# Lecture 4. Random Projections and Johnson-Lindenstrauss Lemma

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#### **Outline**

#### Recall: PCA and MDS

#### Random Projections

Example: Human Genomics Diversity Project

Johnson-Lindenstrauss Lemma

Proofs

#### Applications of Random Projections

Locality Sensitive Hashing

Compressed Sensing

Algorithms: BP, OMP, LASSO, Dantzig Selector, ISS, LBI etc.

From Johnson-Lindenstrauss Lemma to RII

Appendix: A Simple Version of Johnson-Lindenstrauss Lemma

#### PCA and MDS

- ▶ Data matrix:  $X = [x_1, ..., x_n] \in \mathbb{R}^{p \times n}$ 
  - Centering: Y = XH, where  $H = I \frac{1}{n}\mathbf{1}\mathbf{1}^T$
- ▶ Singular Value Decomposition  $Y = USV^T$ ,  $S = \mathbf{diag}(\sigma_j)$ ,  $\sigma_1 \geq \sigma_2 \geq \ldots \geq \sigma_{\min(n,p)}$ 
  - PCA is given by top-k SVD  $(S_k, U_k)$ :  $U_k = (u_1, \dots, u_k) \in \mathbb{R}^{p \times k}$ , with embedding coordinates  $U_k S_k$
  - MDS is given by top-k SVD  $(S_k,V_k)$ :  $V_k=(v_1,\ldots,v_k)\in\mathbb{R}^{n\times k}$ , with embedding coordinates  $V_kS_k$
  - Kernel PCA (MDS): for  $K\succeq 0$ ,  $K_c=HKH^T$ ,  $K_c=U\Lambda U^T$  gives MDS embedding  $U_k\Lambda_k^{1/2}\in\mathbb{R}^{n\times k}$

# Computational Concerns: Big Data and High Dimensionality

- ▶ Big Data: n is large
  - Downsample for approximate PCA:

$$\widehat{\Sigma}_{n'} = \frac{1}{n'} \sum_{i=1}^{n'} (x_i - \widehat{\mu}_{n'}) (x_i - \widehat{\mu}_{n'})^T, \qquad \widehat{\Sigma}_{n'} = U \Lambda U^T$$

- Nyström Approximation for MDS:  $V_k = (v_1, \dots, v_k) \in \mathbb{R}^{n \times k}$  (we'll come to this in Manifold Learning ISOMAP)
- ▶ High Dimensionality: *p* is large
  - Random Projections for PCA:  $RXH = \tilde{U}\tilde{S}\tilde{V}^T$  with random matrix  $R^{d\times p}$  (today):  $\tilde{U}_k = (\tilde{u}_1,\ldots,\tilde{u}_k) \in \mathbb{R}^{d\times k}$
  - Perturbation of MDS:  $\tilde{V}_k = (\tilde{v}_1, \dots, \tilde{v}_k) \in \mathbb{R}^{n \times k}$

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# **Random Projections: Examples**

$$\begin{array}{l} \blacktriangleright \ R=[r_1,\cdots,r_k], \ r_i\sim U(S^{d-1}), \ \text{e.g.} \ r_i=(a_1^i,\cdots,a_d^i)/\parallel a^i\parallel \\ a_k^i\sim N(0,1) \end{array}$$

$$R = A/\sqrt{k} \quad A_{ij} \sim N(0,1)$$

$$R = A/\sqrt{k} \quad A_{ij} = \begin{cases} 1 & p = 1/2 \\ -1 & p = 1/2 \end{cases}$$

▶ 
$$R = A/\sqrt{k/s}$$
  $A_{ij} = \begin{cases} 1 & p = 1/(2s) \\ 0 & p = 1 - 1/s \\ -1 & p = 1/(2s) \end{cases}$  where  $s = 1, 2, \sqrt{D}, D/\log D$ , etc.

# **Example: Human Genomics Diversity Project**

► Now consider a SNPs (Single Nucleid Polymorphisms) dataset in Human Genome Diversity Project (HGDP),

http://www.cephb.fr/en/hgdp\_panel.php

- Data matrix of n-by-p for n=1,064 individuals around the world and p=644,258 SNPs.
- Each entry in the matrix has 0, 1, 2, and 9, representing "AA", "AC", "CC", and "missing value", respectively.
- After removing 21 rows with all missing values, we are left with a matrix X of size  $1,043\times 644,258$ .

# Original MDS (PCA)

- ▶ Projection of 1,043 persons on the top-2 MDS (PCA) coordinates.
  - Define

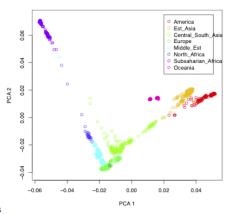
$$K = HXX^TH = U\Lambda U^T, \qquad H = I - \frac{1}{n}\mathbf{1}\mathbf{1}^T$$

which is a positive semi-define matrix as centered Gram matrix whose eigenvalue decomposition is given by  $U\Lambda U^T$ .

– Take the first two eigenvectors  $\sqrt{\lambda_i}u_i$   $(i=1,\ldots,2)$  as the projections of n individuals.

# Figure: Original MDS (PCA)

Projection of 1,043 individuals on the top-2 MDS principal components, shows a continuous trajectory of human migration in history: human origins from Africa, then migrates to the Middle East, followed by one branch to Europe and another branch to Asia, finally spreading into America and Oceania.



# Random Projection MDS (PCA)

▶ To reduce the computational cost due to the high dimensionality p=644,258, we randomly select (without replacement)  $\{n_i,i=1,\ldots,k\}$  from  $1,\ldots,p$  with equal probability. Let  $R\in\mathbb{R}^{k\times p}$  is a Bernoulli random matrix satisfying:

$$R_{ij} = \begin{cases} 1/k & j = n_i, \\ 0 & otherwise. \end{cases}$$

Now define

$$\widetilde{K} = H(XR^T)(RX^T)H$$

whose eigenvectors leads to new principal components of MDS.

