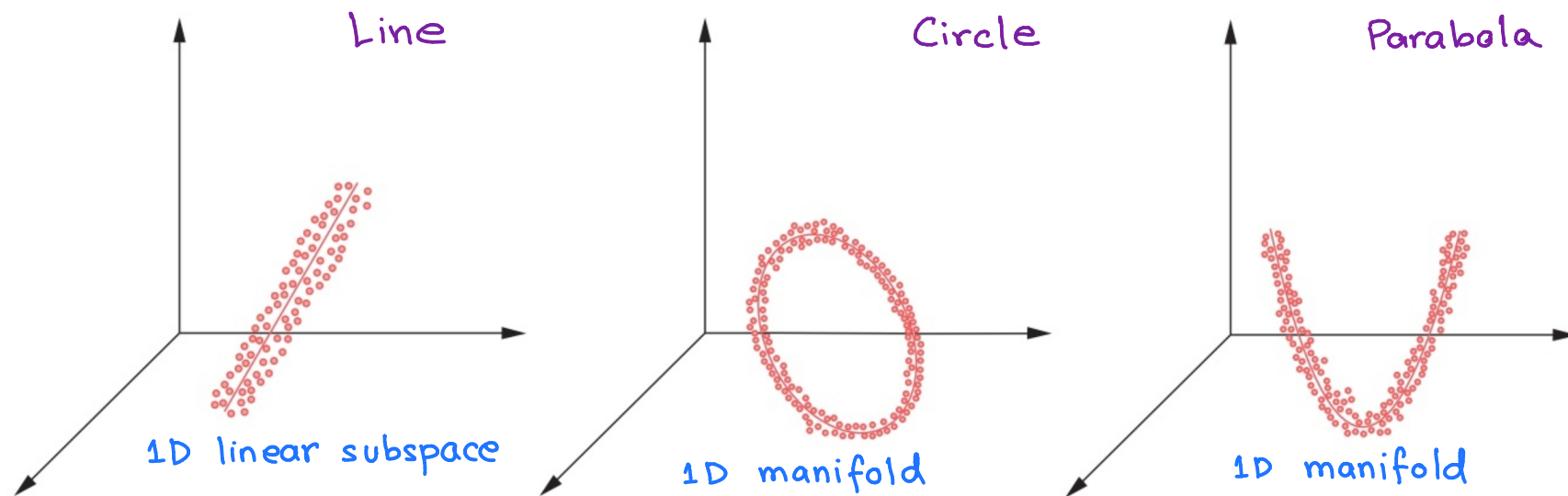


Dimensionality Reduction

- In unsupervised learning, we have seen clustering.
- In this lecture, we will look at dimensionality reduction
- In many practical applications, the input data $\underline{x} \in \mathbb{R}^p$ is a very high-dimensional, however, the **intrinsic dimensionality** may be quite small



In all three cases, the intrinsic dimensionality of data is 1

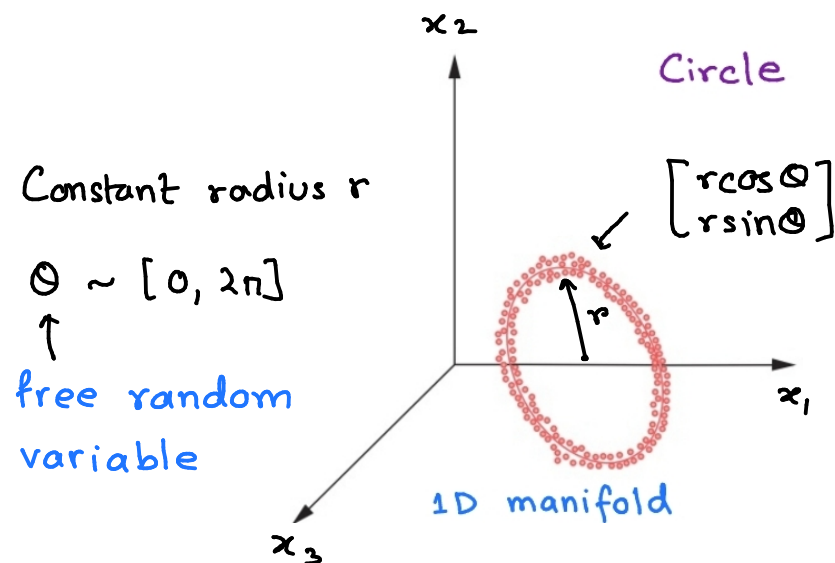
Intrinsic Dimensionality

- A data set $\{\underline{x}_i\}_{i=1}^N$, with $\underline{x} \in \mathbb{R}^P$, is said to have intrinsic dimensionality $M \leq P$, if the dataset can be described effectively in terms of 'M' free random variables

$$\underline{x} = g(\underline{u})$$

$\mathbb{R}^P \leftarrow \quad \quad \quad \rightarrow \mathbb{R}^M$

Example



The data lies along the circumference of a circle of radius r and a single free parameter θ suffices to describe the data

