

COMPSCI 514: Algorithms for Data Science

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University of Massachusetts Amherst. Spring 2026.

Lecture 11

grades open request for 1 week out of 34

- Midterm 1 grades have been posted.
- Average around a 66%. This was a lower than expected. I will have several questions on this week's quiz to try to figure out why.
- If you are worried about your grade, feel free to email me and set up a time to chat.
- Problem Set 2 is graded, and I will post grades very shortly.

Summary

We are now starting on the **Spectral Methods** unit.

- Linear algebraic techniques for working with large and high dimensional datasets.
- Today we will cover something at the intersection of the first two units: randomized methods for compressing high dimensional data.
- Low-distortion embeddings and the Johnson-Lindenstrauss (JL) Lemma.

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- Linear algebraic techniques for working with large and high dimensional datasets.
- Today we will cover something at the intersection of the first two units: randomized methods for compressing high dimensional data.
- Low-distortion embeddings and the Johnson-Lindenstrauss (JL) Lemma.

If you feel shaky on your linear algebra background, you'll want to start doing some review. See the reading material for resources.

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- The human genome contains 3 billion+ base pairs. Genetic datasets often contain information on **100s of thousands+ mutations and genetic markers**.

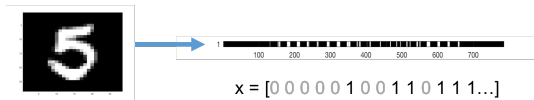
Data as Vectors and Matrices

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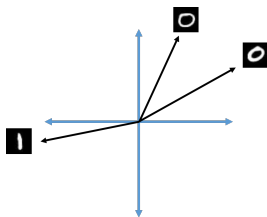
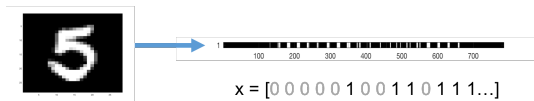
ATAGCCGTAGT \longrightarrow $x = [1\ 2\ 1\ 3\ 4\ 4\ 3\ 2\ 1\ 3\ 4]$



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Similarities/distances between vectors (e.g., $\langle x, y \rangle$, $\|x - y\|_2$) have meaning for underlying data points.

Datasets as Vectors and Matrices

Data points are interpreted as **high dimensional vectors**, with real valued entries. Data set is interpreted as a matrix.

Data Points: $\vec{x}_1, \vec{x}_2, \dots, \vec{x}_n \in \mathbb{R}^d$.

Data Set: $X \in \mathbb{R}^{n \times d}$ with i^{th} row equal to \vec{x}_i^T .

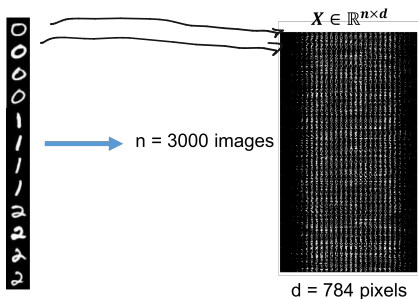
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$$[x_i] \rightarrow [x_i]^T$$

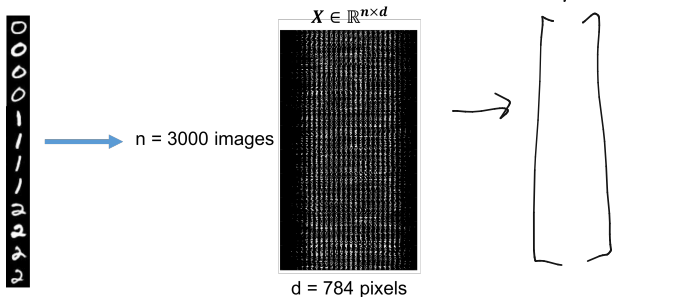


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Many data points $n \implies$ tall. Many dimensions $d \implies$ wide.