# Figure: Comparisons of Random Projected MDS with Original One

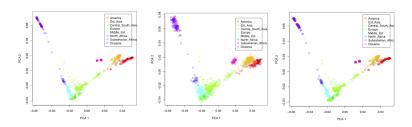


Figure: (Left) Projection of 1043 individuals on the top 2 MDS principal components. (Middle) MDS computed from 5,000 random columns. (Right) MDS computed from 100,000 random columns. Pictures are due to Qing Wang.

### Question

# How does the Random Projection work?

#### **General MDS**

▶ Given pairwise distances  $d_{ij}$  between n sample points, MDS aims to find  $Y := [y_i]_{i=1}^n \in \mathbb{R}^{k \times n}$  such that the following sum of square is minimized,

$$\min_{Y = [y_1, \dots, y_n]} \quad \sum_{i,j} (\|y_i - y_j\|^2 - d_{ij}^2)^2$$
subject to 
$$\sum_{i=1}^n y_i = 0$$
(1)

i.e. the total distortion of distances is minimized.

#### Metric MDS

▶ When  $d_{ij} = \|x_i - x_j\|$  is exactly given by the distances of points in Euclidean space  $x_i \in \mathbb{R}^p$ , classical (metric) MDS defines a positive semidefinite kernel matrix  $K = -\frac{1}{2}HDH$  where  $D = (d_{ij}^2)$  and  $H = I - \frac{1}{n}\mathbf{1}\mathbf{1}^T$ . Then, the minimization (1) is equivalent to

$$\min_{Y \in \mathbb{R}^{k \times n}} \quad \|Y^T Y - K\|_F^2 \tag{2}$$

i.e. the total distortion of distances is minimized by setting the column vectors of Y as the eigenvectors corresponding to k largest eigenvalues of K.

#### MDS toward Minimal Total Distortion

- ▶ The main features of MDS are the following.
  - MDS looks for Euclidean embedding of data whose total or average metric distortion are minimized.
  - MDS embedding basis is adaptive to the data, e.g. as a function of data via spectral decomposition.
- Can we have a tighter control on metric distortions, e.g. uniform distortion control?

# **Uniformly Almost-Isometry?**

▶ What if a *uniform* control on metric distortion: there exists a  $\epsilon \in (0,1)$ , such that for every (i,j) pair,

$$(1 - \epsilon) \le \frac{\|y_i - y_j\|^2}{d_{ij}^2} \le (1 + \epsilon)?$$

It is a uniformly almost isometric embedding or a Lipschitz mapping from metric space  $\mathcal X$  to  $\mathcal Y.$ 

▶ An beautiful answer is given by Johnson-Lindenstrauss Lemma, if  $\mathcal X$  is an Euclidean space (or more generally Hilbert space), that  $\mathcal Y$  can be a subspace of dimension  $k = O(\log n/\epsilon^2)$  via random projections to obtain an almost isometry with high probability.

#### Johnson-Lindenstrauss Lemma

# Theorem (Johnson-Lindenstrauss Lemma)

For any  $0 < \epsilon < 1$  and any integer n, let k be a positive integer such that

$$k \ge (4+2\alpha)(\epsilon^2/2 - \epsilon^3/3)^{-1} \ln n, \quad \alpha > 0.$$

Then for any set V of n points in  $\mathbb{R}^p$ , there is a map  $f:\mathbb{R}^p\to\mathbb{R}^k$  such that for all  $u,v\in V$ 

$$(1 - \epsilon) \parallel u - v \parallel^{2} \le \parallel f(u) - f(v) \parallel^{2} \le (1 + \epsilon) \parallel u - v \parallel^{2}$$
 (3)

Such a f in fact can be found in randomized polynomial time. In fact, inequalities (3) holds with probability at least  $1-1/n^{\alpha}$ .

#### Remark

- Almost isometry is achieved with a uniform metric distortion bound (Bi-Lipschitz bound), with high probability, rather than average metric distortion control;
- ▶ The mapping is **universal**, rather than being adaptive to the data.
- ▶ The theoretical basis of this method was given as a lemma by Johnson and Lindenstrauss (1984) in the study of a Lipschitz extension problem in Banach space.
- ▶ In 2001, Sanjoy Dasgupta and Anupam Gupta, gave a simple proof of this theorem using elementary probabilistic techniques in a four-page paper. Below we are going to present a brief proof of Johnson-Lindenstrauss Lemma based on the work of Sanjoy Dasgupta, Anupam Gupta, and Dimitris Achlioptas.

#### Note

▶ The distributions of the following two events are identical:

 $\mbox{unit vector was randomly projected to $k$-subspace} \iff \mbox{random vector on } S^{d-1} \mbox{ fixed top-$k$ coordinates.}$ 

Based on this observation, we change our target from random k-dimensional projection to random vector on sphere  $S^{d-1}$ .

- Let  $x_i \sim N(0,1)$   $(i=1,\cdots,p)$ , and  $X=(x_1,\cdots,x_p)$ , then  $Y=X/\|x\| \in S^{p-1}$  is uniformly distributed.
- Fixing top-k coordinates, we get  $z=(x_1,\cdots,x_k,0,\cdots,0)^T/\|x\|\in\mathbb{R}^p$ . Let  $L=\|z\|^2$  and  $\mu:=k/p$ . Note that  $\mathbf{E}\,\|(x_1,\cdots,x_k,0,\cdots,0)\|^2=k=\mu\cdot\mathbf{E}\,\|x\|^2$ .
- The following lemma shows that L is concentrated around  $\mu$ .