Intrinsic dimension = 1

Intrinsic Dimensionality

- An important concern in ML is learning from high-dimensional data x
- Success of ML, in particular deep learning, is due to its capability of

learning a useful representation of high-dimensional data

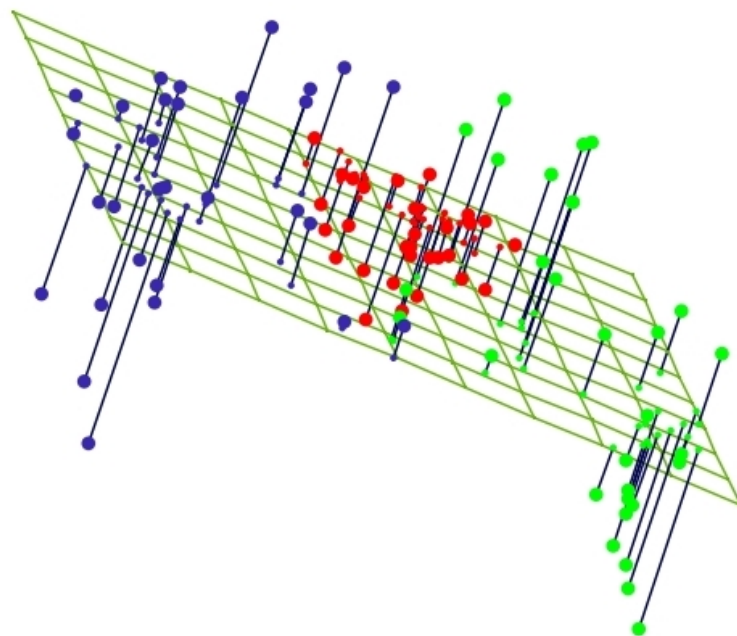
- One of the goals of unsupervised learning:

Learning a lower-dimensional subspace for encoding high-dimensional data set

- Idea of dimensionality reduction: Map data to a lower dimensional space
 - Save computational time in modelling high-dimensional data
 - Visualization in 2-dimensions can offer insights
 - Reduce overfitting and achieve better generalization

Linear Dimensionality Reduction

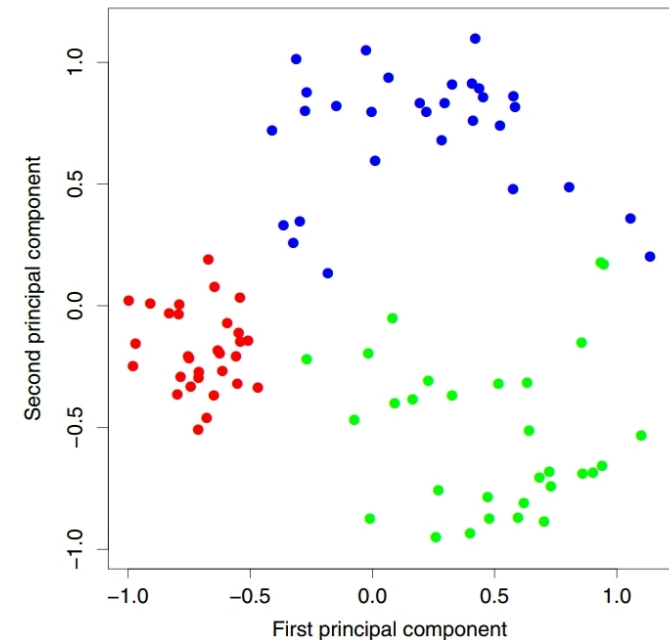
- We will introduce linear dimensionality reduction using Principal Component Analysis (PCA)
- PCA is also known as Karhunen-Loève (KL) transform
 - It falls under linear dimensionality reduction techniques



