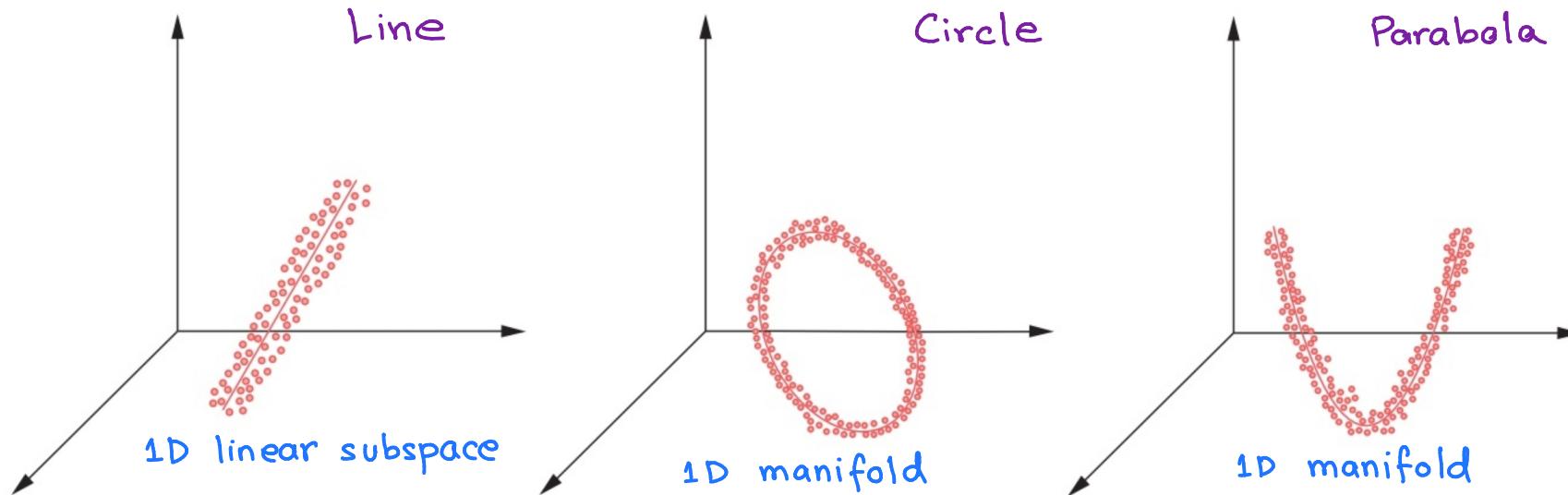


Dimensionality Reduction

- 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

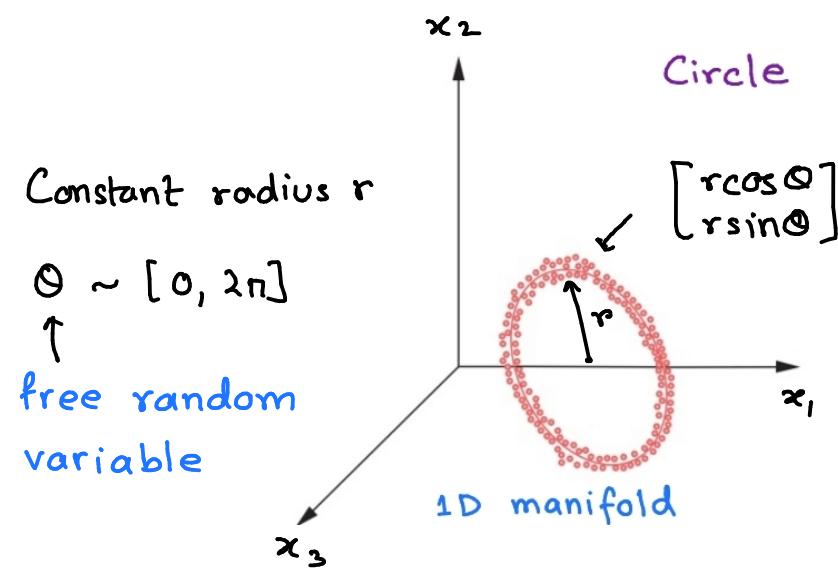
Intrinsic Dimensionality

- 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 ↓ \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