# **Key Lemma**

#### Lemma

For any k < p, there hold

(a) if  $\beta < 1$  then

$$\mathbf{Prob}[L \le \beta \mu] \le \beta^{k/2} \left( 1 + \frac{(1-\beta)k}{p-k} \right)^{(p-k)/2} \le \exp\left(\frac{k}{2} (1-\beta + \ln \beta)\right)$$

(b) if  $\beta > 1$  then

Here  $\mu = k/p$ .

$$\mathbf{Prob}[L \ge \beta \mu] \le \beta^{k/2} \left( 1 + \frac{(1-\beta)k}{p-k} \right)^{(p-k)/2} \le \exp\left(\frac{k}{2} (1-\beta + \ln \beta)\right)$$

#### **Proof of Johnstone-Lindenstrauss Lemma**

- ▶ If  $p \le k$ , the theorem is trivial.
- ▶ Otherwise take a random k-dimensional subspace S, and let  $v_i'$  be the projection of point  $v_i \in V$  into S, then setting  $L = \|v_i' v_j'\|^2$  and  $\mu = (k/p)\|v_i v_j\|^2$  and applying Lemma 1(a), we get that

$$\begin{aligned} \mathbf{Prob}[L &\leq (1-\epsilon)\mu] \leq \exp\left(\frac{k}{2}(1-(1-\epsilon)+\ln(1-\epsilon))\right) \\ &\leq \exp\left(\frac{k}{2}(\epsilon-(\epsilon+\frac{\epsilon^2}{2}))\right), \\ &\qquad \qquad \text{by } \ln(1-x) \leq -x-x^2/2 \text{ for } 0 \leq x < 1 \\ &= \exp\left(-\frac{k\epsilon^2}{4}\right) \leq \exp(-(2+\alpha)\ln n), \\ &\qquad \qquad \text{for } k \geq 4(1+\alpha/2)(\epsilon^2/2)^{-1}\ln n \\ &= \frac{1}{n^{2+\alpha}} \end{aligned}$$

# **Proof of Johnstone-Lindenstrauss Lemma (continued)**

▶ Similarly, we can apply Lemma 1(b) to get

$$\begin{aligned} \mathbf{Prob}[L \geq (1+\epsilon)\mu] &\leq \exp\left(\frac{k}{2}(1-(1+\epsilon)+\ln(1+\epsilon))\right) \\ &\leq \exp\left(\frac{k}{2}(-\epsilon+(\epsilon-\frac{\epsilon^2}{2}+\frac{\epsilon^3}{3}))\right), \\ & \text{by } \ln(1+x) \leq x-x^2/2+x^3/3 \text{ for } x \geq 0 \\ &= \exp\left(-\frac{k}{2}(\epsilon^2/2-\epsilon^3/3)\right) \leq \exp(-(2+\alpha)\ln n), \\ & \text{for } k \geq 4(1+\alpha/2)(\epsilon^2/2-\epsilon^3/3)^{-1}\ln n \\ &= \frac{1}{n^{2+\alpha}} \end{aligned}$$

# **Proof of Johnstone-Lindenstrauss Lemma (continued)**

Now set the map  $f(x) = \sqrt{\frac{d}{k}}x' = \sqrt{\frac{d}{k}}(x_1,\ldots,x_k,0,\ldots,0)$ . By the above calculations, for some fixed pair i,j, the probability that the distortion

$$\frac{\|f(v_i) - f(v_j)\|^2}{\|v_i - v_j\|^2}$$

does not lie in the range  $[(1-\epsilon),(1+\epsilon)]$  is at most  $\frac{2}{n^{(2+\alpha)}}$ . Using the trivial union bound with  $\binom{n}{2}$  pairs, the chance that some pair of points suffers a large distortion is at most:

$$\binom{n}{2} \frac{2}{n^{(2+\alpha)}} = \frac{1}{n^{\alpha}} \left( 1 - \frac{1}{n} \right) \le \frac{1}{n^{\alpha}}.$$

Hence f has the desired properties with probability at least  $1 - \frac{1}{n^{\alpha}}$ . This gives us a randomized polynomial time algorithm.

#### Proof of Lemma 1

▶ For Lemma 1(a),

$$\begin{split} \mathbf{Prob}(L \leq \beta \mu) &= \mathbf{Prob} \left( \sum_{i=1}^k x_i^2 \leq \beta \mu (\sum_{i=1}^p x_i^2) \right) \\ &= \mathbf{Prob} \left( \beta \mu \sum_{i=1}^p x_i^2 - \sum_{i=1}^k x_i^2 \geq 0 \right) \\ &= \mathbf{Prob} \left[ \exp \left( t \beta \mu \sum_{i=1}^p x_i^2 - t \sum_{i=1}^k x_i^2 \right) \geq 1 \right], \quad (t > 0) \\ &\leq \mathbf{E} \left[ \exp \left( t \beta \mu \sum_{i=1}^p x_i^2 - t \sum_{i=1}^k x_i^2 \right) \right] \\ &\qquad \qquad (\text{by Markov's inequality}) \end{split}$$

# **Proof of Lemma 1 (continued)**

$$\begin{split} r.h.s. = & \Pi_{i=1}^k \, \mathbf{E} \exp(t(\beta \mu - 1) x_i^2) \Pi_{i=k+1}^p \, \mathbf{E} \exp(t\beta \mu x_i^2) \\ = & (\mathbf{E} \exp(t(\beta \mu - 1) x^2))^k (\mathbf{E} \exp(t\beta \mu x^2))^{p-k} \\ = & (1 - 2t(\beta \mu - 1))^{-k/2} (1 - 2t\beta \mu)^{-(p-k)/2} =: g(t) \end{split}$$

where the last equation uses the fact that if  $X \sim \mathcal{N}(0,1)$ , then

$$\mathbf{E}[e^{sX^2}] = \frac{1}{\sqrt{(1-2s)}},$$

for 
$$-\infty < s < 1/2$$
.

# **Proof of Lemma 1 (continued)**

- Now we will refer to last expression as g(t).
  - The last line of derivation gives us the additional constraints that  $t\beta\mu \leq 1/2$  and  $t(\beta\mu 1) \leq 1/2$ , and so we have  $0 < t < 1/(2\beta\mu)$ .
  - Now to minimize g(t), which is equivalent to maximize

$$h(t) = 1/g(t) = (1 - 2t(\beta\mu - 1))^{k/2} (1 - 2t\beta\mu)^{(p-k)/2}$$

in the interval  $0 < t < 1/(2\beta\mu)$ . Setting the derivative h'(t) = 0, we get the maximum is achieved at

$$t_0 = \frac{1 - \beta}{2\beta(p - \beta k)}$$

Hence we have

$$h(t_0) = \left(\frac{p-k}{p-k\beta}\right)^{(p-k)/2} \left(\frac{1}{\beta}\right)^{k/2},$$

and this is exactly what we need.