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5 $\rightarrow x = [0\ 0\ 0\ 0\ 0\ 1\ 0\ 0\ 1\ 1\ 0\ 1\ 1\ 1\dots]$ $\rightarrow \tilde{x} = [-5.5\ 4\ 3.2\ -1]$

Dimensionality Reduction

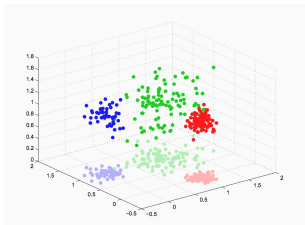
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'Lossy compression' that still preserves important information about the relationships between $\vec{x}_1, \dots, \vec{x}_n$.



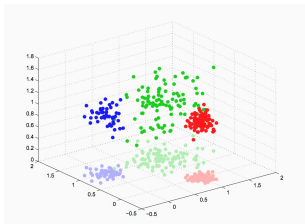
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Generally will not consider directly how well \tilde{x}_i approximates \vec{x}_i .

Dimensionality Reduction

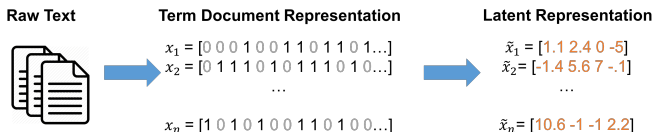
Dimensionality reduction is one of the most important techniques in data science. **What methods have you heard of?**

- PCA (principal component analysis)
 - SVD, LSA, ... - low-rank approx.
- Isomap, V-map, LLE, ...
- Auto encoders $[\tilde{x}] = \hat{f}([x])$
- TSNE

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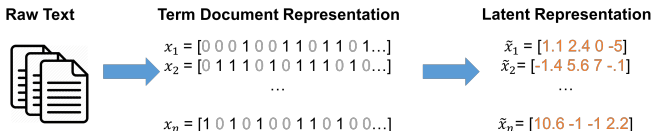


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Compressing data makes it more efficient to work with. May also remove extraneous information/noise.

Embeddings for Euclidean Space

Euclidean Low Distortion Embedding: Given $\vec{x}_1, \dots, \vec{x}_n \in \mathbb{R}^d$ and error parameter $\epsilon \geq 0$, find $\tilde{x}_1, \dots, \tilde{x}_n \in \mathbb{R}^m$ (where $m \ll d$) such that for all $i, j \in [n]$: all distances should be preserved

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

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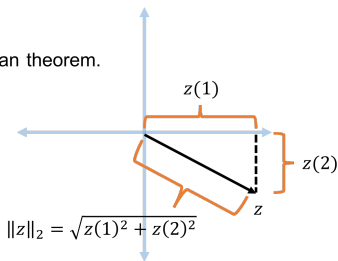
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$\| \quad \|_2$

Recall that for $\vec{z} \in \mathbb{R}^n$, $\|\vec{z}\|_2 = \sqrt{\sum_{i=1}^n \vec{z}(i)^2}$.

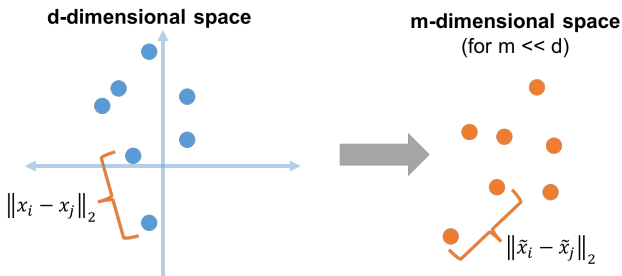
Pythagorean theorem.