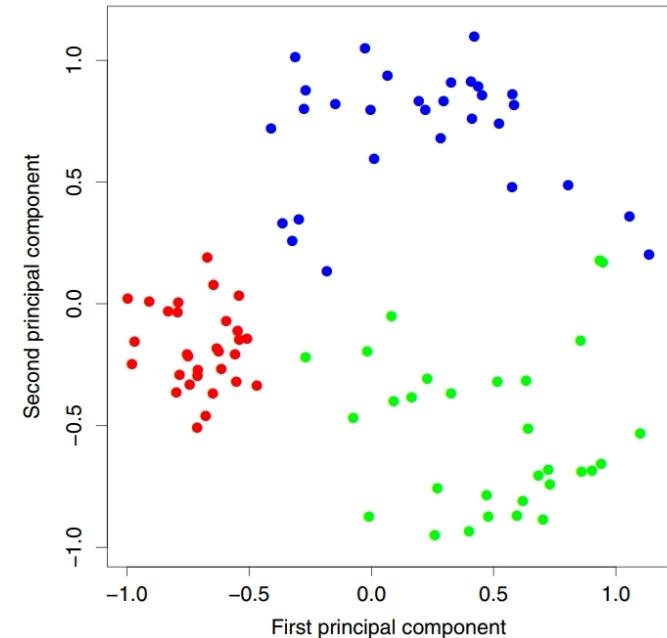
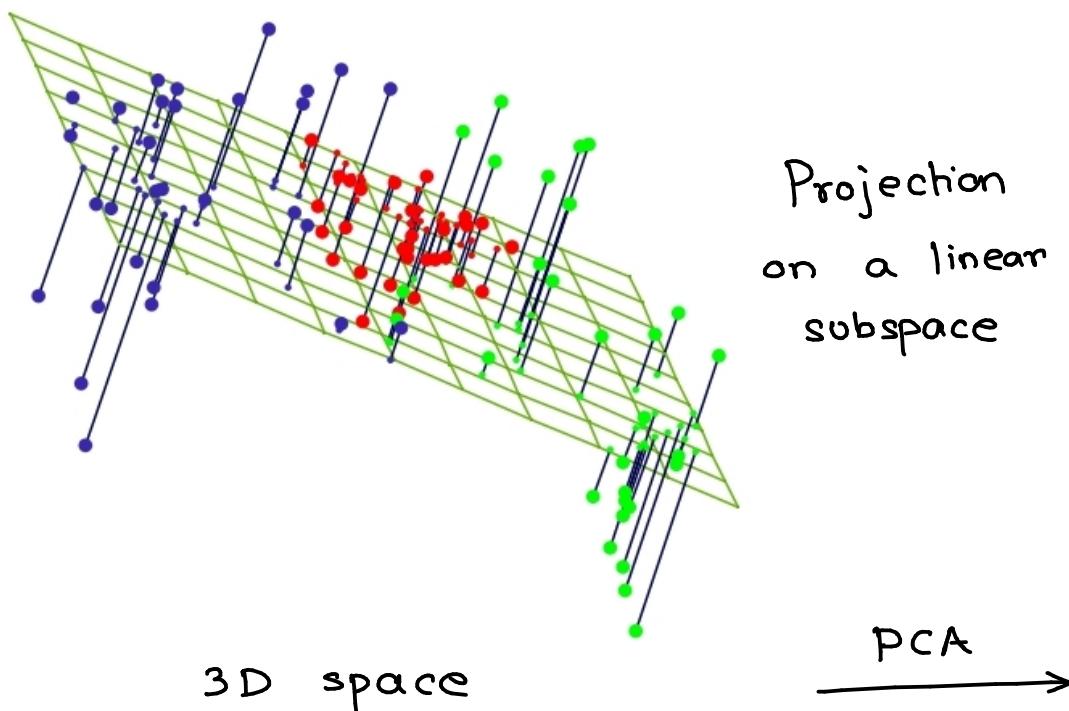
Intrinsic Dimensionality

- An important concern in ML is learning from high-dimensional data \geq
- 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

