



CS182: Introduction to Machine Learning – Dimensionality Reduction

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Recall: K -means Algorithm

- Input: $\mathcal{D} = \{(\mathbf{x}^{(n)})\}_{n=1}^N, K$

1. Initialize cluster centers $\boldsymbol{\mu}_1, \dots, \boldsymbol{\mu}_K$

2. While NOT CONVERGED

a. Assign each data point to the cluster with the nearest cluster center:

$$z^{(n)} = \underset{k}{\operatorname{argmin}} \|\mathbf{x}^{(n)} - \boldsymbol{\mu}_k\|_2$$

b. Recompute the cluster centers:

$$\boldsymbol{\mu}_k = \frac{1}{N_k} \sum_{n: z^{(n)} = k} \mathbf{x}^{(n)}$$

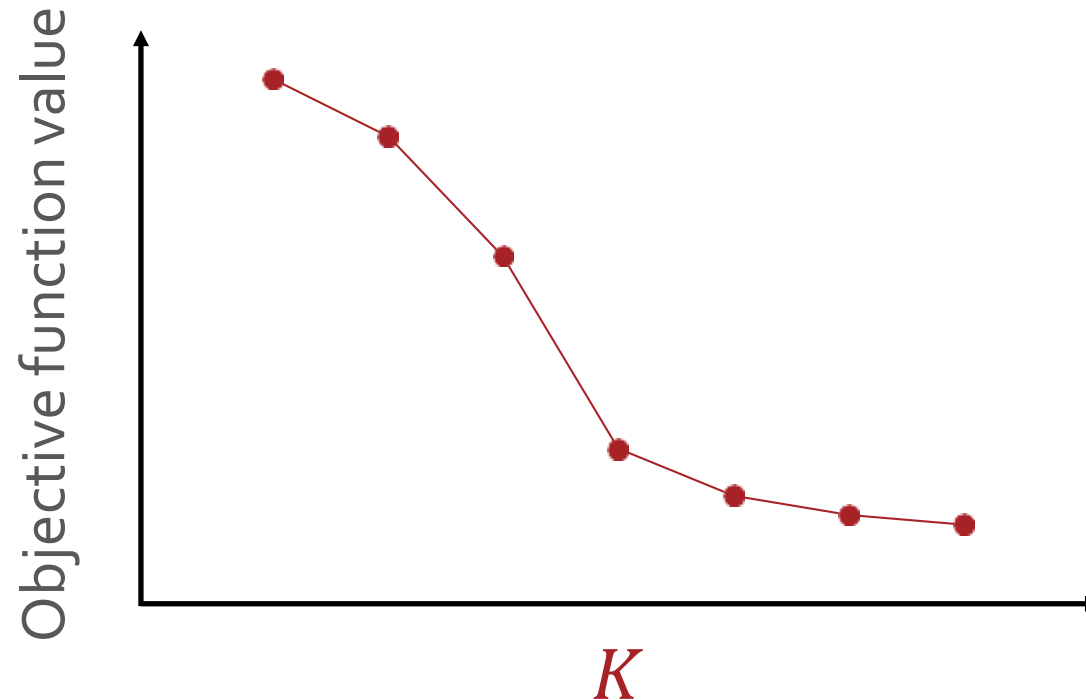
where N_k is the number of data points in cluster k

- Output: cluster centers $\boldsymbol{\mu}_1, \dots, \boldsymbol{\mu}_K$ and cluster assignments $z^{(1)}, \dots, z^{(N)}$

How do we set these hyperparameters?

Setting K

- Idea: choose the value of K that minimizes the objective function

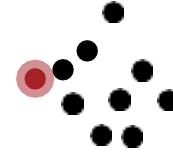
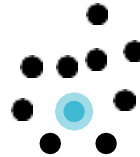
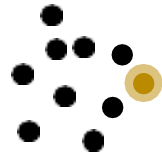


- Better Idea: look for the characteristic “elbow” or largest decrease when going from $K - 1$ to K



Initializing K -means: Lloyd's Method

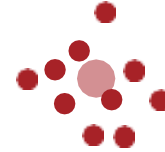
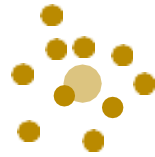
- Randomly choose K data points to be the initial cluster centers





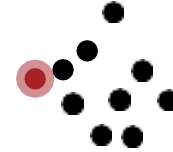
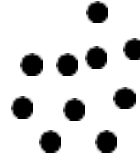
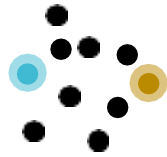
Initializing K -means: Lloyd's Method

- Randomly choose K data points to be the initial cluster centers



Initializing K -means: Lloyd's Method

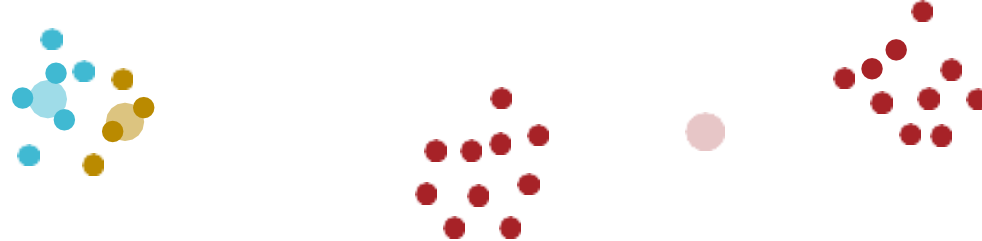
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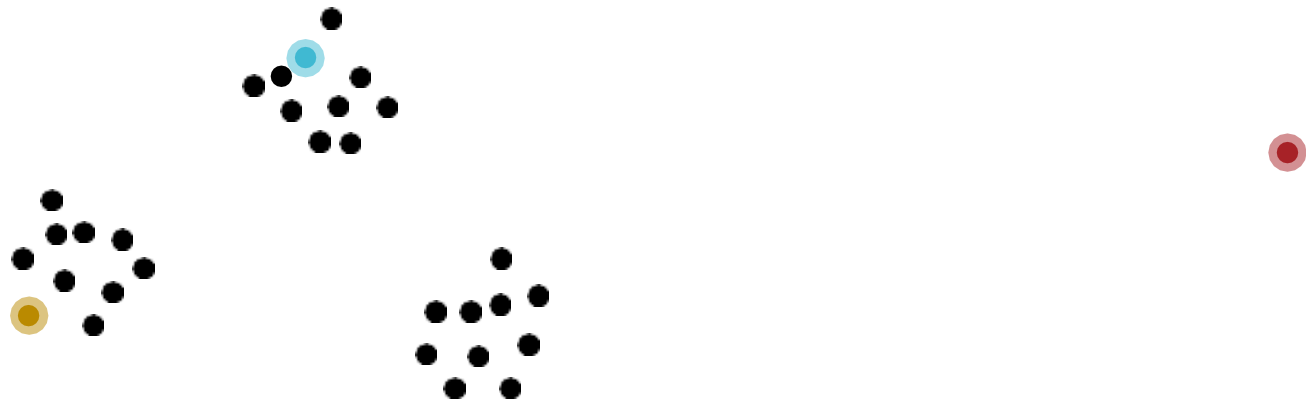
Initializing K -means: Lloyd's Method

- Randomly choose K data points to be the initial cluster centers



Initializing K -means: Furthest Point

1. Choose the first cluster center randomly from the data points
2. For each other data point x , compute $D(x)$, the distance between x and the closest cluster center to x
3. Select the data point with the largest $D(x)$ as the next cluster center
4. Repeat 2 and 3 $K - 1$ times



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Initializing K -means: Furthest Point

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 3. Select the data point with the largest $D(\mathbf{x})$ as the next cluster center
 4. Repeat 2 and 3 $K - 1$ times
- Works great in the case of well-clustered data!
 - Can struggle with outliers...

Initializing K -means: K -means++

1. Choose the first cluster center randomly from the data points
2. For each other data point \mathbf{x} , compute $D(\mathbf{x})$, the distance between \mathbf{x} and the closest cluster center to \mathbf{x}
3. *Sample* the next cluster center with probability proportional to $D(\mathbf{x})^2$
4. Repeat 2 and 3 $K - 1$ times

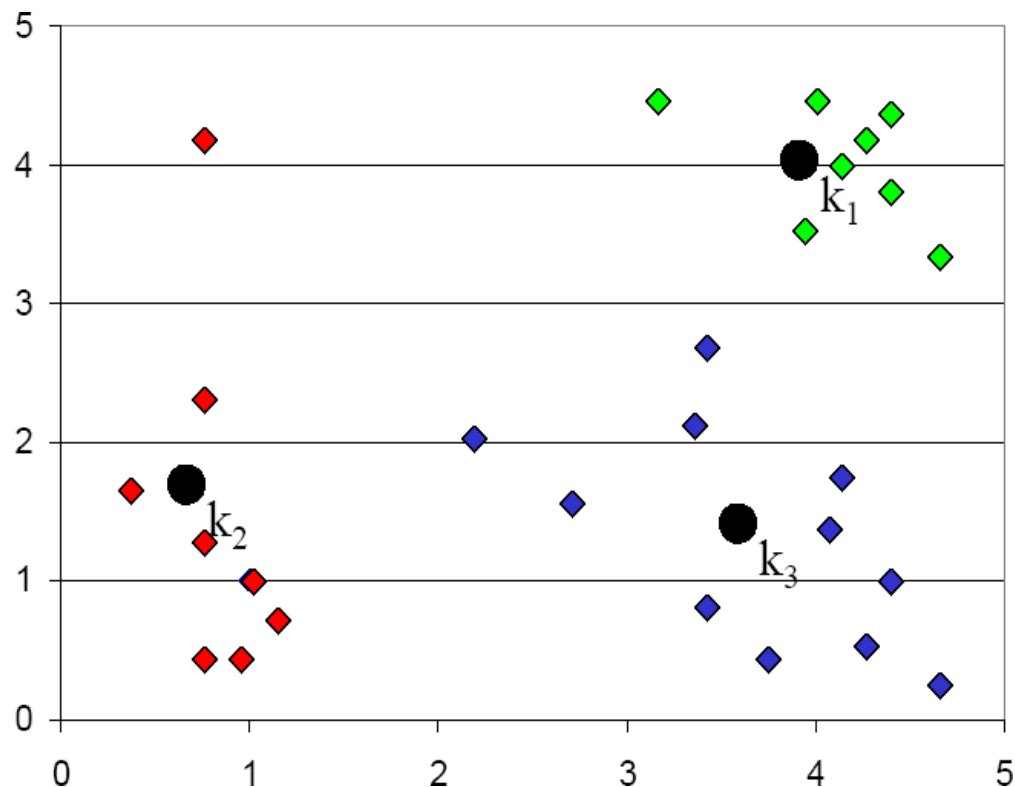
i	$D(\mathbf{x}^{(i)})$	$D(\mathbf{x}^{(i)})^2$	Probability of Being Selected
1	4	16	16/123
2	7	49	49/123
\vdots	\vdots	\vdots	\vdots
N	1	1	1/123
Total		123	123/123 = 1