3D space

Projection
on a linear
subspace

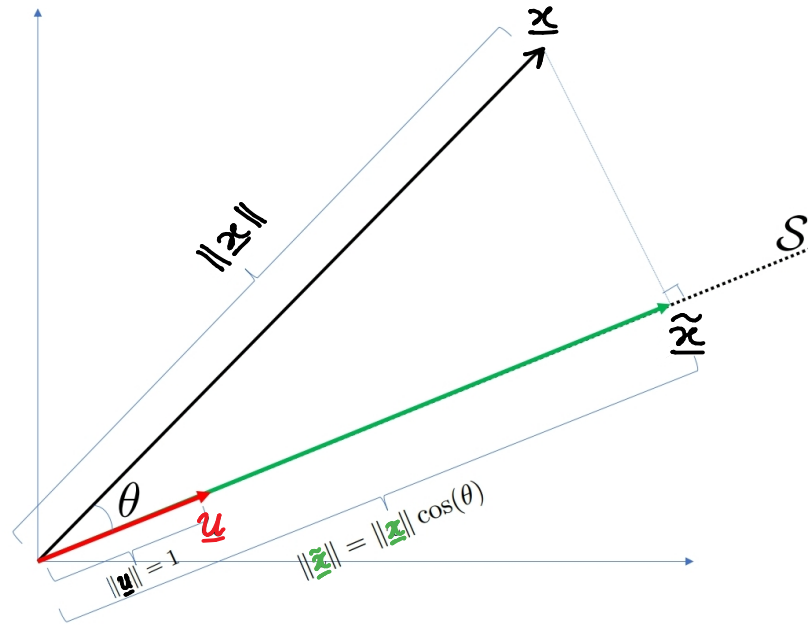
PCA



2D - space

Idea of projection

- Consider projection onto 1-D subspace of 2D



low dimensional

- Subspace S is the line along the unit vector \underline{u}
 - \underline{u} is the basis of S : Any point in S can be written as $z \underline{u}$ for some scalar z

- Projection of vector \underline{x} on S is denoted by $\tilde{\underline{x}} = \text{Proj}_S(\underline{x})$

- Recall that: $\underline{x}^T \underline{u} = \|\underline{x}\| \|\underline{u}\| \overset{1}{\cos(\theta)} = \|\underline{x}\| \cos \theta = \|\tilde{\underline{x}}\|$

- $\tilde{\underline{x}} = \text{Proj}_S(\underline{x}) = \underbrace{\underline{x}^T \underline{u}}_{\text{length of projection}} \cdot \underbrace{\underline{u}}_{\text{direction of projection}} = \|\tilde{\underline{x}}\| \underline{u}$

Idea of projection

from a $D > M$
space

- How to project onto an M -dimensional subspace?

- **Idea**: Choose an orthonormal bases $\{\underline{u}_1, \underline{u}_2, \dots, \underline{u}_M\}$ for S
- Project onto each unit vector individually (as in previous slide) and sum together the projections

- Mathematically, the projection is given as:

$$\tilde{\underline{x}} = \text{Proj}_S(\underline{x}) = \sum_{i=1}^M z_i \underline{u}_i \quad \text{where} \quad z_i = \underline{x}^T \underline{u}_i$$