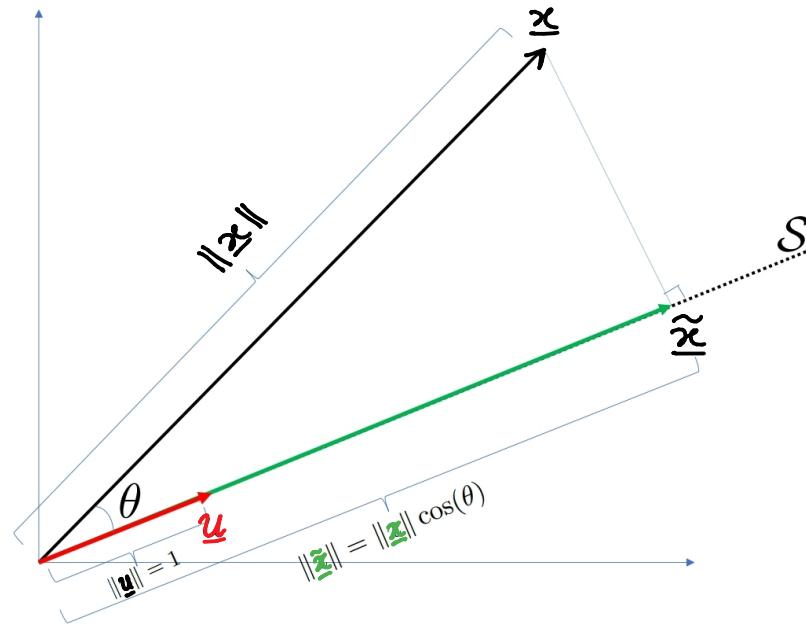
- We will introduce linear dimensionality reduction using Principal Component Analysis (PCA)
- PCA is also known as Karhunen-Loëve (KL) transform
 - It falls under linear dimensionality reduction techniques



2D - space

Idea of projection

- Consider projection onto 1-D subspace (a line)



- 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 $\underline{\tilde{x}} = \text{Proj}_S(\underline{x})$

- Recall that: $\underline{x}^T \underline{u} = \|\underline{x}\| \|\underline{u}\|^{\frac{1}{2}} \cos(\theta) = \|\underline{x}\| \cos \theta = \|\underline{\tilde{x}}\|$

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

Idea of projection

- How to project onto a 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:

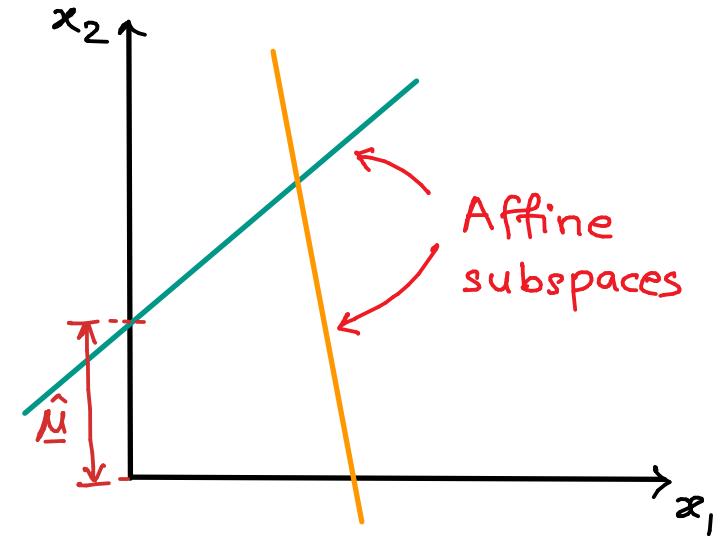
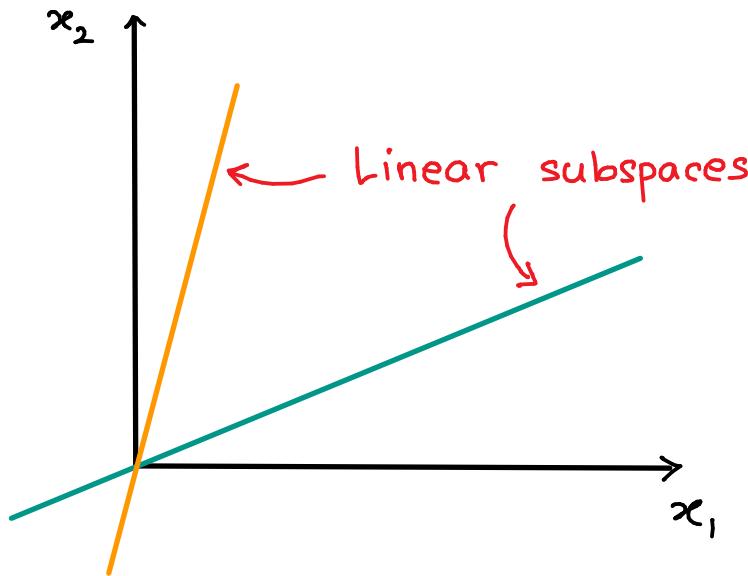
$$\tilde{\underline{x}} = \text{Proj}_S(\underline{x}) = \sum_{i=1}^M z_i \underline{u}_i \quad \text{where} \quad z_i = \underline{x}^T \underline{u}_i$$