Initializing K -means: K -means++

1. Choose the first cluster center randomly from the data points
 2. For each other data point \mathbf{x} , compute $D(\mathbf{x})$, the distance between \mathbf{x} and the closest cluster center to \mathbf{x}
 3. *Sample* the next cluster center with probability proportional to $D(\mathbf{x})^2$
 4. Repeat 2 and 3 $K - 1$ times
- K -means++ achieves a $O(\log K)$ approximation to the optimal clustering in expectation!
 - All initialization methods can benefit from multiple random restarts

Problem with K-means



	Cluster 1	Cluster 2	Cluster 3
Individual 1	1	0	0
Individual 2	0	1	0
Individual 3	0	1	0
Individual 4	1	0	0
Individual 5
Individual 6
Individual 7
Individual 8
Individual 9
Individual 10

Hard assignment of samples into three clusters

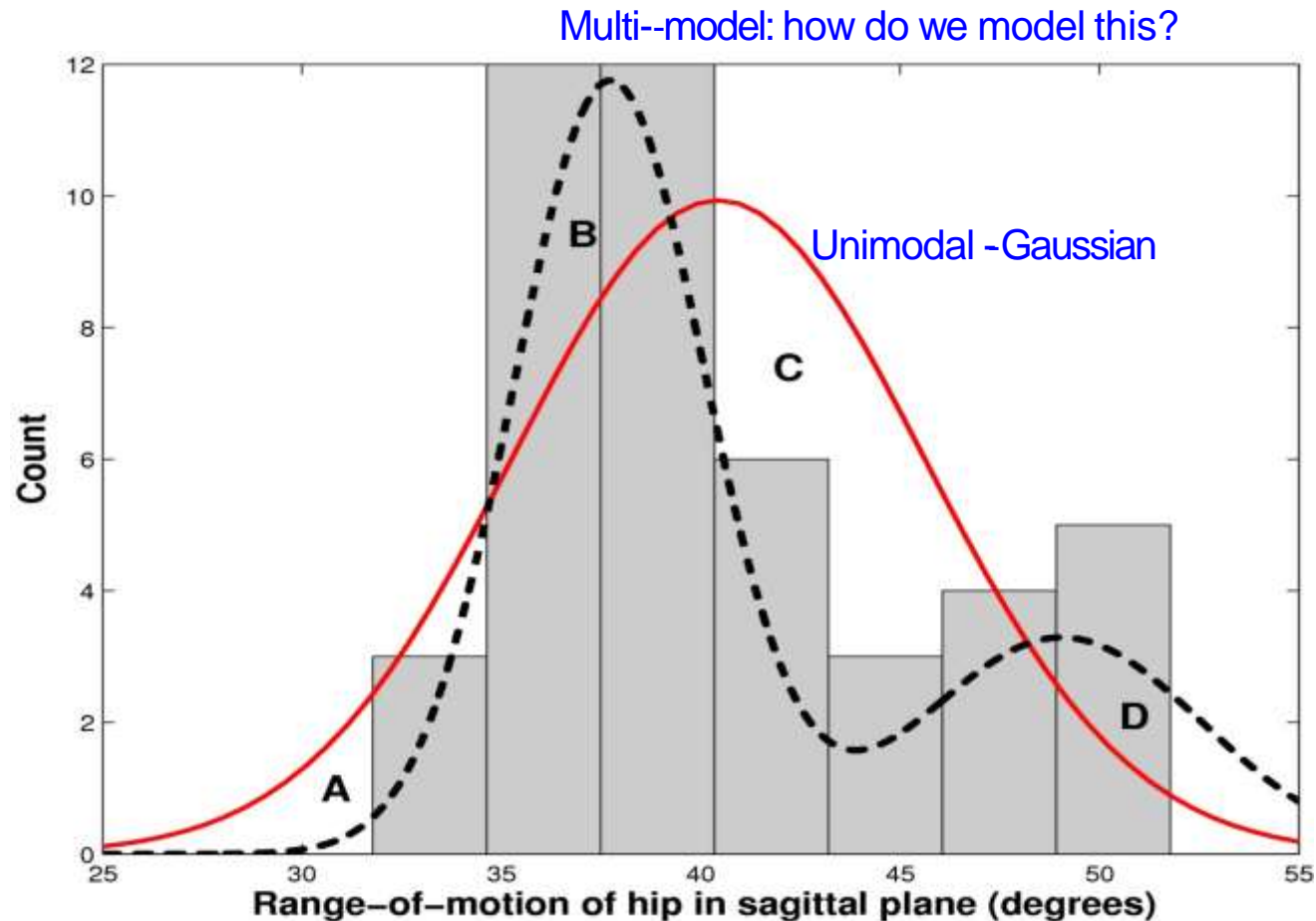
Probabilistic Soft-Clustering of Samples into Three Clusters

Probability of	Cluster 1	Cluster 2	Cluster 3	Sum
Individual 1	0.1	0.4	0.5	1
Individual 2	0.8	0.1	0.1	1
Individual 3	0.7	0.2	0.1	1
Individual 4	0.10	0.05	0.85	1
Individual 5	1
Individual 6	1
Individual 7	1
Individual 8	1
Individual 9	1
Individual 10	1

- Each sample can be assigned to more than one clusters with a certain probability.
- For each sample, the probabilities for all clusters should sum to 1. (i.e., each row should sum to 1.)
- Each cluster is explained by a cluster center variable (i.e., cluster mean)

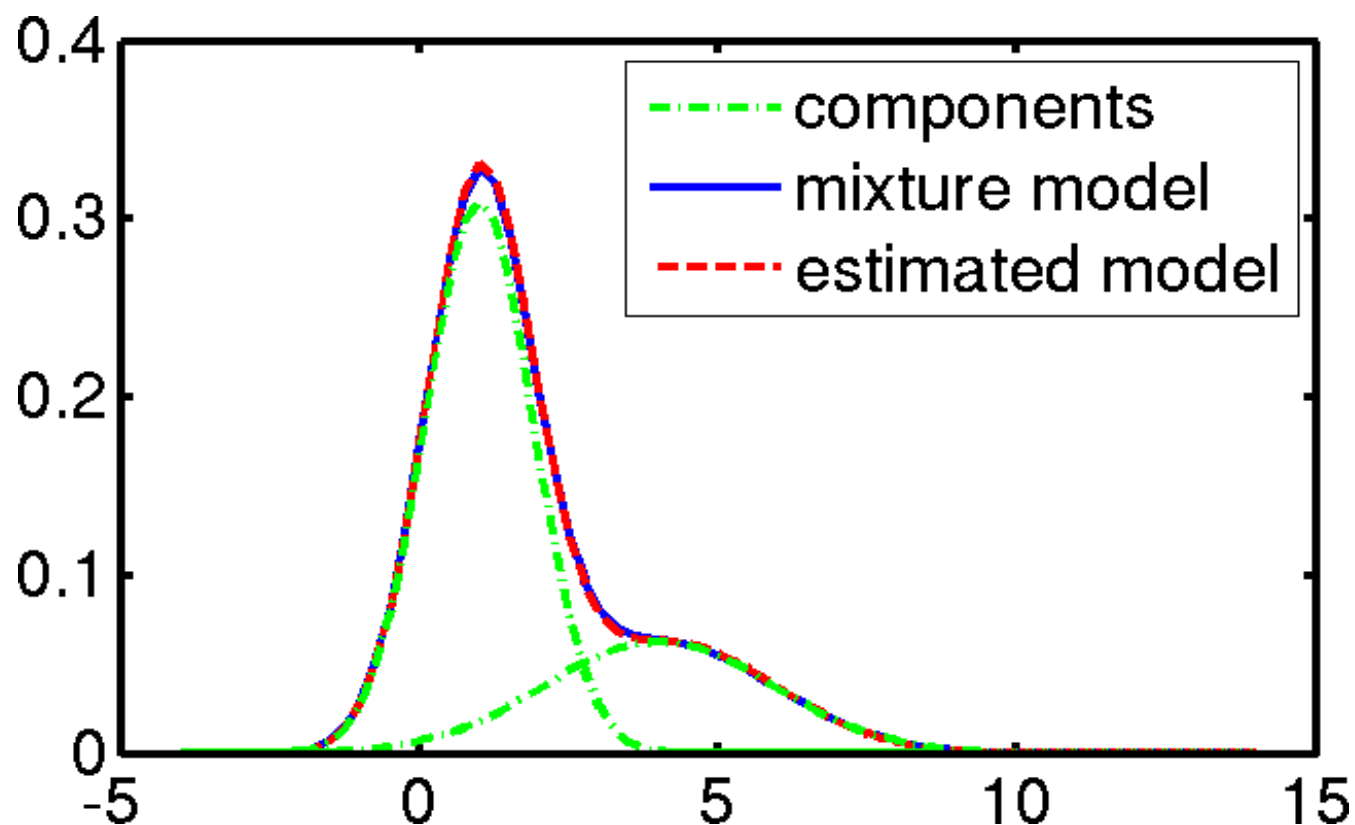
Mixture Model

- A density model $p(\mathbf{x})$ may be multi-modal.