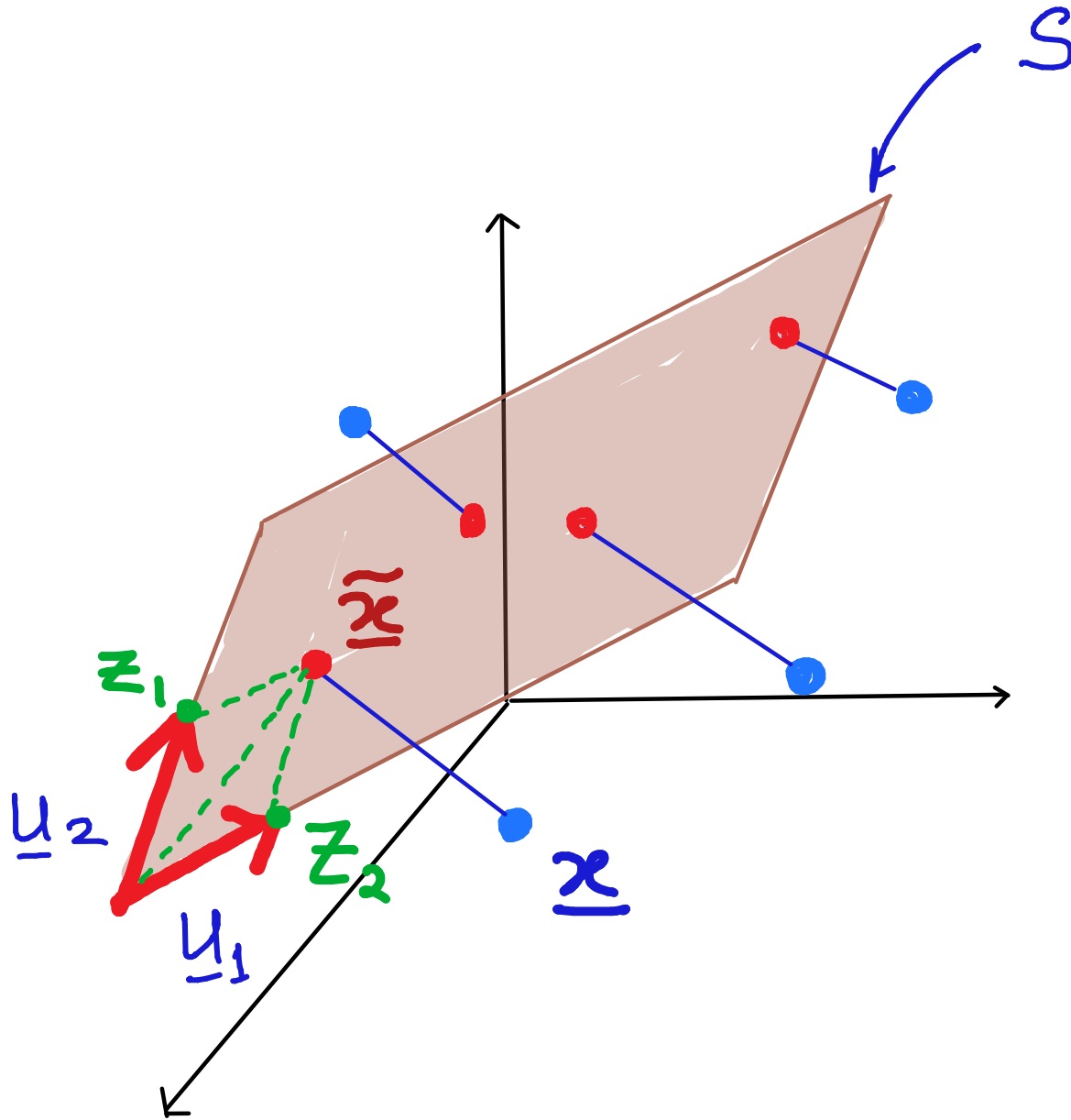
- each \underline{u}_i is the basis vector
- z_i is the magnitude along that projection

- In vector form:

$$\tilde{\underline{x}} = \text{Proj}_S(\underline{x}) = \underline{U} \underline{z} = \begin{bmatrix} | & | & & | \\ \underline{u}_1 & \underline{u}_2 & \dots & \underline{u}_M \\ | & | & & | \end{bmatrix} \begin{bmatrix} z_1 \\ z_2 \\ \vdots \\ z_M \end{bmatrix}, \quad \text{where} \quad \underline{z} = \underline{U}^T \underline{x}$$

