# COMPSCI 514: Algorithms for Data Science

Cameron Musco University of Massachusetts Amherst. Fall 2022. Lecture 12

### Logistics

- · Problem Set 2 is due Friday, 11:59pm.
- · No quiz this week.
- The exam will be held next Thursday in class.
- We will do some midterm review in class on Tuesday. I will also hold additional office hours for midterm prep, TBD.

### Summary

#### Last Class: The Johnson-Lindenstrauss Lemma

- Intro to dimensionality reduction and low-distortion embeddings.
- Statement of the JL Lemma: we can obtain low-distortion embeddings for any set of points via random projection.

#### This Class:

- · Reduction of the JL Lemma to the 'distributional JL Lemma'.
- Proof of the distributional JL lemma.
- · Example application to clustering.

#### The Johnson-Lindenstrauss Lemma

**Johnson-Lindenstrauss Lemma:** For any set of points  $\vec{x}_1, \dots, \vec{x}_n \in \mathbb{R}^d$  and  $\epsilon > 0$  there exists a linear map  $\mathbf{\Pi} : \mathbb{R}^d \to \mathbb{R}^m$  such that  $m = O\left(\frac{\log n}{\epsilon^2}\right)$  and letting  $\tilde{x}_i = \mathbf{\Pi} \vec{x}_i$ :

For all 
$$i, j$$
:  $(1 - \epsilon) \|\vec{x}_i - \vec{x}_j\|_2 \le \|\tilde{x}_i - \tilde{x}_j\|_2 \le (1 + \epsilon) \|\vec{x}_i - \vec{x}_j\|_2$ .

Further, if  $\Pi \in \mathbb{R}^{m \times d}$  has each entry chosen i.i.d. from  $\mathcal{N}(0, 1/m)$ , it satisfies the guarantee with high probability.

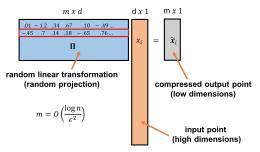
For d=1 trillion,  $\epsilon=.05$ , and n=100,000,  $m\approx 6600$ .

Very surprising! Powerful result with a simple construction: applying a random linear transformation to a set of points preserves distances between all those points with high probability.

### Random Projection

For any  $\vec{x}_1, \dots, \vec{x}_n$  and  $\Pi \in \mathbb{R}^{m \times d}$  with each entry chosen i.i.d. from  $\mathcal{N}(0, 1/m)$ , with high probability, letting  $\tilde{\mathbf{x}}_i = \Pi \vec{x}_i$ :

For all 
$$i, j : (1 - \epsilon) \|\vec{x}_i - \vec{x}_j\|_2 \le \|\tilde{\mathbf{x}}_i - \tilde{\mathbf{x}}_j\|_2 \le (1 + \epsilon) \|\vec{x}_i - \vec{x}_j\|_2$$
.



- • 
   • I is data oblivious. Stark contrast to methods like PCA.

## **Algorithmic Considerations**

- Many alternative constructions:  $\pm 1$  entries, sparse (most entries 0), Fourier structured, etc.  $\Longrightarrow$  more efficient computation of  $\tilde{\mathbf{x}}_i = \mathbf{\Pi} \vec{\mathbf{x}}_i$ .
- Data oblivious property means that once  $\Pi$  is chosen,  $\tilde{x}_1,\ldots,\tilde{x}_n$  can be computed in a stream with little memory.
- Memory needed is just O(d + nm) vs. O(nd) to store the full data set.
- Compression can also be easily performed in parallel on different servers.
- When new data points are added, can be easily compressed, without updating existing points.

#### Distributional JL

The Johnson-Lindenstrauss Lemma is a direct consequence of a closely related lemma:

**Distributional JL Lemma:** Let  $\Pi \in \mathbb{R}^{m \times d}$  have each entry chosen i.i.d. as  $\mathcal{N}(0, 1/m)$ . If we set  $m = O\left(\frac{\log(1/\delta)}{\epsilon^2}\right)$ , then for any  $\vec{y} \in \mathbb{R}^d$ , with probability  $\geq 1 - \delta$   $(1 - \epsilon) \|\vec{y}\|_2 \leq \|\Pi\vec{y}\|_2 \leq (1 + \epsilon) \|\vec{y}\|_2$ 

Applying a random matrix  $\Pi$  to any vector  $\vec{y}$  preserves  $\vec{y}$ 's norm with high probability.

- Like a low-distortion embedding, but for the length of a compressed vector rather than distances between vectors.
- · Can be proven from first principles.