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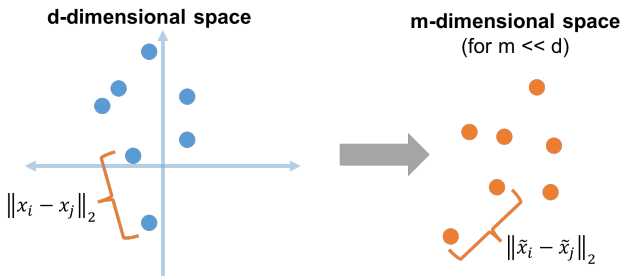
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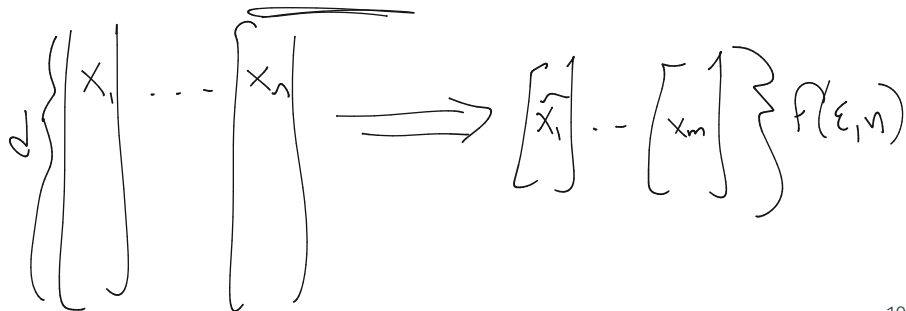
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Can use $\tilde{x}_1, \dots, \tilde{x}_n$ in place of $\vec{x}_1, \dots, \vec{x}_n$ in clustering, SVM, linear classification, near neighbor search, etc.

The Johnson-Lindenstrauss Lemma

The Johnson-Lindenstrauss Lemma tells us that for **any set of points** $\vec{x}_1, \dots, \vec{x}_n \in \mathbb{R}^d$ and any $\epsilon > 0$, we can find an ϵ -distortion embedding into m dimensions, where m depends only on the error parameter ϵ and the number of points n , but not the original dimension d .



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 $\mathbf{\Pi} : \mathbb{R}^d \rightarrow \mathbb{R}^m$ such that $m = O\left(\frac{\log n}{\epsilon^2}\right)$ and letting $\tilde{x}_i = \mathbf{\Pi}\vec{x}_i$:

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

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

(normal w/ mean 0, variance $1/m$)

Handwritten diagram illustrating the Johnson-Lindenstrauss Lemma. It shows a matrix of size m by d with entries 0 , -0.37 , and 0.65 . An arrow labeled x_i points to a vector of size d with entries x_i . A brace indicates the vector is of size d . To the right, a brace indicates the vector is of size $\frac{\log n}{\epsilon^2}$.

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For $d = 1$ trillion, $\epsilon = .05$, and $n = 100,000$, $m \approx 6600$. $= \frac{\log(100000)}{0.05^2}$

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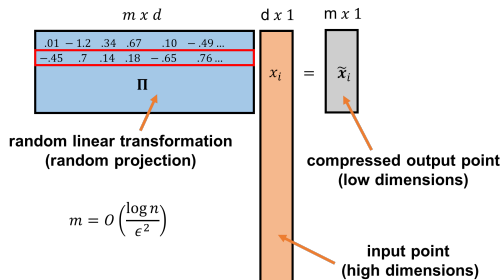
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Very surprising! Powerful result with a simple construction: applying a random linear transformation to a set of points preserves distances between all those points with high probability.

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 $\tilde{x}_i = \Pi \vec{x}_i$:

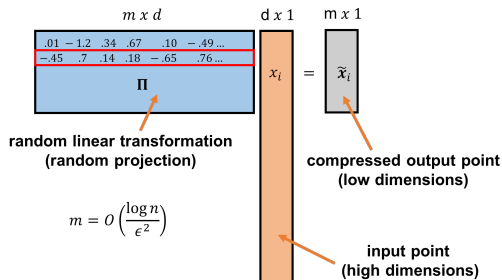
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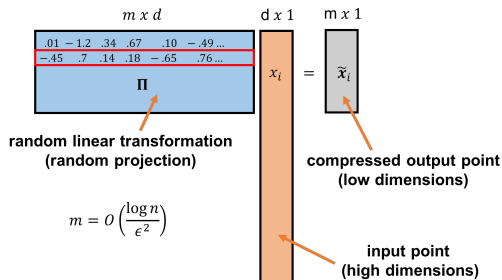


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- Π is known as a **random projection**. It is a random linear function, mapping length d vectors to length m vectors.
- Π is **data oblivious**. Stark contrast to methods like PCA.