\underline{z}

2D subspace in 3D space

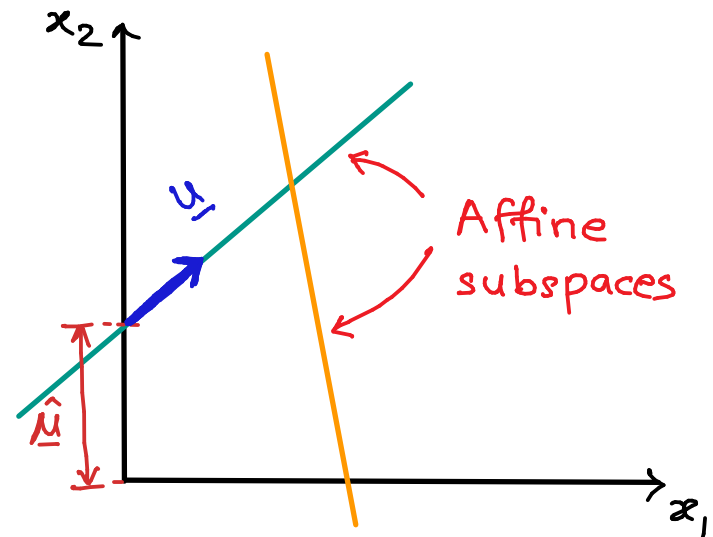
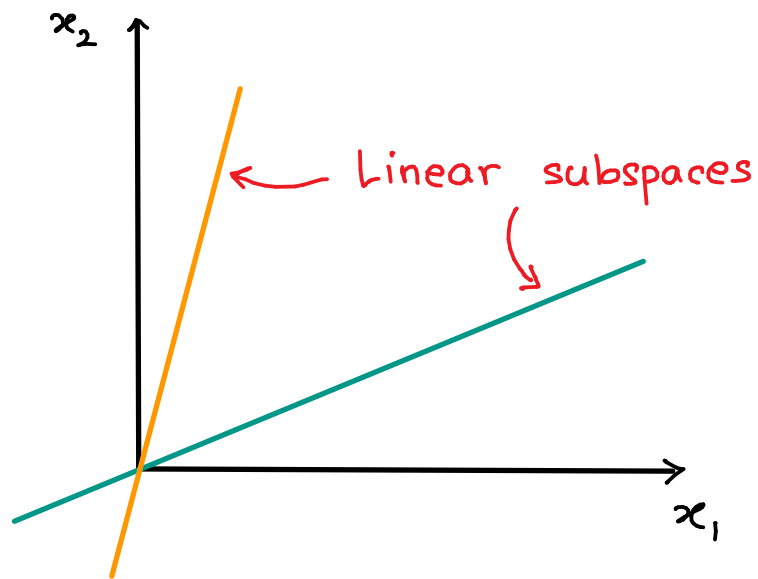


- Recall that every point in subspace S can be represented as $\{\underline{u}_1, \underline{u}_2\}$ & its linear combination

$$\tilde{\underline{x}} = z_1 \underline{u}_1 + z_2 \underline{u}_2$$

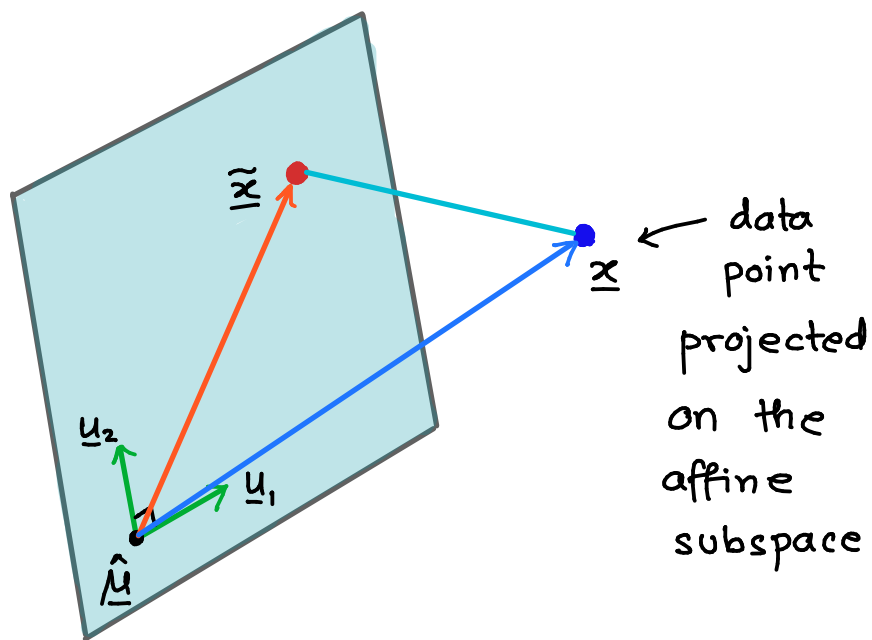
Projection onto an affine subspace

- So far, we have assumed a subspace that passes through zero
- However, the subspaces that we want to project onto can also be **affine subspaces**, which need not pass through zero



The affine subspaces can have an arbitrary origin \hat{u}

Projection onto an affine subspace



$$\begin{aligned}\tilde{x} &= \text{Proj}_S(x) \\ &= \underline{U} \underline{z} + \hat{\underline{\mu}} \\ &= z_1 \underline{u}_1 + z_2 \underline{u}_2 + \hat{\underline{\mu}}\end{aligned}$$
$$\underline{z} = \underline{U}^T (\underline{x} - \hat{\underline{\mu}})$$

($\underline{z} = \underline{U}^T \underline{x}$ earlier)

