two (projection) subproblems need to be solved very efficiently. . .

# First order algorithm

**Regularization.** Let  $\mu > 0$  and  $X \in \mathbf{S}_n$ , we define:

$$f_\mu(X) = \mu \log \text{Tr} \left( \exp \left( \frac{X}{\mu} \right) \right)$$

We then have:

$$\lambda^{\max}(X) \leq f_\mu(X) \leq \lambda^{\max}(X) + \mu \log n,$$

so if we set  $\mu = \epsilon / \log n$ , this becomes a **uniform  $\epsilon$ -approximation** of  $\lambda^{\max}(X)$  with a **Lipschitz continuous gradient** with constant:

$$L = \frac{1}{\mu} = \frac{\log n}{\epsilon}.$$

The gradient  $\nabla f_\mu(X)$  can be computed explicitly in  $O(n^3)$  as:

$$\exp \left( \frac{X - \lambda^{\max}(X)\mathbf{I}}{\mu} \right) / \text{Tr} \left( \exp \left( \frac{X - \lambda^{\max}(X)\mathbf{I}}{\mu} \right) \right)$$

using the same matrix exponential.

# First order algorithm

**Optimal first-order minimization.** The minimization algorithm in Nesterov (1983) then involves the following steps:

Choose  $\epsilon > 0$  and set  $X_0 = \beta I_n$ , **For**  $k = 0, \dots, N$  **do**

1. Compute  $\nabla f_\epsilon(X_k)$
2. Find  $Y_k = \arg \min_{Y \in \mathcal{Q}} \{\text{Tr}(\nabla f_\epsilon(X_k)(Y - X_k)) + \frac{1}{2}L_\epsilon\|Y - X_k\|_F^2\}$ .
3. Find  $Z_k = \arg \min_{X \in \mathcal{Q}} \left\{ L_\epsilon \beta^2 d_1(X) + \sum_{i=0}^k \frac{i+1}{2} \text{Tr}(\nabla f_\epsilon(X_i)(X - X_i)) \right\}$ .
4. Update  $X_k = \frac{2}{k+3}Z_k + \frac{k+1}{k+3}Y_k$ .
5. Test if gap less than target precision.

- **Step 1** requires computing a matrix exponential.
- **Steps 2 and 3** are both Euclidean projections on  $\mathcal{Q} = \{U \in \mathbf{S}^n : |U_{ij}| \leq \rho\}$ .

# First order algorithm

## Complexity:

- The number of iterations to get accuracy  $\epsilon$  is

$$O\left(\frac{n\sqrt{\log n}}{\epsilon}\right)$$

- At each iteration, the cost of computing a matrix exponential up to machine precision is  $O(n^3)$ .

## Computing matrix exponentials:

- Many options, cf. “Nineteen Dubious Ways to Compute the Exponential of a Matrix” by Moler & Van Loan (1978), Moler & Van Loan (2003).
- Padé approximation, full eigenvalue decomposition:  $O(n^3)$  up to machine precision.
- In practice, machine precision is unnecessary and a partial eigenvalue decomposition is enough (see d’Aspremont (2005)).

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# Sparse Eigenvalues

Combining semidefinite and  $\ell_1$  relaxations, we obtained:

$$\begin{aligned}\lambda_{\max}^k(A) \leq & \max. \quad \text{Tr}(AX) \\ \text{s.t.} \quad & \text{Tr}(X) = 1 \\ & \mathbf{1}^T |X| \mathbf{1} \leq k \\ & X \succeq 0,\end{aligned}$$

This relaxation produces **upper bounds** on sparse (or restricted) maximum eigenvalues. Similarly, we can get lower bounds on sparse minimum eigenvalues.

- Used to bound MSE and model consistency in LASSO (sparse least-squares).
- Control recovery rates in compressed sensing.