▶ Similar derivation is for the proof of Lemma 1 (b).

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# **Locality Sensitive Hashing (LSH)**

• (M.S. Charikar 2002) A **locality sensitive hashing** scheme is a distribution on a family  $\mathcal{F}$  of hash functions operating on a collection of objects, such that for two objects x, y

$$\mathbf{Prob}_{h \in \mathcal{F}}[h(x) = h(y)] = \sin(x, y)$$

where  $sim(x,y) \in [0,1]$  is some similarity function defined on the collection of objects.

Such a scheme leads to efficient (sub-linear) algorithms for approximate nearest neighbor search and clustering.

# **LSH** via Random Projections

▶ (Goemans and Williamson (1995); Charikar (2002)) Given a collection of vectors in  $\mathbb{R}^d$ , we consider the family of hash functions defined as follows: We choose a random vector  $\vec{r}$  from the d-dimensional Gaussian distribution (i.e. each coordinate is drawn the 1-dimensional Gaussian distribution). Corresponding to this vector  $\vec{r}$ , we define a hash function  $h_{\vec{r}}$  as follows:

$$h_{\vec{r}}(\vec{u}) = \mathbf{sign}(\vec{r} \cdot \vec{u}) = \begin{cases} 1 & \text{if } \vec{r} \cdot \vec{u} \ge 0 \\ -1 & \text{if } \vec{r} \cdot \vec{u} < 0 \end{cases}$$

Then for vectors  $\vec{u}$  and  $\vec{v}$ 

$$\mathbf{Pr}\left[h_{\vec{r}}(\vec{u}) = h_{\vec{r}}(\vec{v})\right] = 1 - \frac{\theta(\vec{u}, \vec{v})}{\pi}$$

# **Compressed Sensing**

- ▶ Compressive sensing can be traced back to 1950s in signal processing in geography. Its modern version appeared in LASSO (Tibshirani, 1996) and Basis Pursuit (Chen-Donoho-Saunders, 1998), and achieved a highly noticeable status after 2005 due to the work by Candes and Tao et al.
- ▶ The basic problem of compressive sensing can be expressed by the following under-determined linear algebra problem. Assume that a signal  $x^* \in \mathbb{R}^p$  is sparse with respect to some basis (measurement matrix)  $A \in \mathbb{R}^{n \times p}$  or  $A \in \mathbb{R}^{n \times p}$  where n < p, given measurement  $b = Ax^* = Ax^* \in \mathbb{R}^n$ , how can one recover  $x^*$  by solving the linear equation system

$$Ax = b? (4)$$

## **Sparsity**

As n < p, it is an under-determined problem, whence without further constraint, the problem does not have an unique solution. To overcome this issue, one popular assumption is that the signal  $x^*$  is sparse, namely the number of nonzero components  $\|x^*\|_0 := \#\{x_i^* \neq 0: 1 \leq i \leq p\}$  is small compared to the total dimensionality p. Figure below gives an illustration of such sparse linear equation problem.

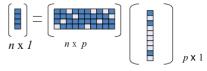


Figure: Illustration of Compressive Sensing (CS). A is a rectangular matrix with more columns than rows. The dark elements represent nonzero elements while the light ones are zeroes. The signal vector  $x^*$ , although high dimensional, is sparse.

Without loss of generality, we assume each column of design matrix  $A=[A_1,\ldots,A_p]$  has being standardized, that is,  $\|A_j\|_2=1$ , j=1,...,p.

▶ With such a sparse assumption above, a simple idea is to find the sparsest solution satisfying the measurement equation:

$$(P_0)$$
 min  $||x||_0$  (5)  
  $s.t.$   $Ax = b.$ 

► This is an NP-hard combinatorial optimization problem.

# A Greedy Algorithm: Orthogonal Matching Pursuit

```
Input A, b.
Output x.
initialization: r_0 = b, x_0 = 0, S_0 = \emptyset.
repeat if \|r_t\|_2 > \varepsilon,
1. \ j_t = \arg\max_{1 \leq j \leq p} |\langle A_j, r_{t-1} \rangle|.
2. \ S_t = S_{t-1} \cup j_t.
3. \ x_t = \arg\min_{x \in \mathbb{R}^p} \|b - A_{S_t}x\|.
4. \ r_t = b - Ax_t.
return x^t
```

- ▶ Stephane Mallat and Zhifeng Zhang (1993), choose the column of maximal correlation with residue, as the steepest descent in residue.
- ▶ Joel Tropp (2004) shows that OMP recovers  $x^*$  under the Incoherence condition; Tony Cai and Lie Wang (2011) extended it to noisy cases.

# Basis Pursuit (BP): $P_1$

► A convex relaxation of (5) is called *Basis Pursuit* (Chen-Donoho-Saunders, 1998),

(P<sub>1</sub>) min 
$$||x||_1 := \sum |x_i|$$
  
s.t.  $Ax = b$ . (6)

This is a tractable linear programming problem.

- Now a natural problem arises, under what conditions the linear programming problem  $(P_1)$  has the solution exactly solves  $(P_0)$ , i.e. exactly recovers the sparse signal  $x^*$ ?
  - Donoho and Huo (2001) proposed Incoherence condition; Joel Tropp (2004) shows that BP recovers  $x^*$  under the Incoherence condition.

#### Illustration

Figure shows different projections of a sparse vector  $x^*$  under  $l_0$ ,  $l_1$  and  $l_2$ , from which one can see in some cases the convex relaxation (6) does recover the sparse signal solution in (5).

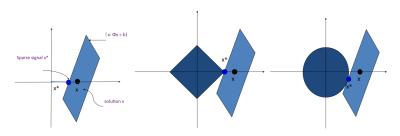


Figure: Comparison between different projections. Left: projection of  $x^*$  under  $\|\cdot\|_0$ ; middle: projection under  $\|\cdot\|_1$  which favors sparse solution; right: projection under Euclidean distance.

# Basis Pursuit De-Noising (BPDN)

When measurement noise exists, i.e.  $b = Ax^* + \varepsilon$  with bound  $\|\varepsilon\|_2$ , the following Basis Pursuit De-Noising (BPDN) are used instead

$$(BPDN) \quad \min \quad ||x||_1$$

$$s.t. \quad ||Ax - b||_2 \le \epsilon.$$

$$(7)$$

It's a convex quadratic programming problem.