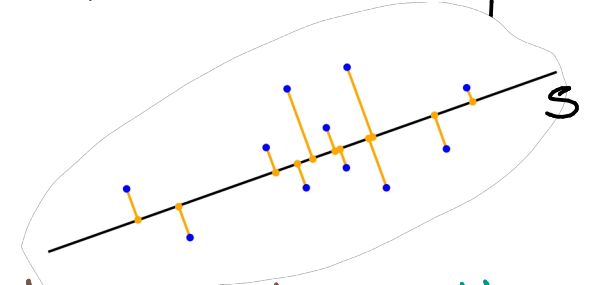
The affine subspace
has an origin $\hat{\underline{\mu}}$

- \tilde{x} is called the **reconstruction** of x
- \underline{z} is its **feature / code**
- If all the data points x lie close to the subspace, we could approximate x with its reconstructions \tilde{x}

$$x \approx \underline{U} \underline{z} + \hat{\underline{\mu}}$$

How to choose a good subspace?

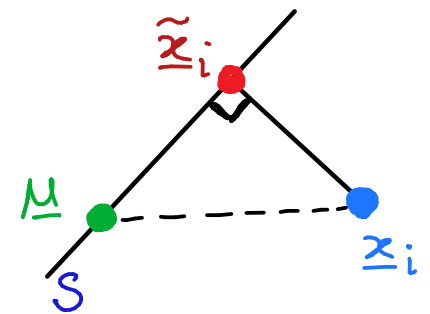
- We want to choose a subspace S which is low-dimensional compared to the dimension of the input space
- How to choose such a subspace S ?
 - We need to find appropriate $\hat{\underline{\mu}}$ and the orthogonal bases \underline{U}
 - origin $\hat{\underline{\mu}}$ can be set equal to the mean of the dataset
- To find \underline{U} , one of the two equivalent criteria could be followed:



- Minimize the reconstruction error:

equivalent (show)

$$\arg \min_{\underline{U}} \frac{1}{N} \sum_{i=1}^N \|\underline{x}_i - \tilde{\underline{x}}_i\|_2^2$$



- Maximize the variance of reconstructions: Find a subspace where the data has the most variability

$$\arg \max_{\underline{U}} \frac{1}{N} \sum_{i=1}^N \|\tilde{\underline{x}}_i - \hat{\underline{\mu}}\|_2^2$$

(You can show that \underline{x} and $\tilde{\underline{x}}$ have same mean)

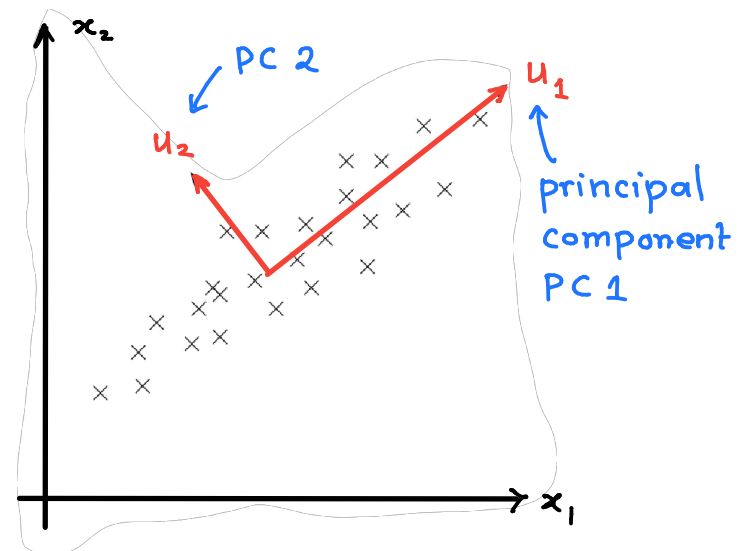
Principal Component Analysis

- Choosing a subspace to maximize the projected variance, or minimize the reconstruction error, is called **PCA**

- Consider the sample covariance matrix:

$$\hat{\Sigma} = \frac{1}{N} \sum_{i=1}^N (\mathbf{x}_i - \hat{\mu})(\mathbf{x}_i - \hat{\mu})^T$$

- $\hat{\Sigma}$ is symmetric and Positive semi-definite (PSD)
- The optimal PCA subspace is spanned by the **top 'M' eigenvectors** of $\hat{\Sigma}$
- These eigenvectors are called **principal components** or principal directions, much like the principal axes of an ellipse