# LASSO

Assume that observations  $(Y_1, \dots, Y_n)$  follow a linear model:

$$Y = X\beta + \epsilon$$

where  $\beta \in \mathbf{R}^p$  and  $\epsilon_i \sim \mathcal{N}(0, \sigma^2)$ . We define the LASSO estimator of  $\beta$  as:

$$\hat{\beta} = \operatorname{argmin}_{\beta} \|Y - X\beta\|_2^2 + \lambda \|\beta\|_1.$$

## Consistency.

- Suppose  $b$  is **sparse** with cardinality  $s(n)$ , Meinshausen & Yu (2007) show that with probability tending to 1 as  $n \rightarrow \infty$ :

$$\|\beta - \hat{\beta}\|_2^2 \leq M \frac{s(n) \log p(n)}{n \lambda_{\min}^{s(n)}(X^T X)}$$

- Meinshausen & Yu (2007) also show sign consistency based on sparse eigenvalues. Similar non-asymptotic result by Candès & Tao (2007).

# Compressed Sensing

Following Candès & Tao (2005) (see also Donoho & Tanner (2005)), recover a signal  $f \in \mathbf{R}^n$  from corrupted measurements  $y$ :

$$y = Af + e,$$

where  $A \in \mathbf{R}^{m \times n}$  is a coding matrix and  $e \in \mathbf{R}^m$  is an unknown **sparse** vector of errors.

This amounts to solving the following (combinatorial) problem:

$$\begin{aligned} &\text{minimize} && \mathbf{Card}(x) \\ &\text{subject to} && Fx = Fy \end{aligned}$$

where  $F \in \mathbf{R}^{p \times m}$  is a matrix such that  $FA = 0$ .

# Compressed Sensing: Restricted Isometry Constant

Given a matrix  $F \in \mathbb{R}^{p \times m}$  and  $0 < S \leq m$ , its **restricted isometry** constant  $\delta_S$  is the smallest number such that for any subset  $I \subset [1, m]$  of cardinality at most  $S$  we have:

$$(1 - \delta_S) \|c\|^2 \leq \|F_I c\|^2 \leq (1 + \delta_S) \|c\|^2,$$

for all  $c \in \mathbb{R}^{|I|}$ , where  $F_I$  is the submatrix of  $F$  formed by keeping only the columns of  $F$  in the set  $I$ .

# Compressed sensing: perfect recovery

The following result then holds.

**Proposition 1.** *Candès & Tao (2005).* Suppose that the restricted isometry constants of a matrix  $F \in \mathbb{R}^{p \times m}$  satisfy :

$$\delta_S + \delta_{2S} + \delta_{3S} < 1 \quad (1)$$

for some integer  $S$  such that  $0 < S \leq m$ , then if  $x$  is an optimal solution of the convex program:

$$\begin{aligned} &\text{minimize} && \|x\|_1 \\ &\text{subject to} && Fx = Fy \end{aligned}$$

such that  $\text{Card}(x) \leq S$  then  $x$  is also an optimal solution of the combinatorial problem:

$$\begin{aligned} &\text{minimize} && \text{Card}(x) \\ &\text{subject to} && Fx = Fy. \end{aligned}$$

## Compressed sensing: restricted isometry

The restricted isometry constant  $\delta_S$  in condition can be computed by solving the following sparse PCA problem:

$$\begin{aligned} (1 + \delta_S) = & \max. && x^T(F^T F)x \\ & \text{s. t.} && \mathbf{Card}(x) \leq S \\ & && \|x\| = 1, \end{aligned}$$

in the variable  $x \in \mathbf{R}^m$  (a similar sparse PCA problem gives the other inequality).

- Candès & Tao (2005) obtain an **asymptotic** proof that some random matrices satisfy the restricted isometry condition with **overwhelming probability** (i.e. exponentially small probability of failure)
- Upper bounds for sparse PCA prove **deterministically** and with **polynomial complexity** that a finite dimensional matrix satisfies the restricted isometry condition.