Mixture Model

- We may be able to model it as a mixture of uni-modal distributions (e.g., Gaussians).
- Each mode may correspond to a different sub-population (e.g., male and female).



GMM

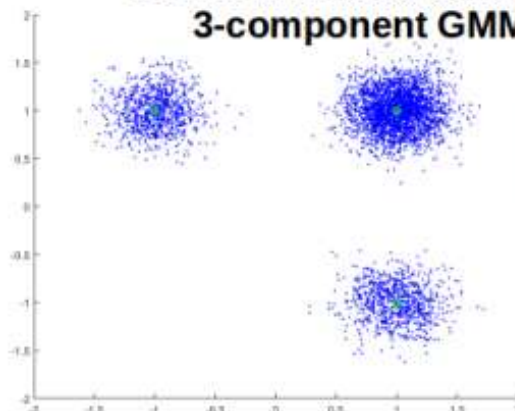
- The generative story for each \mathbf{x}_n , $n = 1, 2, \dots, N$
 - First choose one of the K mixture components as

$$\mathbf{z}_n \sim \text{Multinomial}(\mathbf{z}_n | \boldsymbol{\pi}) \quad (\text{from the prior } p(\mathbf{z}) \text{ over } \mathbf{z})$$

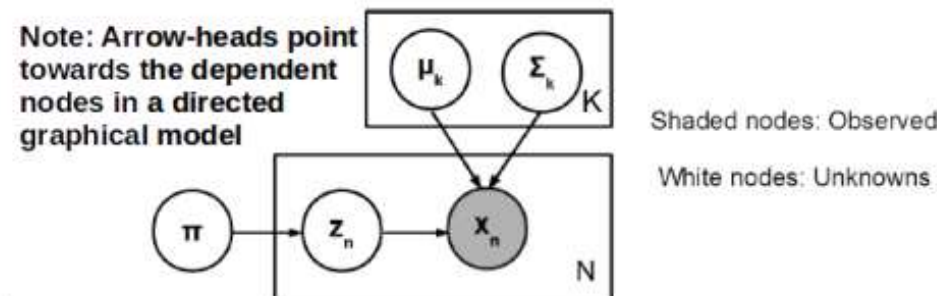
- Suppose $\mathbf{z}_n = k$. Now generate \mathbf{x}_n from the k -th Gaussian as

$$\mathbf{x}_n \sim \mathcal{N}(\mathbf{x}_n | \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k) \quad (\text{from the data distr. } p(\mathbf{x} | \mathbf{z}))$$

Some simulated data from a
3-component GMM



Directed Graphical Model
for a K-component GMM



Parameter Estimation with Latent Variables

Recall...

- Consider the 'incomplete' data log likelihood

$$\begin{aligned}\log p(\mathbf{X}|\Theta) &= \log \sum_{\mathbf{Z}} p(\mathbf{X}, \mathbf{Z}|\Theta) = \log \sum_{\mathbf{Z}} q(\mathbf{Z}) \frac{p(\mathbf{X}, \mathbf{Z}|\Theta)}{q(\mathbf{Z})} \quad (\text{where } q(\mathbf{Z}) \text{ is some dist.}) \\ &\geq \sum_{\mathbf{Z}} q(\mathbf{Z}) \log \frac{p(\mathbf{X}, \mathbf{Z}|\Theta)}{q(\mathbf{Z})} \quad (\text{concave } f, \text{ Jensen's Ineq.: } f(\sum \lambda_i x_i) \geq \sum \lambda_i f(x_i)) \\ \log p(\mathbf{X}|\Theta) &\geq \sum_{\mathbf{Z}} q(\mathbf{Z}) \log p(\mathbf{X}, \mathbf{Z}|\Theta) - \underbrace{\sum_{\mathbf{Z}} q(\mathbf{Z}) \log q(\mathbf{Z})}_{\text{doesn't depend on } \Theta} = \sum_{\mathbf{Z}} q(\mathbf{Z}) \log p(\mathbf{X}, \mathbf{Z}|\Theta) + \text{const.}\end{aligned}$$

- If we set $q(\mathbf{Z}) = p(\mathbf{Z}|\mathbf{X}, \Theta)$, the above inequality becomes equality

$$\begin{aligned}\sum_{\mathbf{Z}} q(\mathbf{Z}) \log \frac{p(\mathbf{X}, \mathbf{Z}|\Theta)}{q(\mathbf{Z})} &= \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta) \log \frac{\cancel{p(\mathbf{Z}|\mathbf{X}, \Theta)} p(\mathbf{X}|\Theta)}{\cancel{p(\mathbf{Z}|\mathbf{X}, \Theta)}} = \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta) \log p(\mathbf{X}|\Theta) \\ &= \log p(\mathbf{X}|\Theta) \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta) = \log p(\mathbf{X}|\Theta)\end{aligned}$$

- Thus for $q(\mathbf{Z}) = p(\mathbf{Z}|\mathbf{X}, \Theta)$, we have

$$\log p(\mathbf{X}|\Theta) = \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta) \log p(\mathbf{X}, \mathbf{Z}|\Theta) + \text{const.} = \mathbb{E}[\log p(\mathbf{X}, \mathbf{Z}|\Theta)] + \text{const.}$$

- Thus $\log p(\mathbf{X}|\Theta)$ is tightly lower-bounded by $\mathbb{E}[\log p(\mathbf{X}, \mathbf{Z}|\Theta)]$ which EM maximizes

The Expectation Maximization (EM) Algorithm

Recall...

Initialize the parameters: Θ^{old} . Then alternate between these steps:

- **E (Expectation) step:**

- Compute the posterior $p(\mathbf{Z}|\mathbf{X}, \Theta^{old})$ over latent variables \mathbf{Z} using Θ^{old}
- Compute the expected complete data log-likelihood w.r.t. *this* posterior

$$Q(\Theta, \Theta^{old}) = \mathbb{E}_{p(\mathbf{Z}|\mathbf{X}, \Theta^{old})} [\log p(\mathbf{X}, \mathbf{Z}|\Theta)] = \sum_{\mathbf{Z}} p(\mathbf{Z}|\mathbf{X}, \Theta^{old}) \log p(\mathbf{X}, \mathbf{Z}|\Theta)$$

- **M (Maximization) step:**

- Maximize the expected complete data log-likelihood w.r.t. Θ

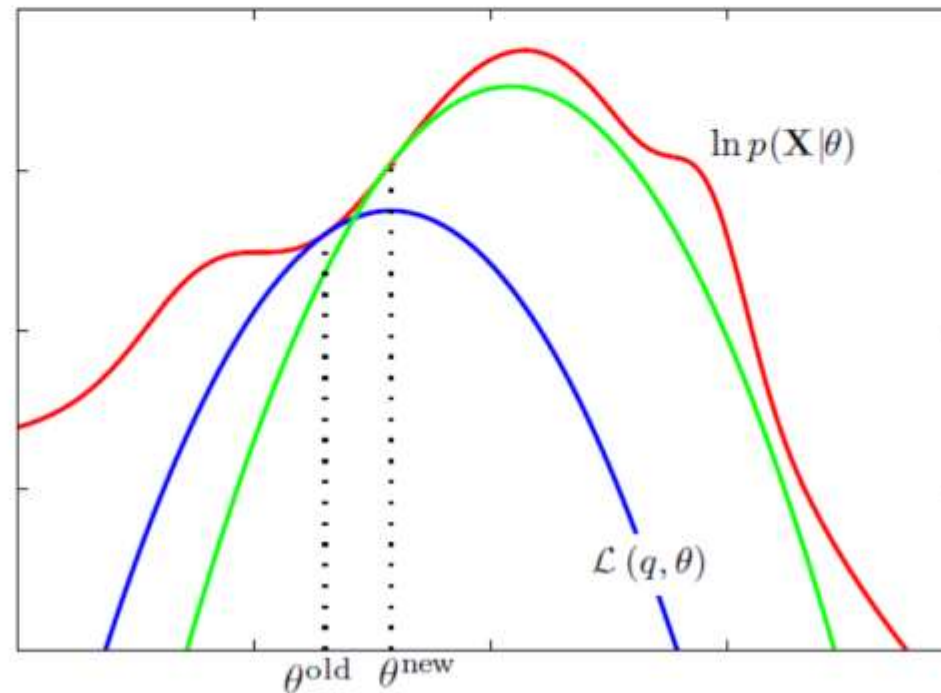
$$\Theta^{new} = \arg \max_{\Theta} Q(\Theta, \Theta^{old}) \quad (\text{if doing MLE})$$

$$\Theta^{new} = \arg \max_{\Theta} \{Q(\Theta, \Theta^{old}) + \log p(\Theta)\} \quad (\text{if doing MAP})$$

- If the log-likelihood or the parameter values not converged then set $\Theta^{old} = \Theta^{new}$ and go to the E step.