Algorithmic Considerations

- Many alternative constructions: ± 1 entries, sparse (most entries 0), Fourier structured, etc. \implies more efficient computation of $\tilde{\mathbf{x}}_j = \mathbf{\Pi} \vec{\mathbf{x}}_j$.

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- Memory needed is just $O(d + nm)$ vs. $O(nd)$ to store the full data set.

$\mathbf{\Pi}$ has $O(dm)$ entries
 \hookrightarrow usually generated using a RNG.

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$$x_1 \dots x_n \quad \downarrow [x_i] \quad \rightarrow \quad [\tilde{x}_1] \dots [\tilde{x}_n]$$

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 "current data point" "stored compressed data"
- 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.

Johnson-Lindenstrauss Lemma Proof

Distributional JL

The Johnson-Lindenstrauss Lemma is a direct consequence of a closely related lemma: \hookrightarrow 1.74

Distributional JL Lemma: Let $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$ 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 $\geq 1 - \delta$

$$(1 - \epsilon) \|\vec{y}\|_2 \leq \|\mathbf{\Pi}\vec{y}\|_2 \leq (1 + \epsilon) \|\vec{y}\|_2 \quad \|\hat{x}_i - \tilde{x}_i\|$$

\downarrow
 $\|\tilde{y}\|_2$

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

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↑↑ show this implication

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↓
 $y_{ij} = x_i - x_j \quad (1 - \epsilon)\|\vec{y}\|_2 \leq \|\mathbf{\Pi}\vec{y}\|_2 \leq (1 + \epsilon)\|\vec{y}\|_2$

Applying a random matrix $\mathbf{\Pi}$ 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.

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Distributional JL \implies JL

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

$\vec{x}_1, \dots, \vec{x}_n$: original points, $\tilde{\vec{x}}_1, \dots, \tilde{\vec{x}}_n$: compressed points, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection matrix. d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

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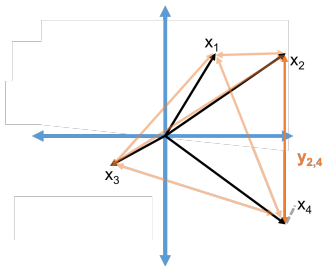
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Since Π is **linear** these are the same thing!

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$\vec{x}_1, \dots, \vec{x}_n$: original points, $\tilde{x}_1, \dots, \tilde{x}_n$: compressed points, $\Pi \in \mathbb{R}^{m \times d}$: random projection matrix. d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

Distributional JL \implies JL

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

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

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

$y_{ij} = x_i - x_j$

$$(1 - \epsilon) \|x_i - x_j\|_2 \quad \dots$$

$\vec{x}_1, \dots, \vec{x}_n$: original points, $\tilde{x}_1, \dots, \tilde{x}_n$: compressed points, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection matrix. d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

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

$$(1 - \epsilon) \|\vec{x}_i - \vec{x}_j\|_2 \leq \|\Pi(\vec{x}_i - \vec{x}_j)\|_2 \leq (1 + \epsilon) \|\vec{x}_i - \vec{x}_j\|_2$$

$$\|\Pi x_i - \Pi x_j\|_2$$

$$\|\hat{x}_i - \tilde{x}_j\|_2$$

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

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With what probability are all pairwise distances preserved?

E_{ij} = event that $\|\mathbf{x}_i - \mathbf{x}_j\|$ is not preserved

$$\Pr\left[\bigcup_{i,j} E_{ij}\right] \leq \sum \Pr[E_{ij}] = \binom{n}{2} \cdot \delta'$$

$$\text{Success prob: } 1 - \binom{n}{2} \cdot \delta'$$

$\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_n$: original points, $\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_n$: compressed points, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection matrix. d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

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

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$$\underline{1 - \delta}$$

Apply the claim with $\delta' = \delta / \binom{n}{2}$.