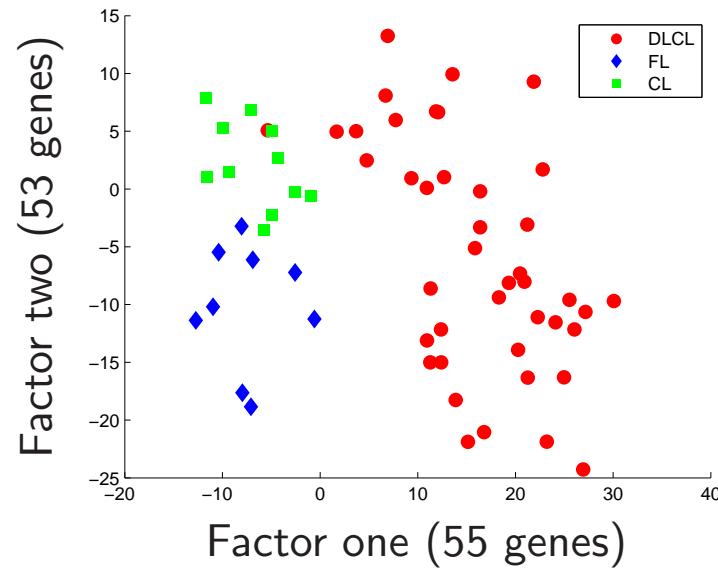
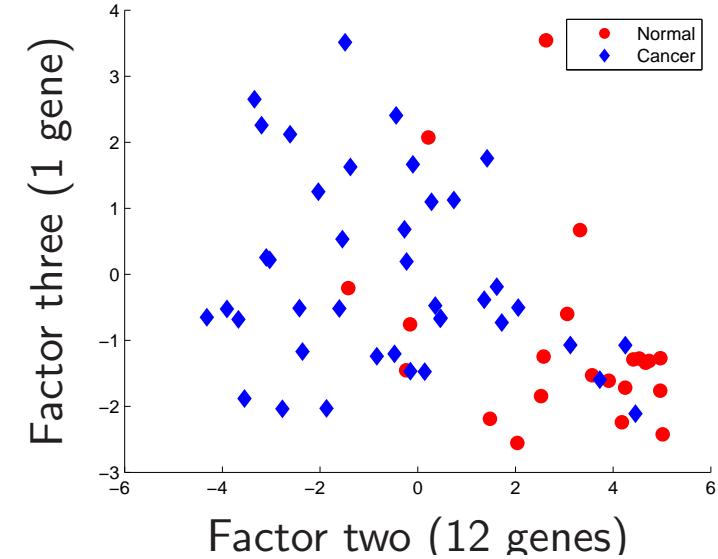
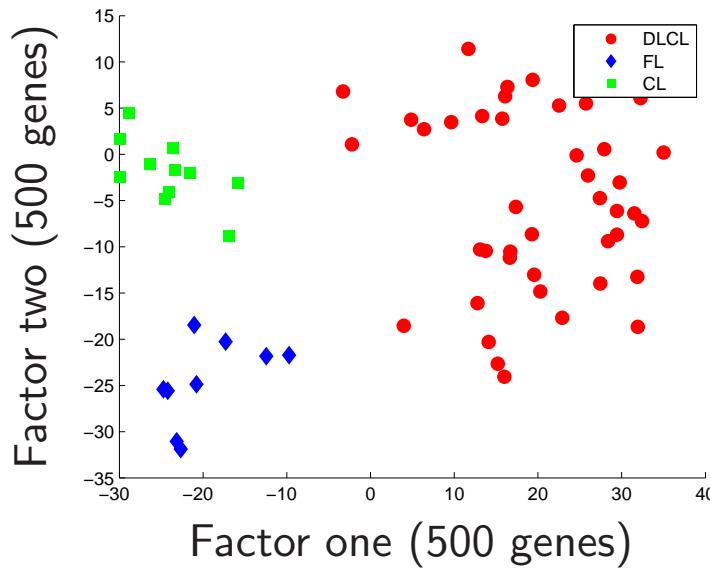
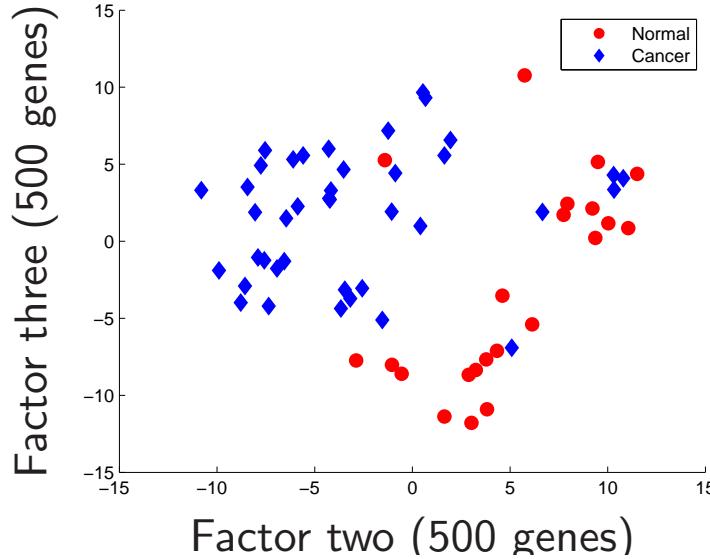
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# Gene Expression Data