EM: A View in the Parameter Space

- E-step: Update of q makes the $\mathcal{L}(q, \Theta)$ curve touch the $\log p(\mathbf{X}|\Theta)$ curve
- M-step gives the maxima Θ^{new} of $\mathcal{L}(q, \Theta)$
- Next E-step readjusts $\mathcal{L}(q, \Theta)$ curve (green) to meet $\log p(\mathbf{X}|\Theta)$ curve again
- This continues until a local maxima of $\log p(\mathbf{X}|\Theta)$ is reached



Learning GMM by EM

1. Initialize the Parameters $\Theta = \{\pi_k, \mu_k, \Sigma_k\}_{k=1}^K$ randomly, or using K-means
2. Iterate until convergence (e.g., when $\log p(X|\Theta)$ ceases to increase

a) E-step:

$$\gamma_{ik} = p(z_i = k | x_i, \Theta^{old}) = \frac{\pi_k^{old} \mathcal{N}(x_i | \mu_k^{old}, \Sigma_k^{old})}{\sum_{j=1}^K \pi_j^{old} \mathcal{N}(x_i | \mu_j^{old}, \Sigma_j^{old})}$$

b) M-step:

$$\pi_k^{new} = \frac{1}{N} \sum_{i=1}^N \gamma_{ik}$$

$$\mu_k^{new} = \frac{\sum_{i=1}^N \gamma_{ik} x_i}{\sum_{i=1}^N \gamma_{ik}}$$

$$\Sigma_k^{new} = \frac{\sum_{i=1}^N \gamma_{ik} (x_i - \mu_k^{new})(x_i - \mu_k^{new})^T}{\sum_{i=1}^N \gamma_{ik}}$$



DIMENSIONALITY REDUCTION

High Dimension Data

Examples of high dimensional data:

- High resolution images (millions of pixels)



High Dimension Data

Examples of high dimensional data:

- Multilingual News Stories
(vocabulary of hundreds of thousands of words)



High Dimension Data

Examples of high dimensional data:

- Brain Imaging Data (100s of MBs per scan)

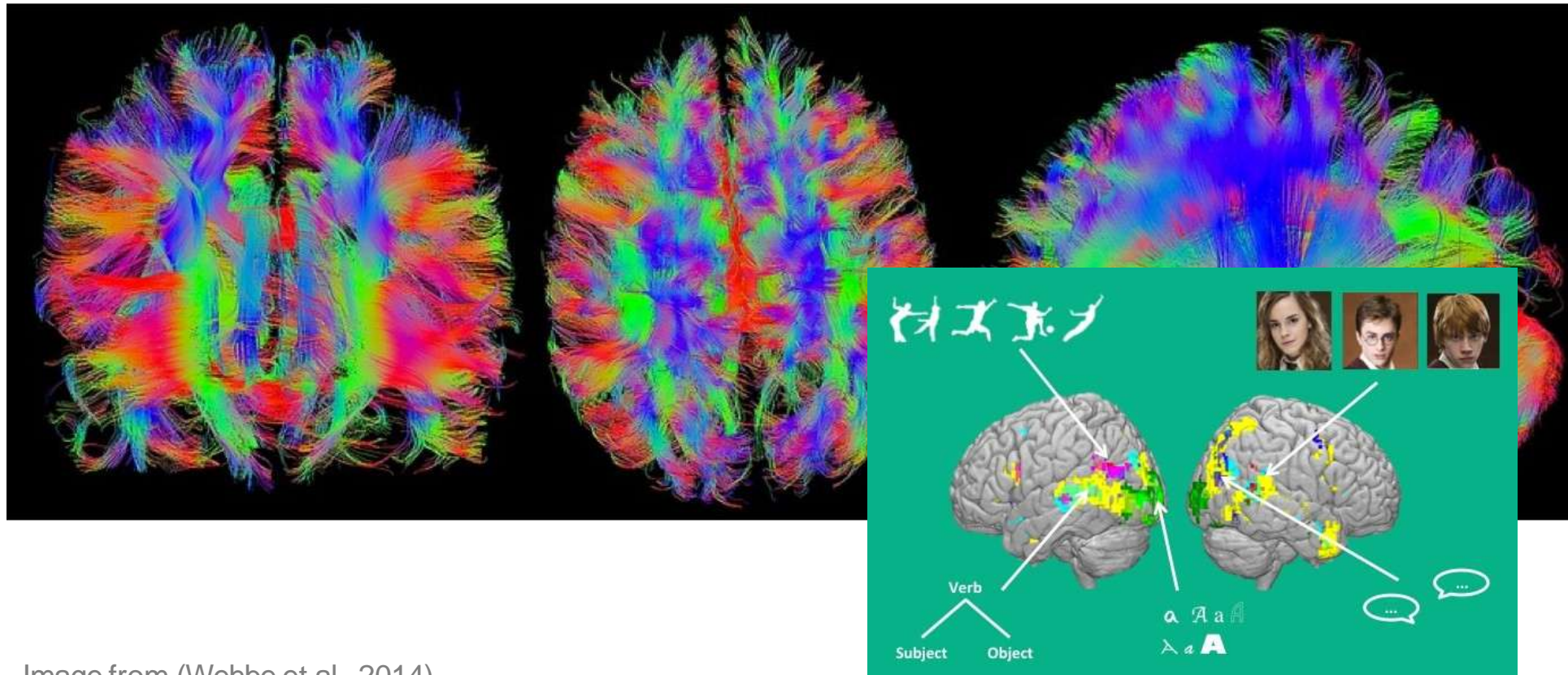


Image from (Wehbe et al., 2014)

Image from <https://pixabay.com/en/brain-mrt-magnetic-resonance-imaging-1728449/>



Dimensionality Reduction Algorithms:

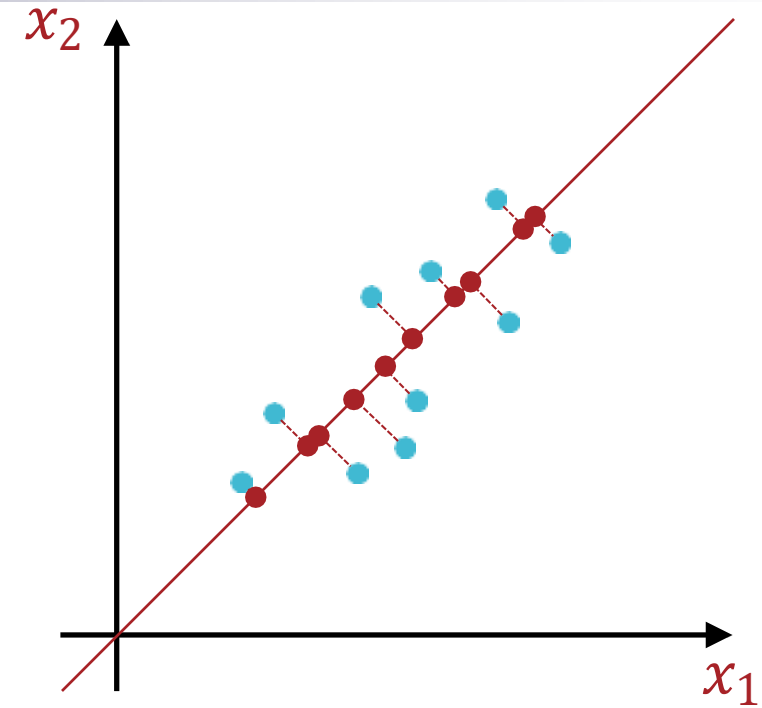
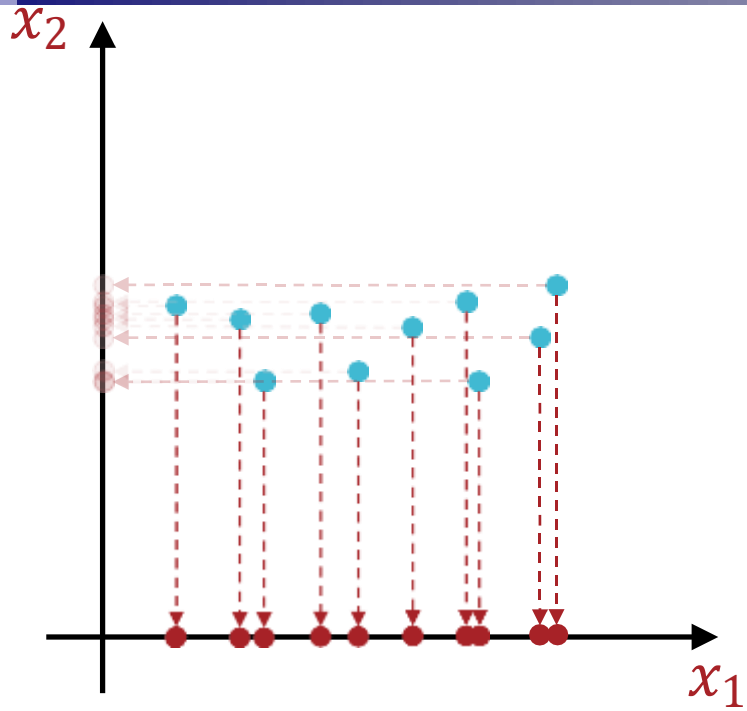
Powerful (often unsupervised) learning techniques for extracting hidden (potentially lower dimensional) structure from high dimensional datasets.

Examples:

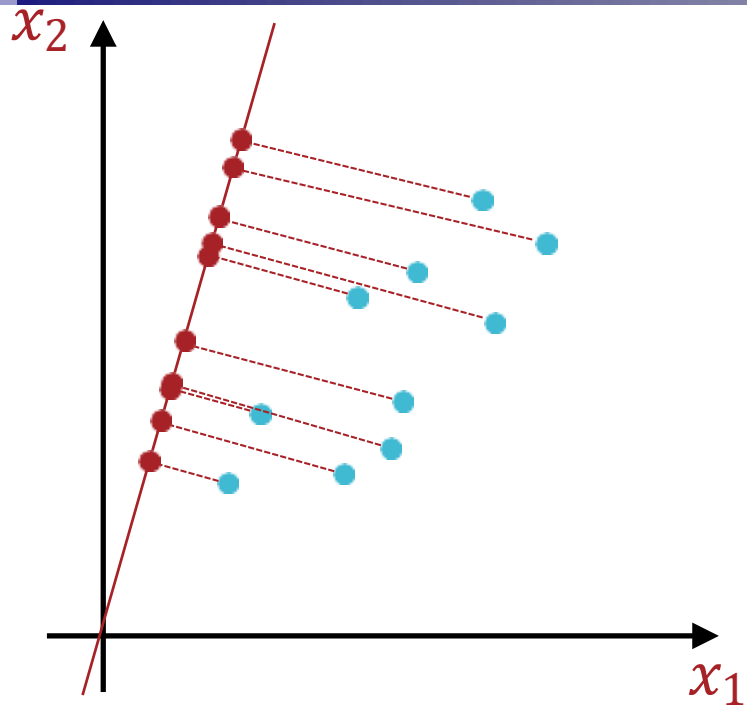
PCA, Kernel PCA, ICA, CCA, t-SNE, Autoencoders, VAEs