$\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_n$: original points, $\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_n$: compressed points, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection matrix. d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

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

$\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_n$: original points, $\tilde{\mathbf{x}}_1, \dots, \tilde{\mathbf{x}}_n$: compressed points, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection matrix. d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

δ' = failure prob for one pairs x_i, x_j

δ = overall failure prob.

Distributional JL \implies JL

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

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

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Yields the JL lemma.

Distributional JL Proof

Distributional JL Lemma: Let $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$ 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 $\geq 1 - \delta$

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

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

Distributional JL Proof

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- Let \tilde{y} denote $\mathbf{\Pi}\vec{y}$ and let $\mathbf{\Pi}(j)$ denote the j^{th} row of $\mathbf{\Pi}$.



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- Let \tilde{y} denote $\mathbf{\Pi}\vec{y}$ and let $\mathbf{\Pi}(j)$ denote the j^{th} row of $\mathbf{\Pi}$.
- For any j , $\tilde{y}(j) = \langle \mathbf{\Pi}(j), \vec{y} \rangle$

The diagram shows a square representing a matrix with a double-headed arrow pointing to it labeled $\mathbf{\Pi}(j)$. To its right is a vertical vector labeled \vec{y} . An equals sign follows, and to the right is a vertical vector labeled $\tilde{y}(j)$. The top element of this vector is also labeled $\tilde{y}(j)$.

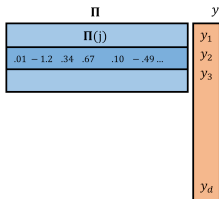
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$$(1 - \epsilon)\|\vec{y}\|_2 \leq \|\mathbf{\Pi}\vec{y}\|_2 \leq (1 + \epsilon)\|\vec{y}\|_2$$

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- For any j , $\tilde{y}(j) = \langle \mathbf{\Pi}(j), \vec{y} \rangle = \sum_{i=1}^d \mathbf{g}_i \cdot \vec{y}(i)$ where $\mathbf{g}_i \sim \mathcal{N}(0, 1/m)$.

$g_i = i^{\text{th}}$ entry of row j ($\mathbf{\Pi}(j)$)

$\mathbf{\Pi}(j)$	y_1
.01 -1.2 .34 .67 .10 -.49...	y_2
	y_3
	y_d

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

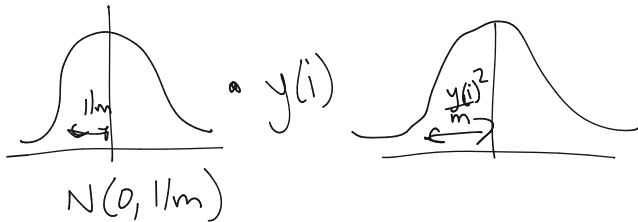
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$\vec{\mathbf{y}} \in \mathbb{R}^d$: arbitrary vector, $\tilde{\mathbf{y}} \in \mathbb{R}^m$: compressed vector, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection mapping $\vec{\mathbf{y}} \rightarrow \tilde{\mathbf{y}}$. $\mathbf{\Pi}(j)$: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

Distributional JL Proof

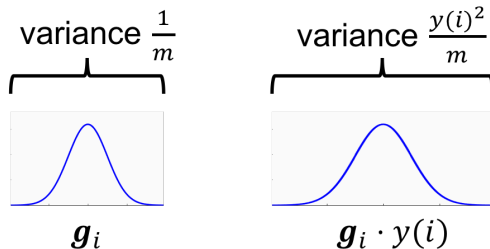
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- $\mathbf{g}_i \cdot \vec{\mathbf{y}}(i) \sim \mathcal{N}(0, \frac{\vec{\mathbf{y}}(i)^2}{m})$: normally distributed with variance $\frac{\vec{\mathbf{y}}(i)^2}{m}$.