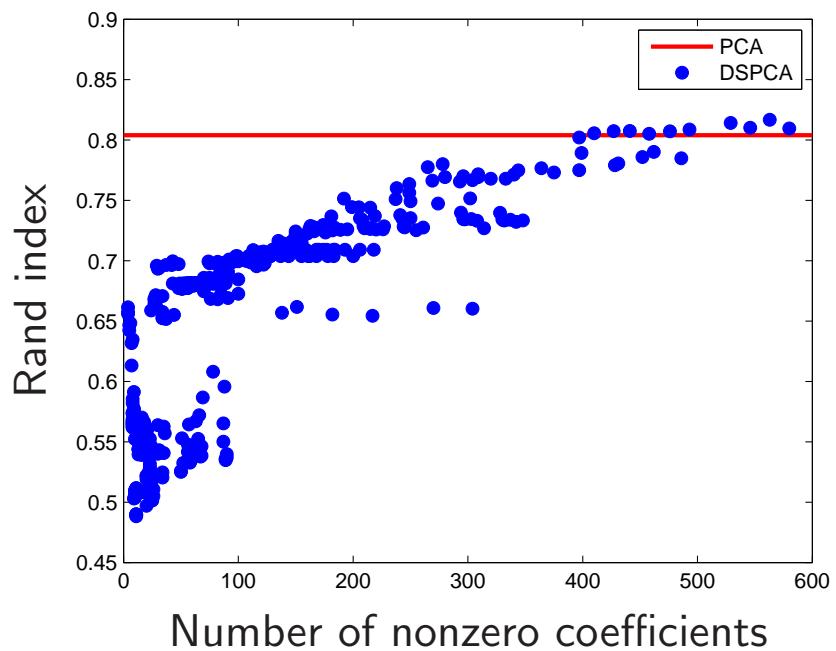
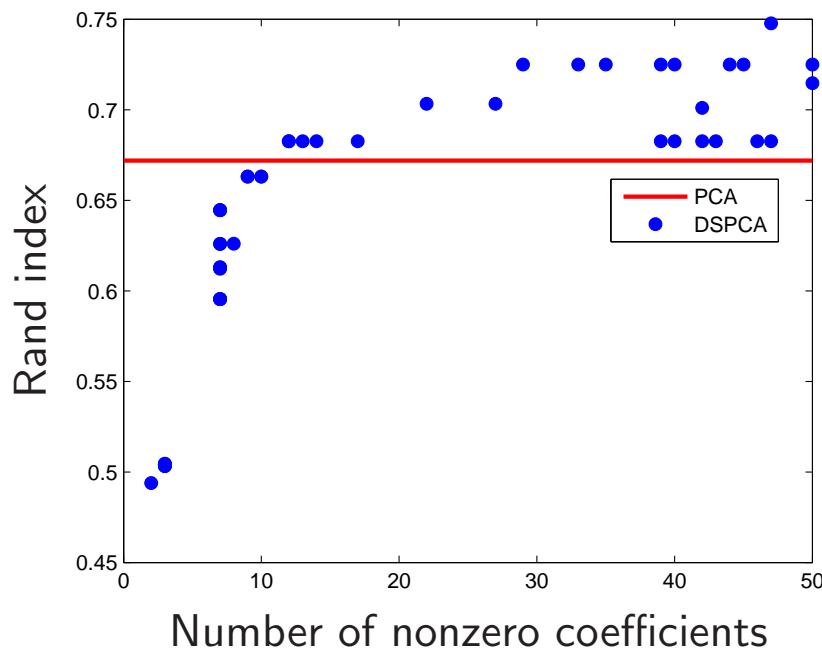
- Use sparse PCA as a crude clustering/variable selection tool (see Luss & d'Aspremont (2007)).
- Use colon cancer data set of Alon, Barkai, Notterman, Gish, Ybarra, Mack & Levine (1999), lymphoma data from Alizadeh, Eisen, Davis, Ma, Lossos & Rosenwald (2000).
- Track clustering quality versus number of genes used.