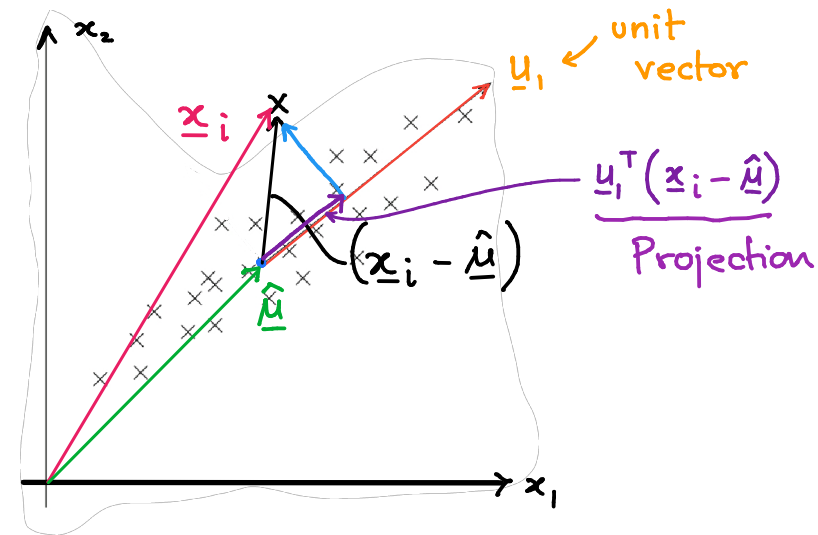


Derivation of PCA

- Let us consider the simplest case of finding a 1-D subspace
 - The goal then is to find a single direction represented by unit vector \underline{u}_1

- Lets maximize the projected variance

$$\begin{aligned} J(\underline{u}_1) &= \frac{1}{N} \sum_{i=1}^N \left(\underline{u}_1^T (\underline{x}_i - \hat{\underline{\mu}}) \right)^2 \\ &= \frac{1}{N} \sum_{i=1}^N \underline{u}_1^T (\underline{x}_i - \hat{\underline{\mu}}) (\underline{x}_i - \hat{\underline{\mu}})^T \underline{u}_1 \\ &= \underline{u}_1^T \hat{\sum} \underline{u}_1 \end{aligned}$$



- So the optimization task is:

$$\begin{aligned} \underline{u}_1 &= \underset{\underline{u}}{\operatorname{argmax}} \quad \underline{u}^T \hat{\sum} \underline{u} \\ \text{s.t.} \quad &\underline{u}^T \underline{u} = 1 \end{aligned}$$

$$\text{Lagrangian: } L(\underline{u}, \lambda) = \underline{u}^T \hat{\sum} \underline{u} - \lambda (\underline{u}^T \underline{u} - 1)$$

Take gradient and set to zero:

$$\hat{\sum} \underline{u} = \lambda \underline{u}$$

eigenvalue ← eigenvector

∴ Principal direction \underline{u}_1 is an eigenvector

- Since $\hat{\Sigma}$ is symmetric and PSD, all eigenvalues are **real** and **non-negative**: $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_p \geq 0$
- The 2nd principal component \underline{u}_2 is selected such that:
 - (a) \underline{u}_2 is orthogonal to \underline{u}_1
 - (b) \underline{u}_2 maximizes the variance after projecting the data onto the direction of \underline{u}_2
 - (c) The **2nd principal component** (or direction) is the eigenvector corresponding to the **2nd largest eigenvalue** of $\hat{\Sigma}$, λ_2
- Similar arguments can be used to show that the '**m**'th **principal component** is the '**m**'th eigenvector of $\hat{\Sigma}$
- The process continues until **M** principal components (corresponding to the **M** largest eigenvalues)

PCA decorrelates features

- The features (or code) are decorrelated by PCA

$$\text{Cov}(\underline{z}) = \text{Cov}(\underline{U}^T (\underline{x} - \hat{\underline{\mu}}))$$

$$= \underline{U}^T \text{Cov}(\underline{x}) \underline{U}$$

$$= \underline{U}^T \hat{\underline{\Sigma}} \underline{U}$$

$$= \underline{U}^T \underline{Q} \underline{\Lambda} \underline{Q}^T \underline{U}$$

$$= \begin{bmatrix} \underline{I} & \underline{0} \end{bmatrix} \underline{\Lambda} \begin{bmatrix} \underline{I} \\ \underline{0} \end{bmatrix}$$

$$= \text{top left } M \times M \text{ block of } \underline{\Lambda}$$

Spectral decomposition

$$\hat{\underline{\Sigma}}_{P \times P} = \underline{Q} \underline{\Lambda} \underline{Q}^T$$

eigenvector matrix

eigenvalues matrix

(orthonormal)

$$\underline{Q}_{P \times P} = \begin{bmatrix} \underline{U} & \vdots & \underline{U}_\perp \end{bmatrix}$$