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Distributional JL Proof

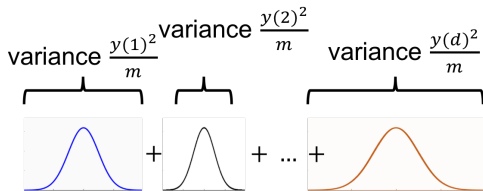
- Let $\tilde{\mathbf{y}}$ denote $\mathbf{\Pi}\vec{\mathbf{y}}$ and let $\mathbf{\Pi}(j)$ denote the j^{th} row of $\mathbf{\Pi}$.
- For any j , $\tilde{\mathbf{y}}(j) = \langle \mathbf{\Pi}(j), \vec{\mathbf{y}} \rangle = \sum_{i=1}^d \mathbf{g}_i \cdot \vec{\mathbf{y}}(i)$ where $\mathbf{g}_i \sim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

- Let $\tilde{\mathbf{y}}$ denote $\mathbf{\Pi}\vec{y}$ 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 $\frac{\vec{y}(i)^2}{m}$.



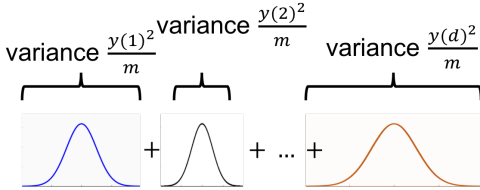
$$\tilde{\mathbf{y}}(j) = [\mathbf{g}_1 \cdot y(1) + \mathbf{g}_2 \cdot y(2) + \dots + \mathbf{g}_n \cdot y(d)]$$

$$\tilde{\mathbf{y}}(j) = \mathcal{N}\left(0, \sum_{i=1}^d \frac{y(i)^2}{m}\right) = \mathcal{N}\left(0, \frac{1}{m} \sum y(i)^2\right) = \mathcal{N}\left(0, \frac{1}{m} \|\mathbf{y}\|_2^2\right)$$

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

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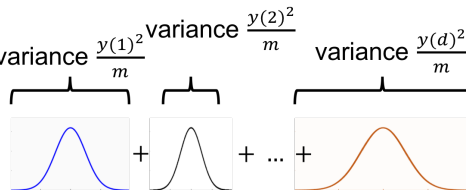

$$\tilde{\mathbf{y}}(j) = [\mathbf{g}_1 \cdot \mathbf{y}(1) + \mathbf{g}_2 \cdot \mathbf{y}(2) + \dots + \mathbf{g}_n \cdot \mathbf{y}(d)]$$

What is the distribution of $\tilde{\mathbf{y}}(j)$?

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

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The diagram illustrates the decomposition of the dot product $\tilde{\mathbf{y}}(j) = \mathbf{g}_1 \cdot y(1) + \mathbf{g}_2 \cdot y(2) + \dots + \mathbf{g}_n \cdot y(d)$. It shows three Gaussian curves representing the terms in the sum. The first curve is blue and has a variance of $\frac{y(1)^2}{m}$. The second curve is black and has a variance of $\frac{y(2)^2}{m}$. The third curve is orange and has a variance of $\frac{y(d)^2}{m}$. Brackets above each curve indicate their respective variances. The curves are separated by plus signs and an ellipsis, indicating they are summed together.

$$\tilde{\mathbf{y}}(j) = [\mathbf{g}_1 \cdot y(1) + \mathbf{g}_2 \cdot y(2) + \dots + \mathbf{g}_n \cdot y(d)]$$

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

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

Distributional JL Proof

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

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

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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:

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

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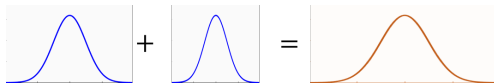
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Thus, $\tilde{\mathbf{y}}(j) \sim \mathcal{N}(0, \frac{\|\vec{y}\|_2^2}{m})$ i.e., $\tilde{\mathbf{y}}$ itself is a random Gaussian vector.

Rotational invariance of the Gaussian distribution.