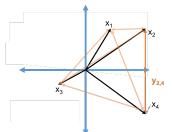
 $\Pi \in \mathbb{R}^{m \times d}$ : random projection matrix. d: original dimension. m: compressed dimension,  $\epsilon$ : embedding error,  $\delta$ : embedding failure prob.

#### Distributional JL $\implies$ JL

**Distributional JL Lemma**  $\Longrightarrow$  **JL Lemma:** Distributional JL show that a random projection  $\Pi$  preserves the norm of any y. The main JL Lemma says that  $\Pi$  preserves distances between vectors.

Since  $\Pi$  is linear these are the same thing!

**Proof:** Given  $\vec{x}_1, \dots, \vec{x}_n$ , define  $\binom{n}{2}$  vectors  $\vec{y}_{ij}$  where  $\vec{y}_{ij} = \vec{x}_i - \vec{x}_j$ .



• If we choose  $\Pi$  with  $m = O\left(\frac{\log 1/\delta}{\epsilon^2}\right)$ , for each  $\vec{y}_{ij}$  with probability  $> 1 - \delta$  we have:

$$(1 \quad c) \| \vec{\sigma} \vec{\nabla} \vec{\nabla} \| \neq \| \mathbf{n} \vec{\sigma} \mathbf{n} (\vec{\nabla} \vec{\nabla}) \vec{\nabla} \vec{\nabla} \| \neq (1 + c) \| \vec{\sigma} \vec{\nabla} \vec{\nabla} \vec{\nabla} \|$$

#### Distributional $JL \implies JL$

Claim: If we choose  $\Pi$  with i.i.d.  $\mathcal{N}(0, 1/m)$  entries and  $m = O\left(\frac{\log(1/\delta')}{\epsilon^2}\right)$ , letting  $\tilde{\mathbf{x}}_i = \Pi \vec{x}_i$ , for each pair  $\vec{x}_i, \vec{x}_j$  with probability  $\geq 1 - \delta'$  we have:

$$(1 - \epsilon) \|\vec{x}_i - \vec{x}_i\|_2 \le \|\tilde{\mathbf{x}}_i - \tilde{\mathbf{x}}_i\|_2 \le (1 + \epsilon) \|\vec{x}_i - \vec{x}_i\|_2.$$

With what probability are all pairwise distances preserved?

**Union bound:** With probability  $\geq 1 - \binom{n}{2} \cdot \delta'$  all pairwise distances are preserved.

Apply the claim with  $\delta' = \delta/\binom{n}{2}$ .  $\Longrightarrow$  for  $m = O\left(\frac{\log(1/\delta')}{\epsilon^2}\right)$ , all pairwise distances are preserved with probability  $\geq 1 - \delta$ .

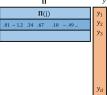
$$m = O\left(\frac{\log(1/\delta')}{\epsilon^2}\right) = O\left(\frac{\log(\binom{n}{2}/\delta)}{\epsilon^2}\right) = O\left(\frac{\log(n^2/\delta)}{\epsilon^2}\right) = O\left(\frac{\log(n/\delta)}{\epsilon^2}\right)$$

Yields the JL lemma.

**Distributional JL Lemma:** Let  $\Pi \in \mathbb{R}^{m \times d}$  have each entry chosen i.i.d. as  $\mathcal{N}(0, 1/m)$ . If we set  $m = O\left(\frac{\log(1/\delta)}{\epsilon^2}\right)$ , then for any  $\vec{y} \in \mathbb{R}^d$ , with probability  $\geq 1 - \delta$ 

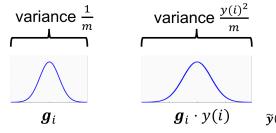
$$(1 - \epsilon) \|\vec{y}\|_2 \le \|\mathbf{\Pi}\vec{y}\|_2 \le (1 + \epsilon) \|\vec{y}\|_2$$

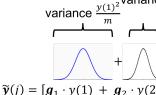
- Let  $\tilde{\mathbf{y}}$  denote  $\mathbf{\Pi}\vec{\mathbf{y}}$  and let  $\mathbf{\Pi}(j)$  denote the  $j^{th}$  row of  $\mathbf{\Pi}$ .
- For any j,  $\tilde{\mathbf{y}}(j) = \langle \mathbf{\Pi}(j), \vec{y} \rangle = \sum_{\mathbf{n}=1}^{d} \mathbf{g}_{i} \cdot \vec{\mathbf{y}}(i)$  where  $\mathbf{g}_{i} \sim \mathcal{N}(0, 1/m)$ .