▶ Similarly, Jiang-Yao-Liu-Guibas (2012) considers  $\ell_{\infty}$ -noise:

$$\min ||x||_1 
s.t. ||Ax - b||_{\infty} \le \epsilon.$$

This is a linear programming problem.

#### **LASSO**

Least Absolute Shrinkage and Selection Operator (LASSO) (Tibshirani, 1996) solves the following problem for noisy measurement:

$$(LASSO) \quad \min_{x \in \mathbb{R}^p} ||Ax - b||_2^2 + \lambda ||x||_1 \tag{8}$$

- A convex quadratic programming problem.
- ▶ Yu-Zhao (2006), Lin-Yuan (2007), Wainwright (2009) show the model selection consistency (support recovery of  $x^*$ ) of LASSO under the Irrepresentable condition.

## **Dantzig Selector**

The Dantzig Selector (Candes and Tao (2007)) is proposed to deal with noisy measurement  $b = Ax^* + \epsilon$ :

$$\min_{x \in \mathbb{R}} \|x\|_1$$

$$s.t. \quad \|A^T (Ax - b)\|_{\infty} \le \lambda$$
(9)

- ▶ A linear programming problem, more scalable than convex quadratic programming (LASSO) for large scale problems.
- Bickel, Ritov, Tsybakov (2009) show that Dantzig Selector and LASSO share similar statistical properties.

## Differential Inclusion: Inverse Scaled Spaces (ISS)

Differential inclusion:

$$\dot{\rho}_t = \frac{1}{n} A^T (b - Ax_t),\tag{10a}$$

$$\rho_t \in \partial \|x_t\|_1. \tag{10b}$$

starting at t=0 and  $\rho_0=\beta_0=0$ .

 $lackbox{ Replace } rac{
ho}{t}$  in KKT condition of LASSO by  $rac{\mathrm{d}
ho}{\mathrm{d}t}$ ,

$$\frac{\rho_t}{t} = \frac{1}{n} A^T (b - Ax_t), \qquad t = \frac{1}{\lambda}$$

to achieve unbiased estimator  $\hat{x}_t$  when it is sign-consistent.

# Differential Inclusion: Inverse Scaled Spaces (ISS) (more)

- ▶ Burger-Gilboa-Osher-Xu (2006) (in image recovery it recovers the objects in an inverse-scale order as t increases (larger objects appear in  $x_t$  first))
- ▶ Osher-Ruan-Xiong-Yao-Yin (2016) shows that its solution is a debiasing regularization path, achieving model selection consistency under nearly the same conditions of LASSO.
  - Note: if  $\hat{x}_{\tau}$  is sign consistent  $\mathbf{sign}(\hat{x}_{\tau}) = \mathbf{sign}(x^*)$ , then  $\hat{x}_{\tau} = x^* + (A^T A)^{-1} A^T \varepsilon$  which is unbiased.
  - However for LASSO, if  $\hat{x}_{\lambda}$  is sign consistent  $\mathbf{sign}(\hat{x}_{\lambda}) = \mathbf{sign}(x^*)$ , then  $\hat{x}_{\lambda} = x^* + \lambda (A^T A)^{-1} \mathbf{sign}(x^*) + (A^T A)^{-1} A^T \varepsilon$  which is biased.

### Example: Regularization Paths of LASSO vs. ISS

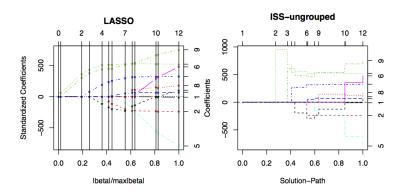


Figure: Diabetes data (Efron et al.'04) and regularization paths are different, yet bearing similarities on the order of parameters being nonzero

## **Linearized Bregman Iterations**

A damped dynamics below has a continuous solution  $x_t$  that converges to the piecewise-constant solution of (10) as  $\kappa \to \infty$ .

$$\dot{\rho}_t + \frac{\dot{x}_t}{\kappa} = -\nabla_x \ell(x_t),\tag{11a}$$

$$\rho_t \in \partial \Omega(x_t), \tag{11b}$$

Its Euler forward discretization gives the *Linearized Bregman Iterations* (LBI, Osher-Burger-Goldfarb-Xu-Yin 2005) as

$$z_{k+1} = z_k - \alpha \nabla_x \ell(x_k), \tag{12a}$$

$$x_{k+1} = \kappa \cdot \operatorname{prox}_{\Omega}(z_{k+1}), \tag{12b}$$

where  $z_{k+1}=\rho_{k+1}+\frac{x_{k+1}}{\kappa}$ , the initial choice  $z_0=x_0=0$  (or small Gaussian), parameters  $\kappa>0,\ \alpha>0,\ \nu>0$ , and the proximal map associated with a convex function  $\Omega$  is defined by

$$\operatorname{prox}_{\Omega}(z) = \arg\min_{x} \frac{1}{2} ||z - x||^2 + \Omega(x).$$

## **Uniform Recovery Conditions**

- ▶ Under which conditions we can recover arbitrary k-sparse  $x^* \in \mathbb{R}^p$  by those algorithms, for  $k = |\mathbf{supp}(x^*)| \ll n < p$ ?
- Now we turn to several conditions presented in literature, under which the algorithms above can recover  $x^*$ . Below  $A_S$  denotes the columns of A corresponding to the indices in  $S = \mathbf{supp}(x^*)$ ;  $A^*$  denotes the conjugate of matrix A, which is  $A^T$  if A is real.

## **Uniform Recovery Conditions: a) Uniqueness**

a) Uniqueness. The following condition ensures the uniqueness of k-sparse  $x^*$  satisfying  $b=Ax^*$ :

$$A_S^*A_S \geq rI, \qquad \text{for some } r>0 \text{,}$$

without which one may have more than one k-sparse solutions in solving  $b=A_Sx$ .