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

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

$$\|\tilde{\mathbf{y}}\|_2^2 \approx \|\mathbf{y}\|_2^2$$

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

$$= \mathbb{E}\left[\sum_{j=1}^m \tilde{y}(j)^2\right] = \sum_{j=1}^m \mathbb{E}[\tilde{y}(j)^2] = \sum_{j=1}^m \frac{\|\mathbf{y}\|_2^2}{m} = \|\mathbf{y}\|_2^2$$

$$\text{Var}(\tilde{y}(j)) = \mathbb{E}[\tilde{y}(j)^2] - \left[\mathbb{E}[\tilde{y}(j)]\right]^2$$

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

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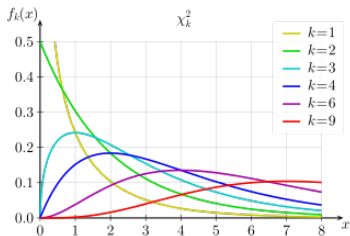
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Lemma: (Chi-Squared Concentration) Letting Z be a Chi-Squared random variable with m degrees of freedom,

$$\Pr[|Z - \mathbb{E}Z| \geq \epsilon \mathbb{E}Z] \leq 2e^{-m\epsilon^2/8}$$

apply to show $m = O\left(\frac{\log(1/\delta)}{\epsilon^2}\right)$
then $(1-\epsilon)\|\vec{y}\|_2 \leq \|\tilde{\mathbf{y}}\|_2 \leq (1+\epsilon)\|\vec{y}\|_2$

$\vec{y} \in \mathbb{R}^d$: arbitrary vector, $\tilde{\mathbf{y}} \in \mathbb{R}^m$: compressed vector, $\mathbf{\Pi} \in \mathbb{R}^{m \times d}$: random projection mapping $\vec{y} \rightarrow \tilde{\mathbf{y}}$. $\mathbf{\Pi}(j)$: j^{th} row of $\mathbf{\Pi}$, d : original dimension. m : compressed dimension, ϵ : embedding error, δ : embedding failure prob.

Distributional JL Proof

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

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

$\|\tilde{\mathbf{y}}\|_2^2 = \sum_{j=1}^m \tilde{y}(j)^2$ 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,

$$\Pr[|Z - \mathbb{E}Z| \geq \epsilon \mathbb{E}Z] \leq 2e^{-m\epsilon^2/8}.$$

If we set $m = O\left(\frac{\log(1/\delta)}{\epsilon^2}\right)$, with probability $1 - O(e^{-\log(1/\delta)}) \geq 1 - \delta$:

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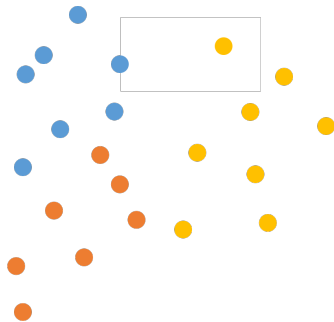
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Gives the distributional JL Lemma and thus the classic JL Lemma!

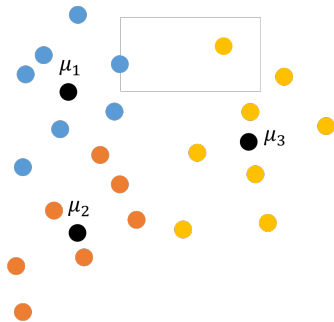
Example Application: k -means clustering

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



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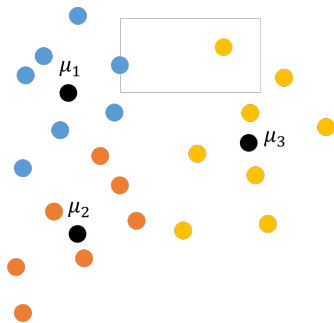
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k-means Objective: $Cost(C_1, \dots, C_k) = \min_{C_1, \dots, C_k} \sum_{j=1}^k \sum_{\vec{x} \in C_k} \|\vec{x} - \mu_j\|_2^2.$

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Write in terms of distances:

$$Cost(\mathcal{C}_1, \dots, \mathcal{C}_k) = \min_{\mathcal{C}_1, \dots, \mathcal{C}_k} \sum_{j=1}^k \sum_{\vec{x}_1, \vec{x}_2 \in \mathcal{C}_k} \|\vec{x}_1 - \vec{x}_2\|_2^2$$

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