Useful for:

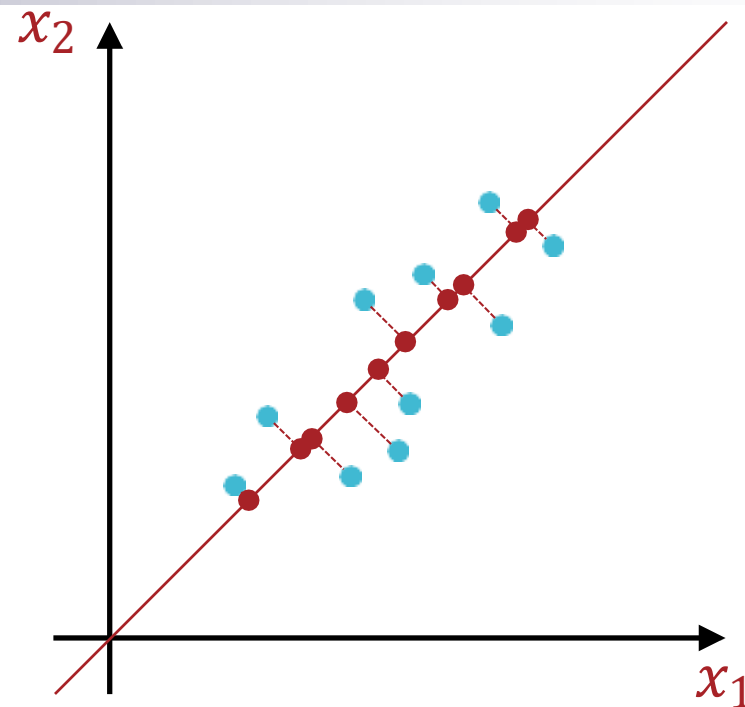
- Visualization
- More efficient use of resources (e.g., time, memory, communication)
- Statistical: fewer dimensions \rightarrow better generalization
- Noise removal (improving data quality)



Feature Elimination \in Dimensionality Reduction



Option A



Option C

Option B (TOXIC)

Which projection do you prefer (Q1) and why (Q2)?

Background: Sample Variance and Covariance

- Given a collection of N 1-dimensional samples $[x^{(1)}, x^{(2)}, \dots, x^{(N)}]$ from some random variable, the **sample variance** is

$$\hat{\sigma}^2 = \frac{1}{N} \sum_{i=1}^N (x^{(i)} - \hat{\mu})^2 = \frac{1}{N} \sum_{i=1}^N \left(x^{(i)} - \frac{1}{N} \sum_{n=1}^N x^{(n)} \right)^2$$

- Given a collection of N D -dimensional samples $[x^{(1)}, x^{(2)}, \dots, x^{(N)}]$ from some random variable, the **sample covariance** between dimension j and k is

$$\Sigma_{jk} = \frac{1}{N} \sum_{i=1}^N (x_j^{(i)} - \hat{\mu}_j) (x_k^{(i)} - \hat{\mu}_k) \text{ where } \hat{\mu}_d = \frac{1}{N} \sum_{n=1}^N x_d^{(n)}$$

Background: Sample Variance and Covariance

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- Given a collection of N D -dimensional samples $[\mathbf{x}^{(1)}, \mathbf{x}^{(2)}, \dots, \mathbf{x}^{(N)}]$ from some random variable, the **sample covariance matrix** is

$$\Sigma = \frac{1}{N} X^T X \quad \text{where} \quad X = \begin{bmatrix} (\mathbf{x}^{(1)} - \boldsymbol{\mu})^T \\ (\mathbf{x}^{(2)} - \boldsymbol{\mu})^T \\ \vdots \\ (\mathbf{x}^{(N)} - \boldsymbol{\mu})^T \end{bmatrix}$$

Centering the Data



- To be consistent, we will constrain principal components to be *orthogonal unit vectors* that begin at the origin
- Preprocess data to be centered around the origin:

$$1. \mu = \frac{1}{N} \sum_{n=1}^N \mathbf{x}^{(n)}$$

$$2. \tilde{\mathbf{x}}^{(n)} = \mathbf{x}^{(n)} - \mu \quad \forall n$$

$$3. X = \begin{bmatrix} \tilde{\mathbf{x}}^{(1)T} \\ \tilde{\mathbf{x}}^{(2)T} \\ \vdots \\ \tilde{\mathbf{x}}^{(N)T} \end{bmatrix}$$

Reconstruction Error

- The projection of $\tilde{\mathbf{x}}^{(n)}$ onto a vector \mathbf{v} is

$$\mathbf{z}^{(n)} = \left(\frac{\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}}{\|\mathbf{v}\|_2} \right) \frac{\mathbf{v}}{\|\mathbf{v}\|_2}$$

Length of projection

Direction of projection

Reconstruction Error

- The projection of $\tilde{\mathbf{x}}^{(n)}$ onto a unit vector \mathbf{v} is

$$\mathbf{z}^{(n)} = (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v}$$

$$\hat{\mathbf{v}} = \operatorname{argmin}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \sum_{n=1}^N \|\tilde{\mathbf{x}}^{(n)} - (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v}\|_2^2$$

$$\|\tilde{\mathbf{x}}^{(n)} - (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v}\|_2^2$$

$$= \tilde{\mathbf{x}}^{(n)T} \tilde{\mathbf{x}}^{(n)} - 2(\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v}^T \tilde{\mathbf{x}}^{(n)} + (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v}^T \mathbf{v}$$

$$= \tilde{\mathbf{x}}^{(n)T} \tilde{\mathbf{x}}^{(n)} - (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v}^T \tilde{\mathbf{x}}^{(n)}$$

$$= \|\tilde{\mathbf{x}}^{(n)}\|_2^2 - (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)})^2$$

Minimizing the
Reconstruction
Error



Maximizing the
Variance

$$\hat{\mathbf{v}} = \operatorname{argmin}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \sum_{n=1}^N \left\| \tilde{\mathbf{x}}^{(n)} - (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)}) \mathbf{v} \right\|_2^2$$

$$= \operatorname{argmin}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \sum_{n=1}^N \left\| \tilde{\mathbf{x}}^{(n)} \right\|_2^2 - (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)})^2$$

$$= \operatorname{argmax}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \sum_{n=1}^N (\mathbf{v}^T \tilde{\mathbf{x}}^{(n)})^2 \quad \leftarrow \text{Variance of projections } (\tilde{\mathbf{x}}^{(n)} \text{ are centered})$$

$$= \operatorname{argmax}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \mathbf{v}^T \left(\sum_{n=1}^N \tilde{\mathbf{x}}^{(n)} \tilde{\mathbf{x}}^{(n)T} \right) \mathbf{v}$$

$$= \operatorname{argmax}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \mathbf{v}^T (X^T X) \mathbf{v}$$

Maximizing the Variance

$$\hat{\mathbf{v}} = \operatorname{argmax}_{\mathbf{v}: \|\mathbf{v}\|_2^2 = 1} \mathbf{v}^T (X^T X) \mathbf{v}$$

$$\begin{aligned} \mathcal{L}(\mathbf{v}, \lambda) &= \mathbf{v}^T (X^T X) \mathbf{v} - \lambda (\|\mathbf{v}\|_2^2 - 1) \\ &= \mathbf{v}^T (X^T X) \mathbf{v} - \lambda (\mathbf{v}^T \mathbf{v} - 1) \end{aligned}$$

$$\frac{\partial \mathcal{L}}{\partial \mathbf{v}} = 2(X^T X) \mathbf{v} - 2\lambda \mathbf{v}$$

$$\rightarrow 2(X^T X) \hat{\mathbf{v}} - 2\lambda \hat{\mathbf{v}} = 0 \rightarrow (X^T X) \hat{\mathbf{v}} = \lambda \hat{\mathbf{v}}$$

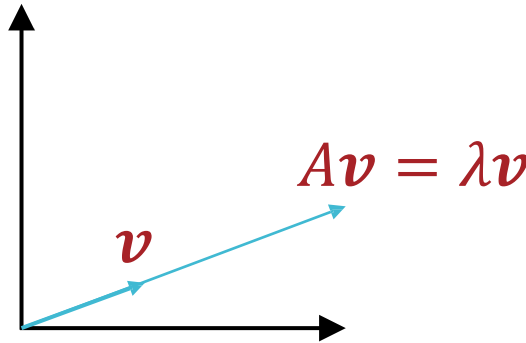
- $\hat{\mathbf{v}}$ is an eigenvector of $X^T X$ and λ is the corresponding eigenvalue!

Background: Eigenvectors & Eigenvalues

- Given a square matrix $A \in \mathbb{R}^{N \times N}$, a vector $v \in \mathbb{R}^{N \times 1}$ is an

eigenvector of A iff there exists some scalar λ such that

$$Av = \lambda v$$



Intuition: A scales or stretches v but does not rotate it

- Key property: the eigenvectors of symmetric matrices (e.g., the covariance matrix of a data set) are orthogonal!

Maximizing the Variance

$$\hat{\mathbf{v}} = \operatorname{argmax}_{\mathbf{v}: \|\mathbf{v}\|_2^2 = 1} \mathbf{v}^T (X^T X) \mathbf{v}$$

$$\begin{aligned} \mathcal{L}(\mathbf{v}, \lambda) &= \mathbf{v}^T (X^T X) \mathbf{v} - \lambda (\|\mathbf{v}\|_2^2 - 1) \\ &= \mathbf{v}^T (X^T X) \mathbf{v} - \lambda (\mathbf{v}^T \mathbf{v} - 1) \end{aligned}$$

$$\frac{\partial \mathcal{L}}{\partial \mathbf{v}} = 2(X^T X) \mathbf{v} - 2\lambda \mathbf{v}$$

$$\rightarrow 2(X^T X) \hat{\mathbf{v}} - 2\lambda \hat{\mathbf{v}} = 0 \rightarrow (X^T X) \hat{\mathbf{v}} = \lambda \hat{\mathbf{v}}$$

- $\hat{\mathbf{v}}$ is an eigenvector of $X^T X$ and λ is the corresponding eigenvalue!
- But which one?