## **Uniform Recovery Conditions: b) Incoherence**

b) Incoherence. Donoho-Huo (2001) shows the following sufficient condition

$$\mu(A) := \max_{i \neq j} |\langle A_i, A_j \rangle| < \frac{1}{2k - 1},$$

for sparse recovery by BP, which is later improved by Elad-Bruckstein (2001) to be

$$\mu(A) < \frac{\sqrt{2} - \frac{1}{2}}{k}.$$

This condition is numerically **verifiable**, so the simplest condition.

## Uniform Recovery Conditions: c) Irrepresentable

c) Irrepresentable condition. It is also called the Exact Recovery
Condition (ERC) by Joel Tropp (2004), which shows that
under the following condition

$$M =: ||A_{S^c}^* A_S (A_S^* A_S)^{-1}||_{\infty} < 1,$$

both OMP and BP recover  $x^*$ .

- This condition is unverifiable since the true support set S is unknown.
- "Irrepresentable" is due to Yu and Zhao (2006) for proving LASSO's model selection consistency under noise, based on the fact that the regression coefficients of  $A_j \sim A_S \beta + \varepsilon$  for  $j \in S^c$ , are the row vectors of  $A_{S^c}^*A_S(A_S^*A_S)^{-1}$ , suggesting that columns of  $A_S$  can not be linearly represented by columns of  $A_{S^c}$ .

### Incoherence vs. Irrepresentable

► Tropp (2004) also shows that Incoherence condition is strictly stronger than the Irrepresentable condition in the following sense:

$$\mu < \frac{1}{2k-1} \Rightarrow M \le \frac{k\mu}{1 - (k-1)\mu} < 1.$$
 (13)

▶ On the other hand, Tony Cai et al. (2009, 2011) shows that the Irrepresentable and the Incoherence condition are **both tight** in the sense that if it fails, there exists data A,  $x^*$ , and b such that sparse recovery is not possible.

# Uniform Recovery Conditions: d) Restricted Isometry Property

d) Restricted-Isometry-Property (RIP) For all k-sparse  $x \in \mathbb{R}^p$ ,  $\exists \delta_k \in (0,1)$ , s.t.

$$(1 - \delta_k) \|x\|_2^2 \le \|Ax\|_2^2 \le (1 + \delta_k) \|x\|_2^2.$$

- ▶ This is the most popular condition by Candes-Romberg-Tao (2006).
- Although RIP is not easy to be verified, Johnson-Lindestrauss
   Lemma says some suitable random matrices will satisfy RIP with high probability.

# Restricted Isometry Property for Uniform Exact Recovery

Candes (2008) shows that under RIP, uniqueness of  $P_0$  and  $P_1$  can be guaranteed for all k-sparse signals, often called *uniform exact recovery*.

#### **Theorem**

The following holds for all k-sparse  $x^*$  satisfying  $Ax^* = b$ .

- ▶ If  $\delta_{2k} < 1$ , then problem  $P_0$  has a unique solution  $x^*$ ;
- ▶ If  $\delta_{2k} < \sqrt{2} 1$ , then the solution of  $P_1$  (BP) has a unique solution  $x^*$ , i.e. recovers the original sparse signal  $x^*$ .

## Restricted Isometry Property for Stable Noisy Recovery

Under noisy measurement  $b = Ax^* + \varepsilon$ , Candes (2008) also shows that RIP leads to stable recovery of the true sparse signal  $x^*$  using BPDN.

#### **Theorem**

Suppose that  $\|\varepsilon\|_2 \leq \epsilon$ . If  $\delta_{2k} < \sqrt{2} - 1$ , then

$$\|\hat{x} - x^*\|_2 \le C_1 k^{-1/2} \sigma_k^1(x^*) + C_2 \epsilon,$$

where  $\hat{x}$  is the solution of BPDN and

$$\sigma_k^1(x^*) = \min_{\mathbf{supp}(y) \le k} ||x^* - y||_1$$

is the best k-term approximation error in  $l_1$  of  $x^*$ .

#### $JL \Rightarrow RIP$

- Johnson-Lindenstrauss Lemma ensures RIP with high probability.
- ▶ Baraniuk, Davenport, DeVore, and Wakin (2008) show that in the proof of Johnson-Lindenstrauss Lemma, one essentially establishes that a random matrix  $A \in \mathbb{R}^{n \times p}$  with each element i.i.d. sampled according to some distribution satisfying certain bounded moment conditions, has  $||Ax||_2^2$  concentrated around its mean  $\mathbf{E} \|Ax\|_2^2 = \|x\|_2^2$  (see Appendix), *i.e.*

$$\mathbb{E} \|Ax\|_2^2 = \|x\|_2^2$$
 (see Appendix), *i.e.*

$$\mathbf{Prob}\left(\left|\|Ax\|_{2}^{2} - \|x\|_{2}^{2}\right| \ge \epsilon \|x\|_{2}^{2}\right) \le 2e^{-nc_{0}(\epsilon)}.\tag{14}$$

With this one can establish a bound on the action of A on k-sparse x by an union bound via covering numbers of k-sparse signals.

## JL ⇒ RIP: Key Lemma

#### Lemma

Let  $A \in \mathbb{R}^{n \times p}$  be a random matrix satisfying the concentration inequality (14). Then for any  $\delta \in (0,1)$  and any set all T with |T|=k < n, the following holds

$$(1 - \delta) \|x\|_2 \le \|Ax\|_2 \le (1 + \delta) \|x\|_2 \tag{15}$$

for all x whose support is contained in T, with probability at least

$$1 - 2\left(\frac{12}{\delta}\right)^k e^{-c_0(\delta/2)n}.\tag{16}$$

#### **Proof of Lemma**

It suffices to prove the results when  $||x||_2 = 1$  as A is linear.