 $\vec{y} \in \mathbb{R}^d$ : arbitrary vector,  $\tilde{\mathbf{y}} \in \mathbb{R}^m$ : compressed vector,  $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$ : random projection. d: original dim. m: compressed dim,  $\epsilon$ : error,  $\delta$ : failure prob.

- Let  $\tilde{\mathbf{y}}$  denote  $\mathbf{\Pi}\vec{\mathbf{y}}$  and let  $\mathbf{\Pi}(j)$  denote the  $j^{th}$  row of  $\mathbf{\Pi}$ .
- For any j,  $\tilde{\mathbf{y}}(j) = \langle \mathbf{\Pi}(j), \vec{y} \rangle = \sum_{i=1}^d \mathbf{g}_i \cdot \vec{y}(i)$  where  $\mathbf{g}_i \sim \mathcal{N}(0, 1/m)$ .
- $\mathbf{g}_i \cdot \vec{y}(i) \sim \mathcal{N}(0, \frac{\vec{y}(i)^2}{m})$ : normally distributed with variance  $\frac{\vec{y}(i)^2}{m}$ .





What is the distribution of  $\tilde{\mathbf{v}}(i)$ ? Also Gaussian!

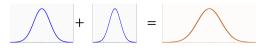
 $\vec{y} \in \mathbb{R}^d$ : arbitrary vector,  $\tilde{\mathbf{y}} \in \mathbb{R}^m$ : compressed vector,  $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$ : random projection mapping  $\vec{y} \to \tilde{\mathbf{y}}$ .  $\mathbf{\Pi}(j)$ :  $j^{th}$  row of  $\mathbf{\Pi}$ , d: original dimension. m: compressed dimension,  $\mathbf{g}_i$ : normally distributed random variable.

Letting  $\tilde{\mathbf{y}} = \mathbf{\Pi} \vec{\mathbf{y}}$ , we have  $\tilde{\mathbf{y}}(j) = \langle \mathbf{\Pi}(j), \vec{\mathbf{y}} \rangle$  and:

$$\tilde{\mathbf{y}}(j) = \sum_{i=1}^{d} \mathbf{g}_{i} \cdot \vec{\mathbf{y}}(i) \text{ where } \mathbf{g}_{i} \cdot \vec{\mathbf{y}}(i) \sim \mathcal{N}\left(0, \frac{\vec{\mathbf{y}}(i)^{2}}{m}\right).$$

Stability of Gaussian Random Variables. For independent  $a \sim \mathcal{N}(\mu_1, \sigma_1^2)$  and  $b \sim \mathcal{N}(\mu_2, \sigma_2^2)$  we have:

$$a + b \sim \mathcal{N}(\mu_1 + \mu_2, \sigma_1^2 + \sigma_2^2)$$



Thus,  $\tilde{\mathbf{y}}(j) \sim \mathcal{N}(0, \frac{\vec{\mathbf{y}}(1)^2}{m} + \frac{\vec{\mathbf{y}}(2)^2}{m} + \ldots + \frac{\vec{\mathbf{y}}(d)^2}{m} \frac{\|\vec{\mathbf{y}}\|_2^2}{m})$  I.e.,  $\tilde{\mathbf{y}}$  itself is a random Gaussian vector. Rotational invariance of the Gaussian distribution.

 $\vec{y} \in \mathbb{R}^d$ : arbitrary vector,  $\tilde{\mathbf{y}} \in \mathbb{R}^m$ : compressed vector,  $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$ : random projection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ :  $\vec{y} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ :  $\vec{y} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}} \in \mathbb{R}^d$ : random rejection mapping  $\vec{y} = \tilde{\mathbf{y}}$ 

So far: Letting  $\Pi \in \mathbb{R}^{d \times m}$  have each entry chosen i.i.d. as  $\mathcal{N}(0, 1/m)$ , for any  $\vec{y} \in \mathbb{R}^d$ , letting  $\tilde{\mathbf{y}} = \Pi \vec{y}$ :

$$\tilde{\mathbf{y}}(j) \sim \mathcal{N}(0, \|\vec{\mathbf{y}}\|_2^2/m).$$

What is  $\mathbb{E}[\|\tilde{\mathbf{y}}\|_2^2]$ ?

$$\mathbb{E}[\|\tilde{\mathbf{y}}\|_{2}^{2}] = \mathbb{E}\left[\sum_{j=1}^{m} \tilde{\mathbf{y}}(j)^{2}\right] = \sum_{j=1}^{m} \mathbb{E}[\tilde{\mathbf{y}}(j)^{2}]$$
$$= \sum_{j=1}^{m} \frac{\|\vec{\mathbf{y}}\|_{2}^{2}}{m} = \|\vec{\mathbf{y}}\|_{2}^{2}$$

So  $\tilde{\mathbf{y}}$  has the right norm in expectation.