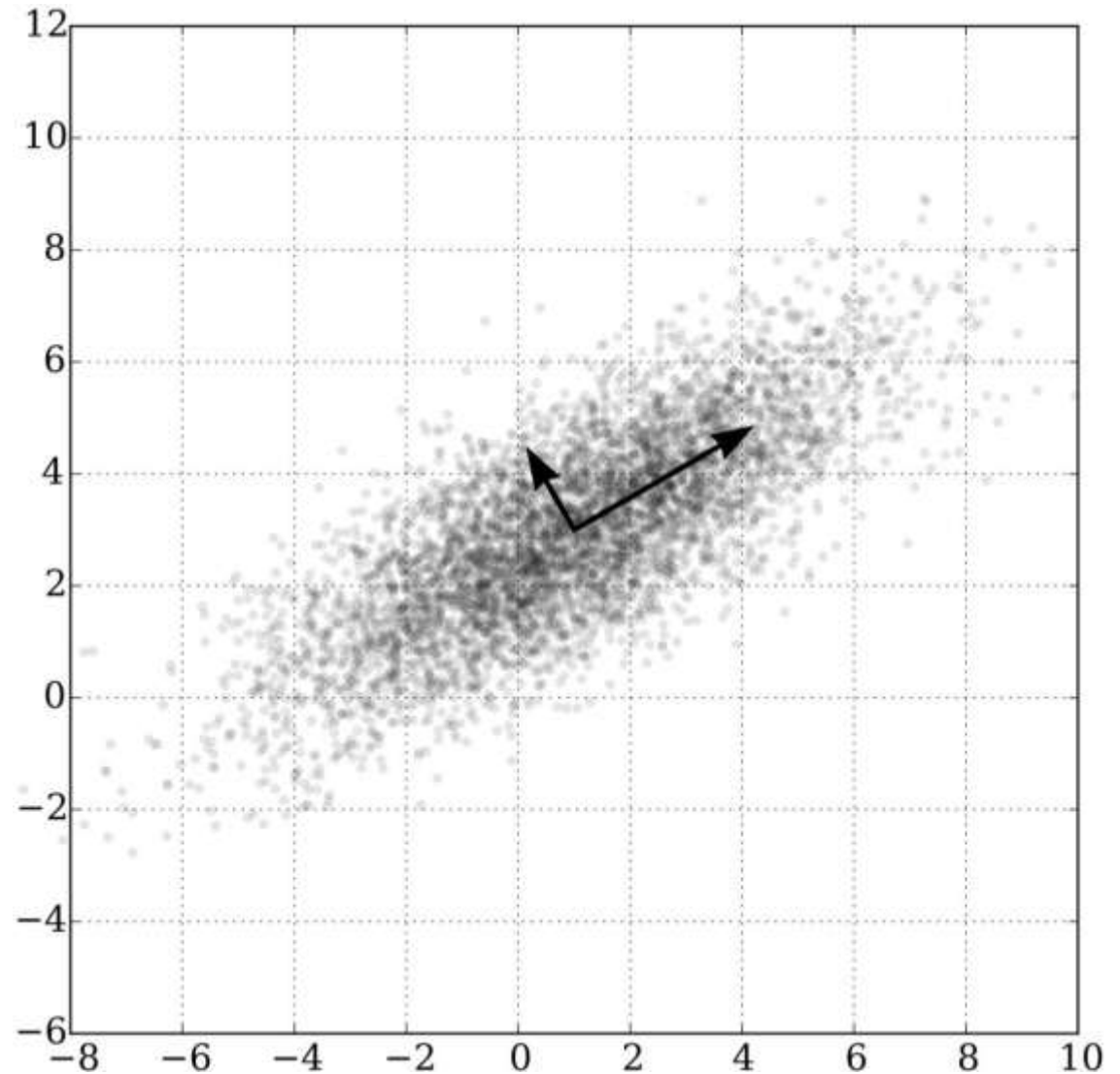
Maximizing the Variance

$$\hat{\mathbf{v}} = \operatorname{argmax}_{\mathbf{v}: \|\mathbf{v}\|_2^2=1} \mathbf{v}^T (X^T X) \mathbf{v}$$

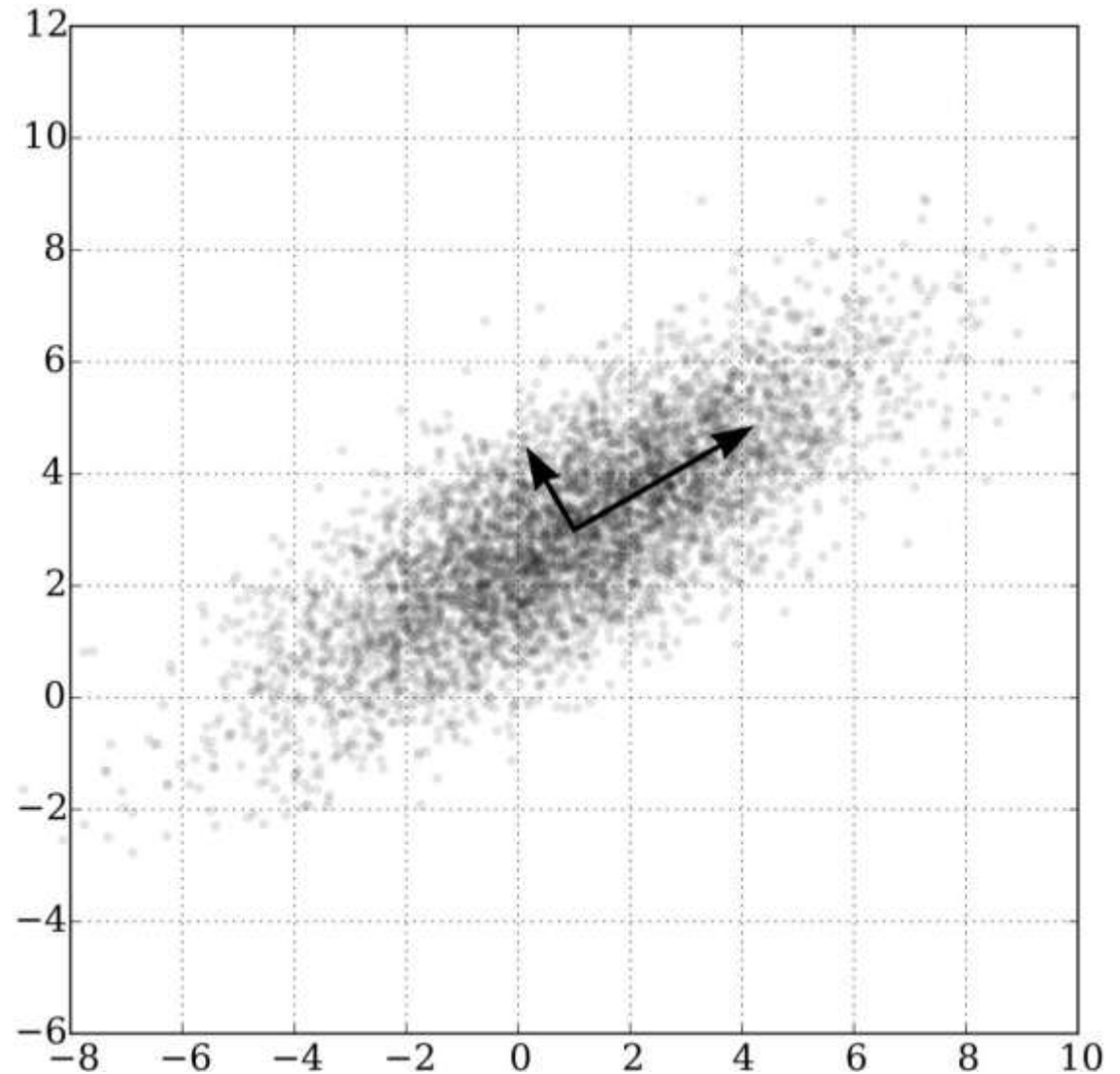
$$(X^T X) \hat{\mathbf{v}} = \lambda \hat{\mathbf{v}} \rightarrow \hat{\mathbf{v}}^T (X^T X) \hat{\mathbf{v}} = \lambda \hat{\mathbf{v}}^T \hat{\mathbf{v}} = \lambda$$

- The first principal component is the eigenvector $\hat{\mathbf{v}}_1$ that corresponds to the largest eigenvalue λ_1
- The second principal component is the eigenvector $\hat{\mathbf{v}}_2$ that corresponds to the second largest eigenvalue λ_2
 - $\hat{\mathbf{v}}_1$ and $\hat{\mathbf{v}}_2$ are orthogonal
- Etc ...
- λ_i is a measure of how much variance falls along $\hat{\mathbf{v}}_i$

Principal Components: Example



How can we
efficiently find
principal
components
(eigenvectors)?



Singular Value Decomposition (SVD) for PCA



- Every real-valued matrix $X \in \mathbb{R}^{N \times D}$ can be expressed as

$$X = USV^T$$

where:

1. $U \in \mathbb{R}^{N \times N}$ - columns of U are eigenvectors of XX^T
2. $V \in \mathbb{R}^{D \times D}$ - columns of V are eigenvectors of $X^T X$
3. $S \in \mathbb{R}^{N \times D}$ - diagonal matrix whose entries are the eigenvalues of $X \rightarrow$ squared entries are the eigenvalues of XX^T and $X^T X$

PCA Algorithm



- Input: $\mathcal{D} = \{(\mathbf{x}^{(n)})\}_{n=1}^N, \rho$
 1. Center the data
 2. Use SVD to compute the eigenvalues and eigenvectors of $X^T X$
 3. Collect the top ρ eigenvectors (corresponding to the ρ largest eigenvalues), $V_\rho \in \mathbb{R}^{D \times \rho}$
 4. Project the data into the space defined by V_ρ , $Z = X V_\rho$
- Output: Z , the transformed (potentially lower-dimensional) data

How many PCs
should we use?