- Let  $X_T:=\{x: \mathbf{supp}(x)=T, \|x\|_2=1\}$ . We first choose  $Q_T$ , a  $\delta/4$ -cover of  $X_T$ , such that for every  $x\in X_T$  there exists  $q\in Q_T$  satisfying  $\|q-x\|_2\leq \delta/4$ . Since  $X_T$  has dimension at most k, it is well-known from covering numbers that the capacity  $\#(Q_T)\leq (12/\delta)^k$ .
- Now we are going to apply the union bound of (14) to the set  $Q_T$  with  $\epsilon = \delta/2$ . For each  $q \in Q_T$ , with probability at most  $2e^{-c_0(\delta/2)n}$ ,  $|Aq||_2^2 \|q\|_2^2 \ge \delta/2\|q\|_2^2$ . Hence for all  $q \in Q_T$ , the same bound holds with probability at most

$$2\#(Q_T)e^{-c_0(\delta/2)n} \le 2\left(\frac{12}{\delta}\right)^k e^{-c_0(\delta/2)n}.$$

## **Proof Lemma (continued)**

 $\blacktriangleright$  Now we define  $\alpha$  to be the smallest constant such that

$$||Ax||_2 \le (1+\alpha)||x||_2$$
, for all  $x \in X_T$ .

We can show that  $\alpha \leq \delta$  with the same probability.

► For this, pick up a  $q \in Q_T$  such that  $||q - x||_2 \le \delta/4$ , whence by the triangle inequality

$$||Ax||_2 \le ||Aq||_2 + ||A(x-q)||_2 \le 1 + \delta/2 + (1+\alpha)\delta/4.$$

This implies that  $\alpha \leq \delta/2 + (1+\alpha)\delta/4$ , whence  $\alpha \leq 3\delta/4/(1-\delta/4) \leq \delta$ . This gives the upper bound. The lower bound also follows this since

$$||Ax||_2 \ge ||Aq||_2 - ||A(x-q)||_2 \ge 1 - \delta/2 - (1+\delta)\delta/4 \ge 1 - \delta,$$

which completes the proof.

#### RIP Theorem

▶ With this lemma, note that there are at most (<sup>p</sup><sub>k</sub>) subspaces of k-sparse, an union bound leads to the following result for RIP.

#### **Theorem**

Let  $A \in \mathbb{R}^{n \times p}$  be a random matrix satisfying the concentration inequality (14) and  $\delta \in (0,1)$ . There exists  $c_1,c_2>0$  such that if

$$k \le c_1 \frac{n}{\log(p/k)}$$

the following RIP holds for all k-sparse x,

$$(1 - \delta_k) \|x\|_2^2 \le \|Ax\|_2^2 \le (1 + \delta_k) \|x\|_2^2$$

with probability at least  $1 - 2e^{-c_2n}$ .

#### **Proof of RIP Theorem**

#### Proof.

For each of k-sparse signal  $(X_T)$ , RIP fails with probability at most

$$2\left(\frac{12}{\delta}\right)^k e^{-c_0(\delta/2)n}.$$

There are  $\binom{p}{k} \leq (ep/k)^k$  such subspaces. Hence, RIP fails with probability at most

$$2\left(\frac{ep}{k}\right)^k \left(\frac{12}{\delta}\right)^2 e^{-c_0(\delta/2)n} = 2e^{-c_0(\delta/2)n + k[\log(ep/k) + \log(12/\delta)]}.$$

Thus for a fixed  $c_1 > 0$ , whenever  $k \le c_1 n / \log(p/k)$ , the exponent above will be  $\le -c_2 n$  provided that

$$c_2 \le c_0(\delta/2) - c_1(1 + (1 + \log(12/\delta))/\log(p/k).$$

Note that one can always choose  $c_2 > 0$  if  $c_1 > 0$  is small enough.

## **Summary**

The following results are about mean estimation under noise:

- ▶ Johnson-Lindenstrauss Lemma tells: random projections give a universal basis to achieve uniformly almost isometric embedding, using  $O(\varepsilon^{-2}\log n)$  number of projections
- Various Applications
  - Dimensionality reduction: PCA or MDS
  - Locality Sensitive Hashing: clustering, nearest neighbor search, etc.
  - Compressed Sensing: random design satisfying Restricted Isometry Property with high probability

#### **Outline**

Recall: PCA and MDS

#### Random Projections

Example: Human Genomics Diversity Project

Johnson-Lindenstrauss Lemma

Proofs

#### Applications of Random Projections

Locality Sensitive Hashing

Compressed Sensing

Algorithms: BP, OMP, LASSO, Dantzig Selector, ISS, LBI etc.

From Johnson-Lindenstrauss Lemma to RII

#### Appendix: A Simple Version of Johnson-Lindenstrauss Lemma

## A Simple Version of Johnson-Lindenstrauss Lemma

## Theorem (Simplified Johnson-Lindenstrauss Lemma)

Let  $A=[A_{ij}]^{k\times d}$  where  $A_{ij}\sim\mathcal{N}(0,1)$  and  $R=A/\sqrt{k}$ . For any  $0<\epsilon<1$  and any positive integer k, the following holds for all  $0\neq x\in\mathbb{R}^d$ ,

$$(1 - \epsilon) \le \frac{\parallel Rx \parallel^2}{\|x\|^2} \le (1 + \epsilon),$$
 (17)

or for all  $x \neq y \in \mathbb{R}^d$ ,

$$1 - \epsilon \le \frac{\|Rx - Ry\|^2}{\|x - y\|^2} \le 1 + \epsilon \tag{18}$$

with probability at least  $1-2\exp\left(-\frac{k\varepsilon^2}{4}(1-2\varepsilon/3)\right)$ .

#### Remark

- This version of JL-Lemma is essentially used in the derivation of RIP in compressed sensing.
- Extension to sub-Gaussian distributions with bounded moment conditions can be found in Joseph Salmon's lecture notes.
- ▶ Given n sample points  $x_i \in V$ . If we let

$$k \ge 4(1 + \alpha/2)(\epsilon^2/2 - \epsilon^3/3)^{-1} \ln n,$$

then

$$\mathbb{P}\left(\|Ru\|^2 \ge 1 + \varepsilon\right) \le \exp(-(2 + \alpha)\log n) = \left(\frac{1}{n}\right)^{2+\alpha},$$

a union of  $\binom{n}{2}$  probabilistic bounds gives the full JL-Lemma.

#### A Basic Lemma

#### Lemma

Let  $X \sim \mathcal{N}(0, 1)$ .