# Sparse PCA: clustering



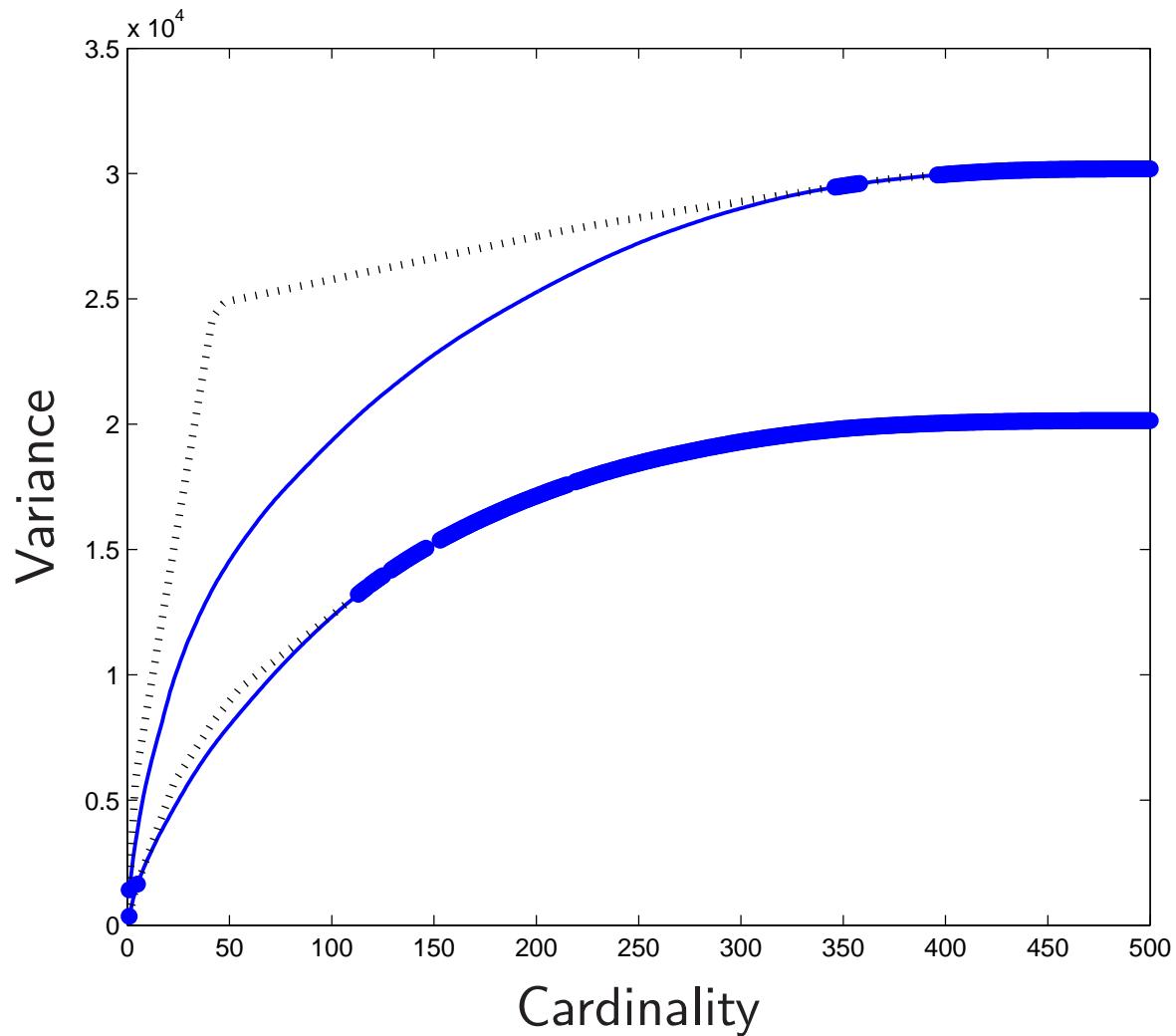
PCA (left) and DSPCA (right), colon cancer (top) and lymphoma (bottom).