- Input: $\mathcal{D} = \{(\mathbf{x}^{(n)})\}_{n=1}^N, \rho$
 1. Center the data
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Choosing the number of PCs



- Define a percentage of explained variance for the i^{th} PC:

$$\lambda_i / \sum \lambda_j$$

- Select all PCs above some threshold of explained variance, e.g., 5%
- Keep selecting PCs until the total explained variance exceeds some threshold, e.g., 90%
- Evaluate on some downstream metric

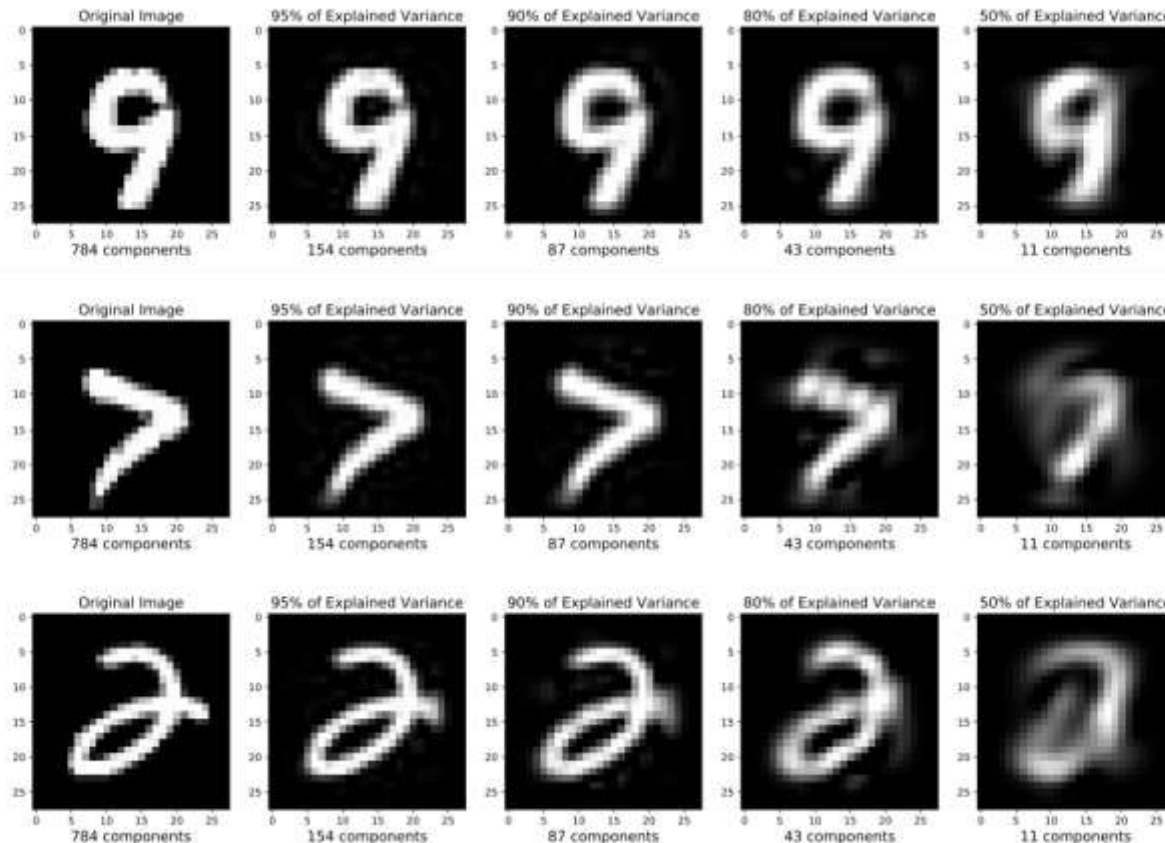


PCA EXAMPLES

Projecting MNIST digits

Task Setting:

1. Take each 28x28 image of a digit (i.e. a vector $\mathbf{x}^{(i)}$ of length 784) and project it down to K components (i.e. a vector $\mathbf{u}^{(i)}$)
2. Report percent of variance explained for K components
3. Then project back up to 28x28 image (i.e. a vector $\tilde{\mathbf{x}}^{(i)}$ of length 784) to visualize how much information was preserved



Takeaway:
Using fewer principal components K leads to higher reconstruction error.

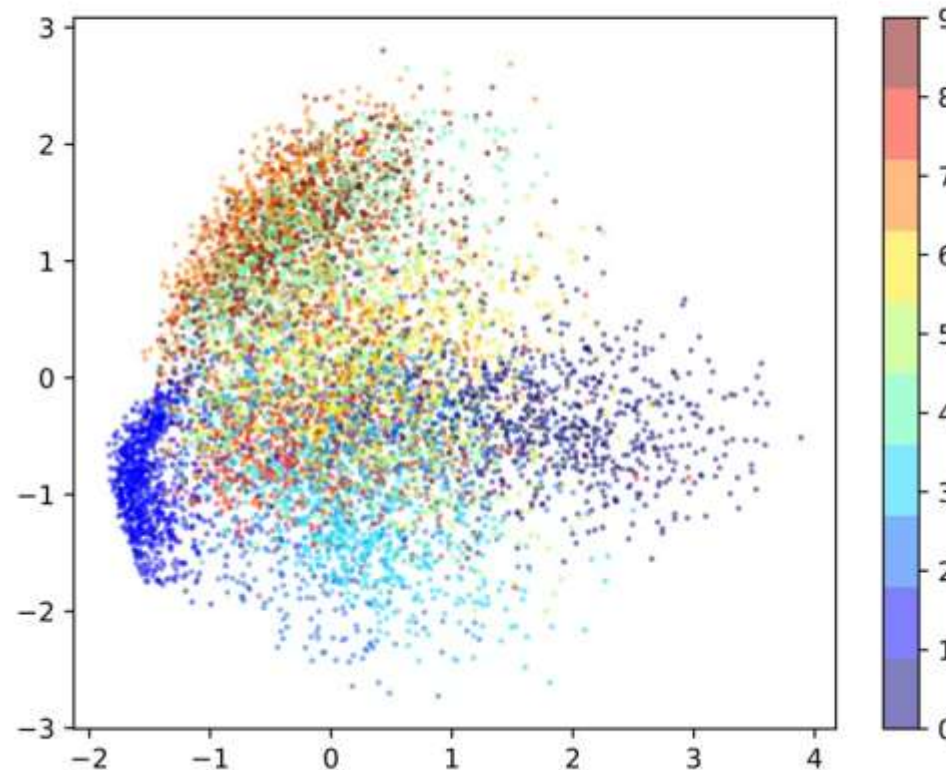
But even a small number (say 43) still preserves a lot of information about the original image.

Projecting MNIST digits



Task Setting:

1. Take each 28x28 image of a digit (i.e. a vector $\mathbf{x}^{(i)}$ of length 784) and project it down to $K=2$ components (i.e. a vector $\mathbf{u}^{(i)}$)
2. Plot the 2 dimensional points $\mathbf{u}^{(i)}$ and label with the (unknown to PCA) label $y^{(i)}$ as the color
3. Here we look at all ten digits 0 - 9



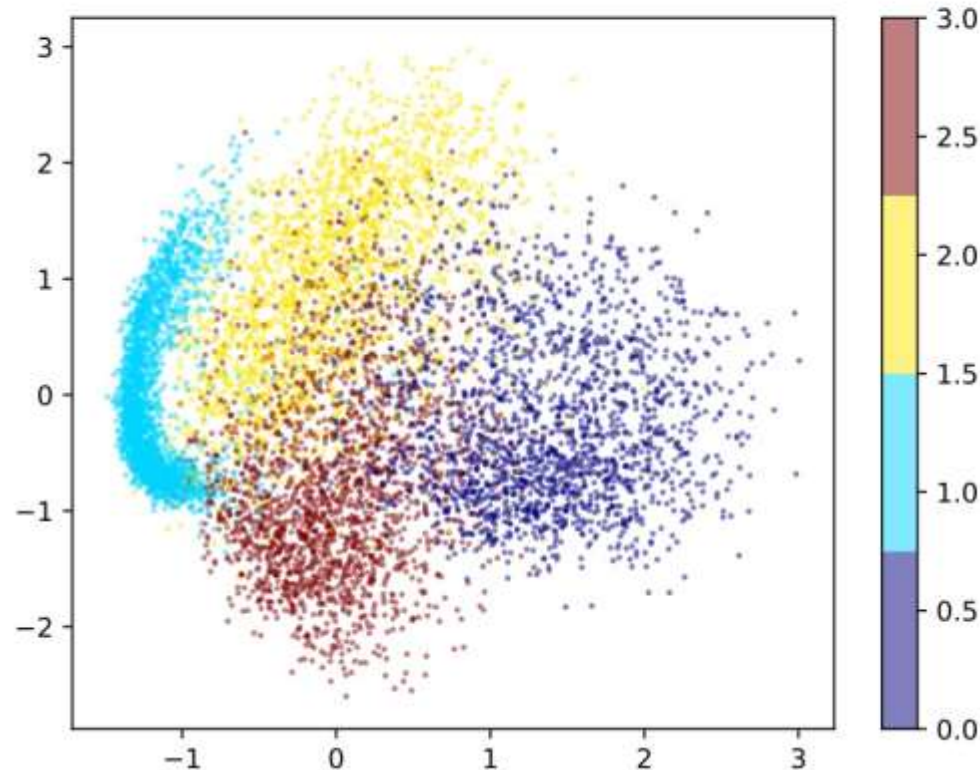
Takeaway:
Even with a tiny number of principal components $K=2$, PCA learns a representation that captures the *latent* information about the type of digit

Projecting MNIST digits



Task Setting:

1. Take each 28x28 image of a digit (i.e. a vector $\mathbf{x}^{(i)}$ of length 784) and project it down to $K=2$ components (i.e. a vector $\mathbf{u}^{(i)}$)
2. Plot the 2 dimensional points $\mathbf{u}^{(i)}$ and label with the (unknown to PCA) label $y^{(i)}$ as the color
3. Here we look at just four digits 0, 1, 2, 3



Takeaway:
Even with a tiny number of principal components $K=2$, PCA learns a representation that captures the *latent* information about the type of digit

Learning Objectives

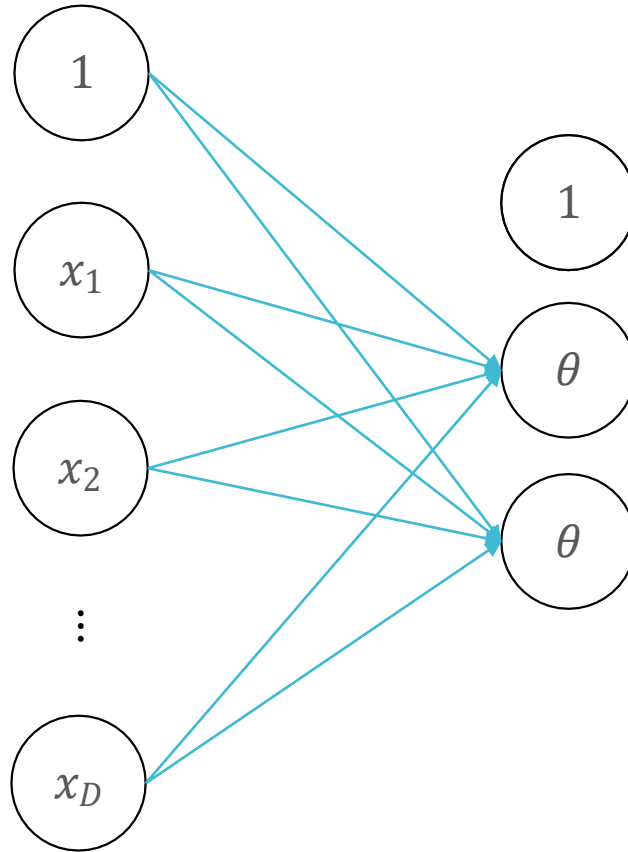


Dimensionality Reduction / PCA

You should be able to...

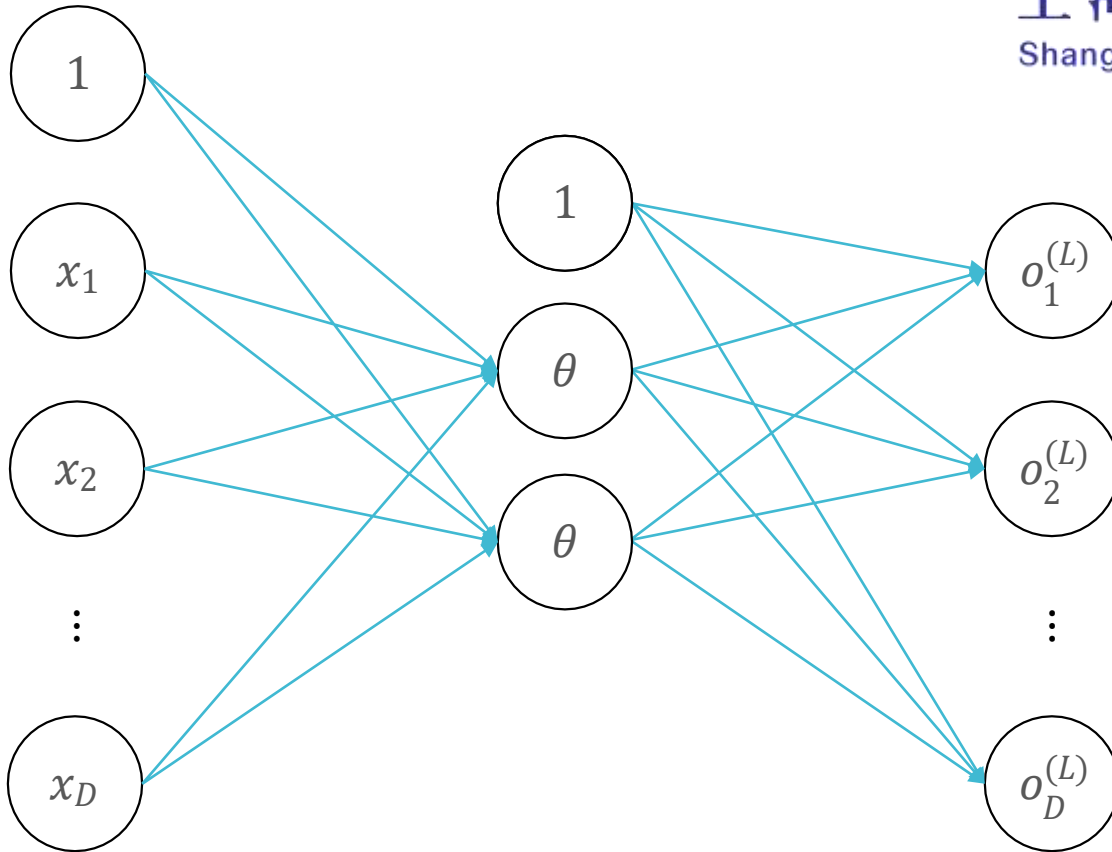
1. Define the sample mean, sample variance, and sample covariance of a vector-valued dataset
2. Identify examples of high dimensional data and common use cases for dimensionality reduction
3. Draw the principal components of a given toy dataset
4. Establish the equivalence of minimization of reconstruction error with maximization of variance
5. Given a set of principal components, project from high to low dimensional space and do the reverse to produce a reconstruction
6. Explain the connection between PCA, eigenvectors, eigenvalues, and covariance matrix
7. Use common methods in linear algebra to obtain the principal components

Autoencoders



Insight: neural networks implicitly learn low-dimensional representations of inputs in hidden layers

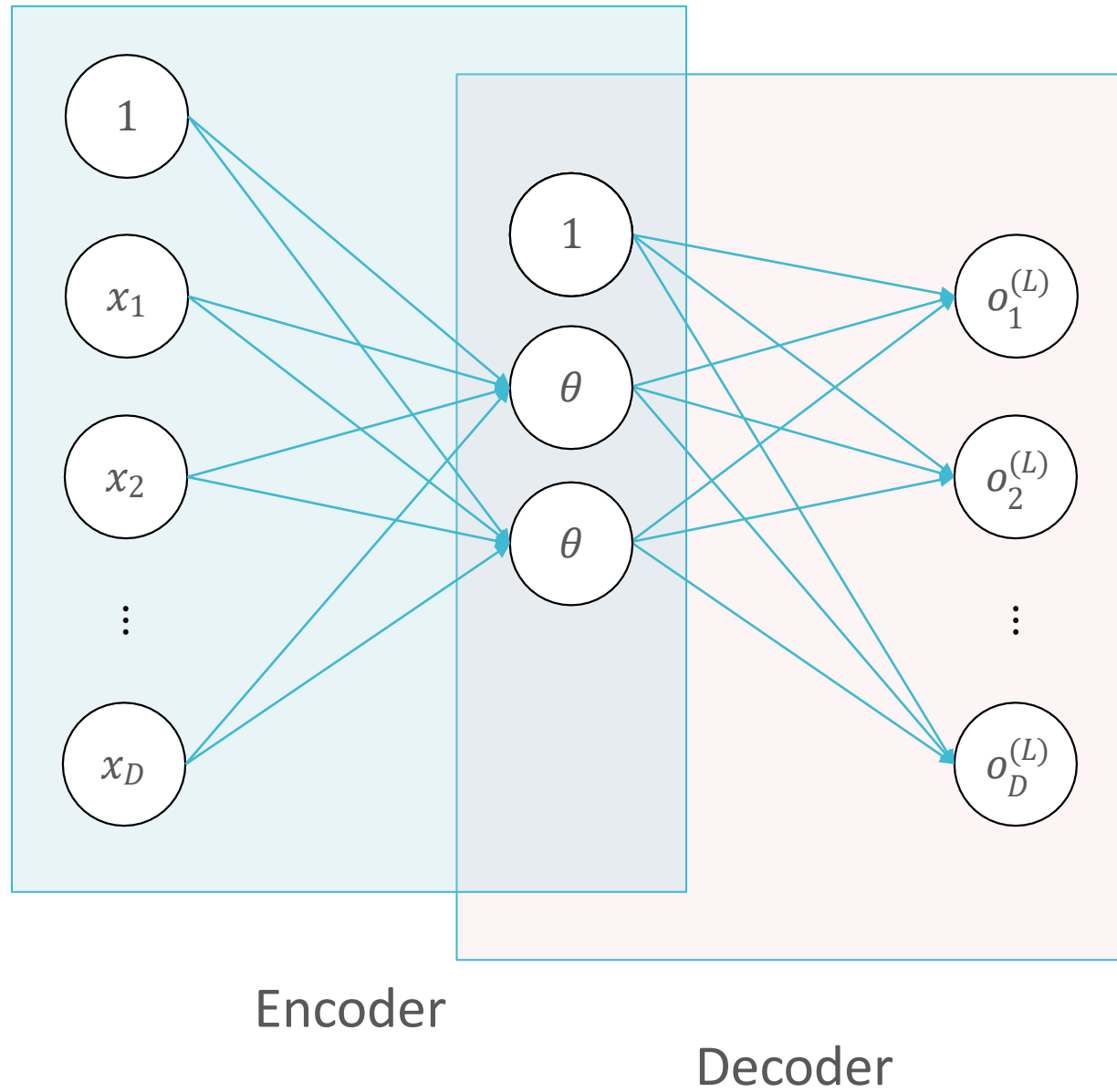
Autoencoders



- Learn the weights by minimizing the reconstruction loss:

$$e(\mathbf{x}) = \|\mathbf{x} - \mathbf{o}^{(L)}\|_2^2$$

Autoencoders



Deep Autoencoders