(a) For all  $t \in (-\infty, 1/2)$ ,

$$\mathbf{E}(e^{tX^2}) = \frac{1}{1 - 2t}.$$

#### Proof.

(a) follows from Gaussian integral.

#### **Proof of JL Lemma**

Let us denote  $x\in\mathbb{R}^d, u=\frac{x}{\|x\|}$  and  $Y_i$  the column values of the output, i.e  $Y_i=(Ru)_i=\sum_{j=1}^d R_{i,j}u_j$ . Then,

$$\mathbb{E}(Y_i) = \mathbb{E}\left(\sum_{j=1}^d R_{i,j}u_j\right) = \sum_{j=1}^d \mathbb{E}(R_{i,j}u_j) = \sum_{j=1}^d u_j \mathbb{E}(R_{i,j}) = 0$$

$$\operatorname{Var}(Y_i) = \operatorname{Var}\left(\sum_{j=1}^d R_{i,j}u_j\right) = \mathbb{E}\left(\sum_{j=1}^d R_{i,j}u_j\right)^2 = \sum_{j=1}^d \operatorname{Var}(R_{i,j}u_j)$$

$$= \sum_{j=1}^d u_j^2 \operatorname{Var}(R_{i,j}) = \frac{1}{k}$$

(Upper) . Defining  $Z_i = \sqrt{k}Y_i \sim \mathcal{N}(0,1)$ , one can state the following bound:

$$\begin{split} \mathbb{P}\left(\|Ru\|^2 \geq 1 + \varepsilon\right) &= \mathbb{P}(\sum_{i=1}^k ((\sqrt{k}Y_i)^2 - 1) \geq \varepsilon k) \\ &= \mathbb{P}(\sum_{i=1}^k (Z_i^2 - 1) \geq \varepsilon k) \\ &\leq e^{-t\varepsilon k} \prod_{i=1}^k \mathbf{E} \exp\left(t(Z_i^2 - 1)\right), \quad \text{(Markov Ineq.)} \\ &= e^{-tk(1+\varepsilon)} [\mathbf{E} \, e^{tZ^2}]^k \\ &= e^{-tk(1+\varepsilon)} (1-2t)^{-k/2} =: g(t) \quad \text{(Lemma (a))} \end{split}$$

Let

$$h(t) := 1/g(t) = e^{tk(1+\varepsilon)}(1-2t)^{k/2}.$$

Hence  $\min_t g(t)$  is equivalent to  $\max_t h(t)$ . Taking derivative of h(t),

$$\begin{split} 0 &= h'(t)|_{t^*} = k(1+\varepsilon)e^{tk(1+\varepsilon)}(1-2t)^{k/2} - ke^{tk(1+\varepsilon)}(1-2t)^{k/2-1})\Big|_{t^*} \\ &= ke^{t^*k(1+\varepsilon)}(1-2t^*)^{k/2-1}\left[(1+\varepsilon)(1-2t^*)-1\right] \\ &\Rightarrow t^* = \frac{1}{2} - \frac{1}{2(1+\varepsilon)} \\ &\Rightarrow g(t^*) = e^{-t^*k(1+\varepsilon)}(1-2t^*)^{-k/2} = e^{-k\varepsilon/2}(1+\varepsilon)^{k/2} \\ &= \exp\left(-\frac{k\varepsilon}{2} + \frac{k}{2}\ln(1+\varepsilon)\right) \\ &\leq \exp\left(-\frac{k\varepsilon}{2} + \frac{k}{2}(\varepsilon - \frac{\varepsilon^2}{2} + \frac{\varepsilon^3}{3})\right), \quad \text{using } \ln(1+x) \leq x - \frac{x^2}{2} + \frac{x^3}{3} \\ &= \exp\left(-\frac{k\varepsilon^2}{4} + \frac{k\varepsilon^3}{6}\right), \quad \varepsilon \in (0,1) \end{split}$$

(Lower) . Similarly

$$\begin{split} \mathbb{P}\left(\|Ru\|^2 \leq 1 - \varepsilon\right) &= \mathbb{P}(\sum_{i=1}^k (1 - (\sqrt{k}Y_i)^2) \geq \varepsilon k) \\ &= \mathbb{P}(\sum_{i=1}^k (1 - Z_i^2) \geq \varepsilon k) \\ &\leq e^{-t\varepsilon k} \prod_{i=1}^k \mathbf{E} \exp\left(t(1 - Z_i^2)\right), \quad \text{(Markov Ineq.)} \\ &= e^{tk(1-\varepsilon)} [\mathbf{E} \, e^{-tZ^2}]^k \\ &= e^{tk(1-\varepsilon)} (1 + 2t)^{-k/2} =: g(t) \quad \text{(Lemma (a))} \end{split}$$

Let

$$h(t) := 1/g(t) = e^{tk(\varepsilon - 1)}(1 + 2t)^{k/2}.$$

Taking derivative of h(t),

$$0 = h'(t)|_{t^*} = k(\varepsilon - 1)e^{tk(\varepsilon - 1)}(1 + 2t)^{k/2} + ke^{tk(\varepsilon - 1)}(1 + 2t)^{k/2 - 1}|_{t^*}$$

$$= ke^{t^*k(\varepsilon - 1)}(1 + 2t^*)^{k/2 - 1}\left[(\varepsilon - 1)(1 + 2t^*) + 1\right]$$

$$\Rightarrow t^* = \frac{1}{2(1 - \varepsilon)} - \frac{1}{2}$$

$$\Rightarrow g(t^*) = e^{t^*k(1 - \varepsilon)}(1 + 2t^*)^{-k/2} = e^{k\varepsilon/2}(1 - \varepsilon)^{k/2}$$

$$= \exp\left(\frac{k\varepsilon}{2} + \frac{k}{2}\ln(1 - \varepsilon)\right)$$

$$\leq \exp\left(\frac{k\varepsilon}{2} + \frac{k}{2}(-\varepsilon - \frac{\varepsilon^2}{2})\right), \quad \text{using } \ln(1 - x) \leq -x - \frac{x^2}{2}$$

$$= \exp\left(-\frac{k\varepsilon^2}{4}\right), \quad \varepsilon \in (0, 1)$$