# Sparse PCA: clustering



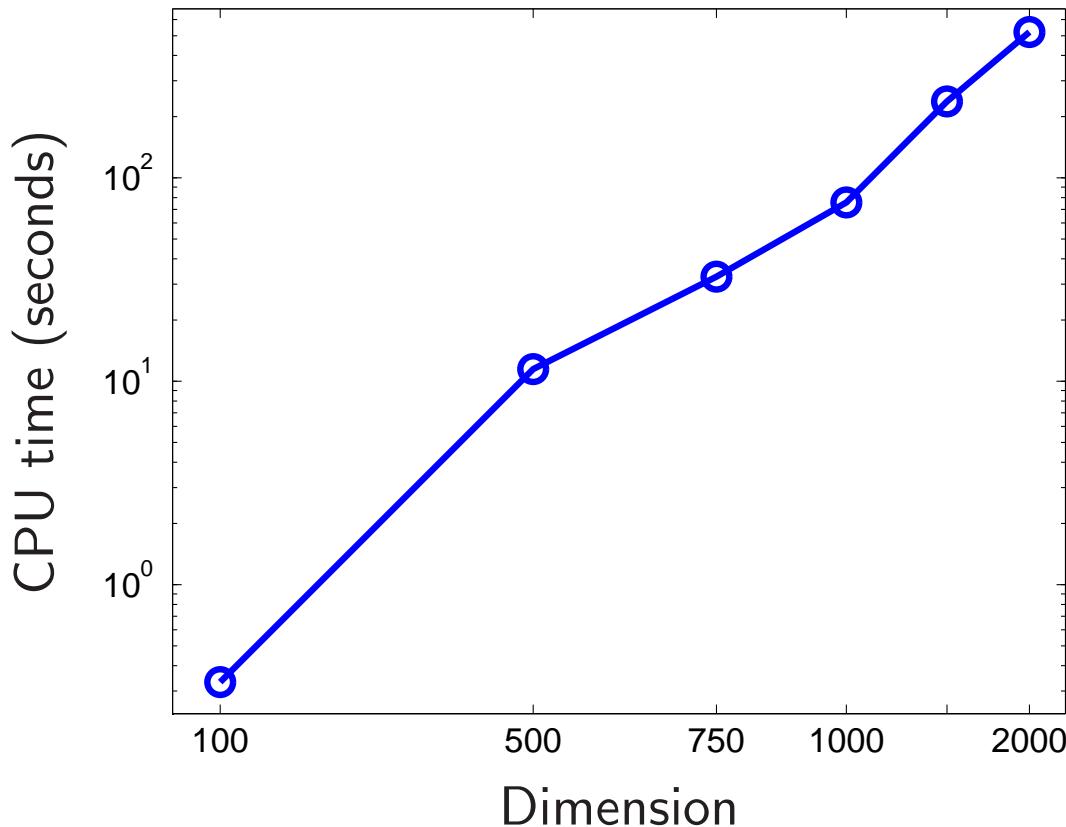
Rand index (clustering) versus sparsity: colon cancer (left) & lymphoma (right).

# Tradeoff



Variance (solid lines) versus cardinality tradeoff curve for two gene expression data sets, lymphoma (top) and colon cancer (bottom).

# CPU time



$n$	CPU time (secs)
100	0 m 1 s
500	0 m 11 s
750	1 m 33 s
1000	1 m 16 s
1500	4 m 57 s
2000	9 m 41 s

Using the data in Alon et al. (1999), with  $\rho = 1$ , we plot CPU time to get a  $10^2$  decrease in duality gap.

# Conclusion

- The tradeoff between sparsity and explained variance is often favorable.
- Dense semidefinite programs solved efficiently for matrices with  $n \sim 10^3$
- Slides online.
- Source code, binaries and test data available at:

[www.princeton.edu/~aspremon/DSPCA.htm](http://www.princeton.edu/~aspremon/DSPCA.htm)

- More results in d'Aspremont, Bach & El Ghaoui (2007) and Sriperumbudur et al. (2007).

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