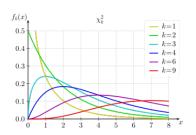
How is  $\|\tilde{\mathbf{y}}\|_2^2$  distributed? Does it concentrate?

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So far: Letting  $\Pi \in \mathbb{R}^{d \times m}$  have each entry chosen i.i.d. as  $\mathcal{N}(0, 1/m)$ , for any  $\vec{y} \in \mathbb{R}^d$ , letting  $\tilde{\mathbf{y}} = \Pi \vec{y}$ :

$$\tilde{\mathbf{y}}(j) \sim \mathcal{N}(0, \|\vec{\mathbf{y}}\|_2^2/m)$$
 and  $\mathbb{E}[\|\tilde{\mathbf{y}}\|_2^2] = \|\vec{\mathbf{y}}\|_2^2$ 

 $\|\tilde{\mathbf{y}}\|_2^2 = \sum_{i=1}^m \tilde{\mathbf{y}}(j)^2$  a Chi-Squared random variable with m degrees of freedom (a sum of m squared independent Gaussians)

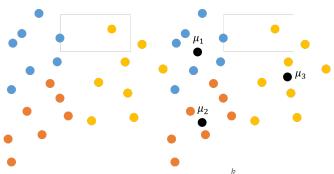


**Lemma:** (Chi-Squared Concentration) Letting **Z** be a Chi-Squared random variable with *m* degrees of freedom,

$$\Pr[|\mathbf{Z} - \mathbb{E}\mathbf{Z}| > \epsilon \mathbb{E}\mathbf{Z}] < 2e^{-m\epsilon^2/8}$$

## Example Application: k-means clustering

**Goal:** Separate *n* points in *d* dimensional space into *k* groups.



k-means Objective: 
$$Cost(C_1, ..., C_k) = \min_{C_1, ..., C_k} \sum_{j=1}^{n} \sum_{\vec{x} \in C_k} ||\vec{x} - \mu_j||_2^2$$
.

Write in terms of distances:

$$Cost(C_1, ..., C_k) = \min_{C_1, ... C_k} \sum_{j=1}^{\kappa} \sum_{\vec{x}_1, \vec{x}_2 \in C_k} ||\vec{x}_1 - \vec{x}_2||_2^2$$

### Example Application: k-means clustering

k-means Objective: 
$$Cost(C_1, \dots, C_k) = \min_{C_1, \dots C_k} \sum_{j=1}^k \sum_{\vec{x}_1, \vec{x}_2 \in C_k} ||\vec{x}_1 - \vec{x}_2||_2^2$$

If we randomly project to  $m = O\left(\frac{\log n}{\epsilon^2}\right)$  dimensions, for all pairs  $\vec{x}_1, \vec{x}_2$ ,

$$(1 - \epsilon) \|\vec{\mathbf{x}}_1 - \vec{\mathbf{x}}_2\|_2^2 \le \|\tilde{\mathbf{x}}_1 - \tilde{\mathbf{x}}_2\|_2^2 \le (1 + \epsilon) \|\vec{\mathbf{x}}_1 - \vec{\mathbf{x}}_2\|_2^2 \implies$$

Letting 
$$\overline{Cost}(\mathcal{C}_1, \dots, \mathcal{C}_k) = \min_{\mathcal{C}_1, \dots \mathcal{C}_k} \sum_{j=1}^R \sum_{\tilde{\mathbf{x}}_1, \tilde{\mathbf{x}}_2 \in \mathcal{C}_k} \|\tilde{\mathbf{x}}_1 - \tilde{\mathbf{x}}_2\|_2^2$$

$$(1-\epsilon)$$
Cost $(C_1,\ldots,C_k) \leq \overline{\text{Cost}}(C_1,\ldots,C_k) \leq (1+\epsilon)$ Cost $(C_1,\ldots,C_k)$ .

**Upshot:** Can cluster in m dimensional space (much more efficiently) and minimize  $\overline{Cost}(\mathcal{C}_1,\ldots,\mathcal{C}_k)$ . The optimal set of clusters will have true cost within  $1+c\epsilon$  times the true optimal. Good exercise to prove this.