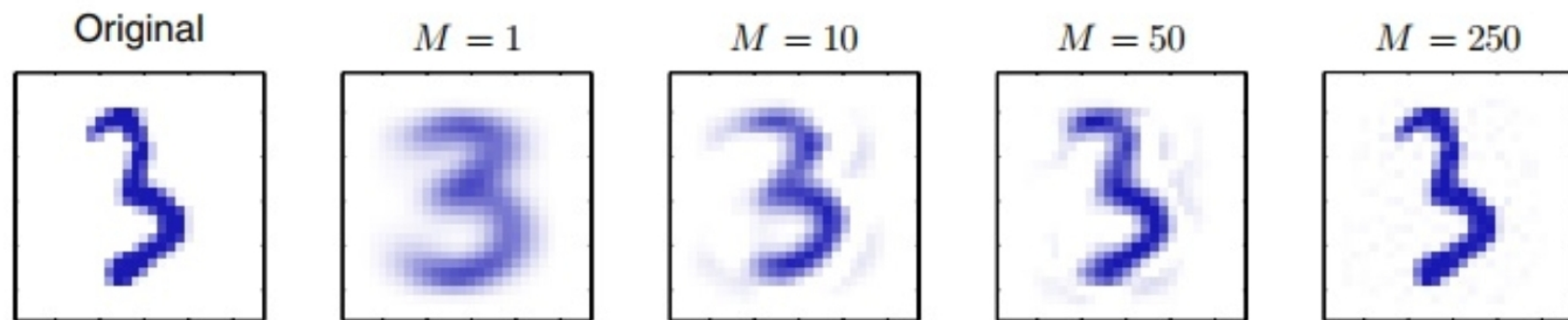
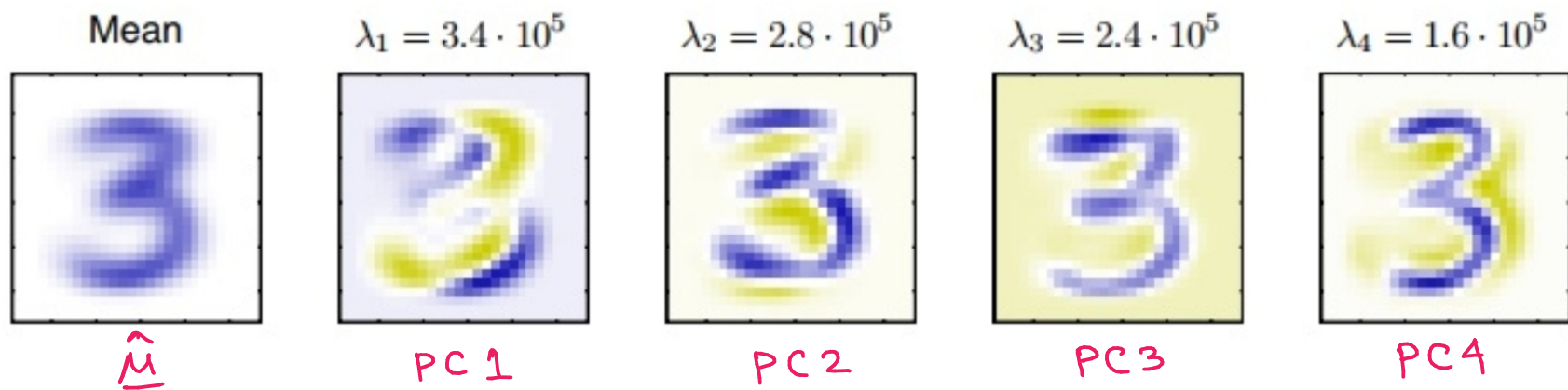
$P \times M$ $P \times (P-M)$

- Covariance of feature \underline{z} is diagonal \rightarrow uncorrelated

Summary of PCA

- Dimensionality reduction aims to find a low-dimensional representation of the data
- PCA projects the data onto an affine subspace that maximizes projected variance or minimizes the reconstruction error
- The optimal subspace is given by the top M eigenvectors of the sample covariance matrix, corresponding to the M largest eigenvalues
- PCA gives a set of decorrelated features

Example of data compression



Original
digit

PCA reconstructions