- In vector form:

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

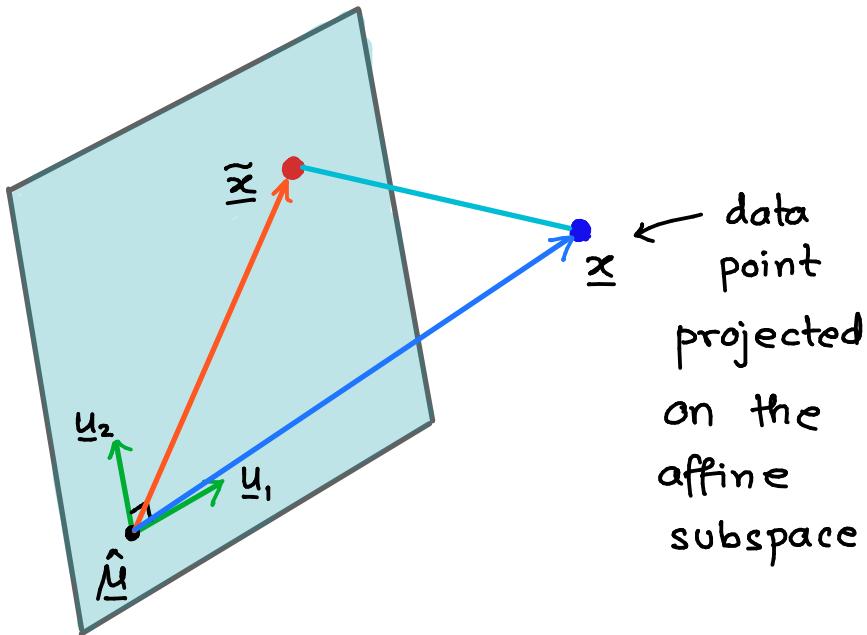
Projection onto an affine subspace

- 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



$$\tilde{x} = \text{Proj}_S(x)$$

$$= \underline{U} \underline{z} + \hat{\underline{u}}$$

$$= x_1 \underline{u}_1 + x_2 \underline{u}_2 + \hat{\underline{u}}$$

$$\underline{z} = \underline{U}^T (\underline{x} - \hat{\underline{u}})$$

The affine subspace
has an origin \hat{u}

- \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}

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

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

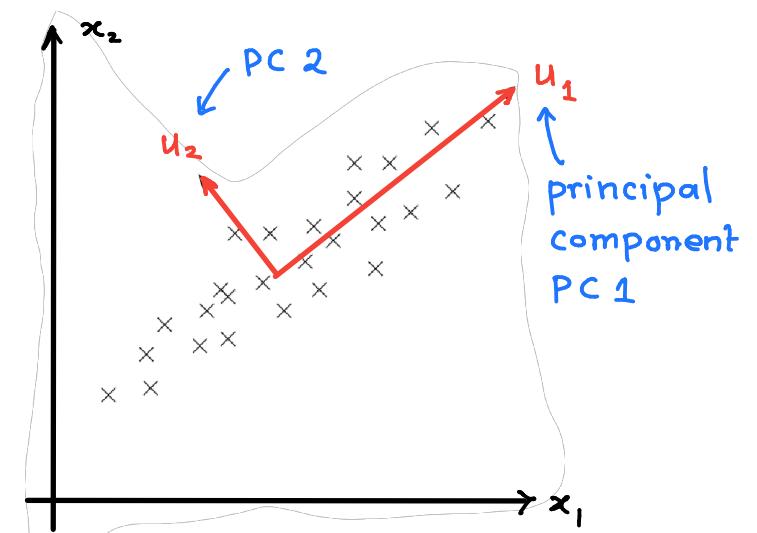
$$\arg \min_{\underline{U}} \frac{1}{N} \sum_{i=1}^N \| \underline{x}^{(i)} - \tilde{\underline{x}}^{(i)} \|_2^2$$

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

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

Principal Component Analysis

- Choosing a subspace to maximize the projected variance, or minimize the reconstruction error, is called PCA
- Consider the sample covariance matrix:
$$\hat{\Sigma} = \frac{1}{N} \sum_{i=1}^N (\underline{x}^{(i)} - \hat{\mu})(\underline{x}^{(i)} - \hat{\mu})^T$$
- $\hat{\Sigma}$ 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

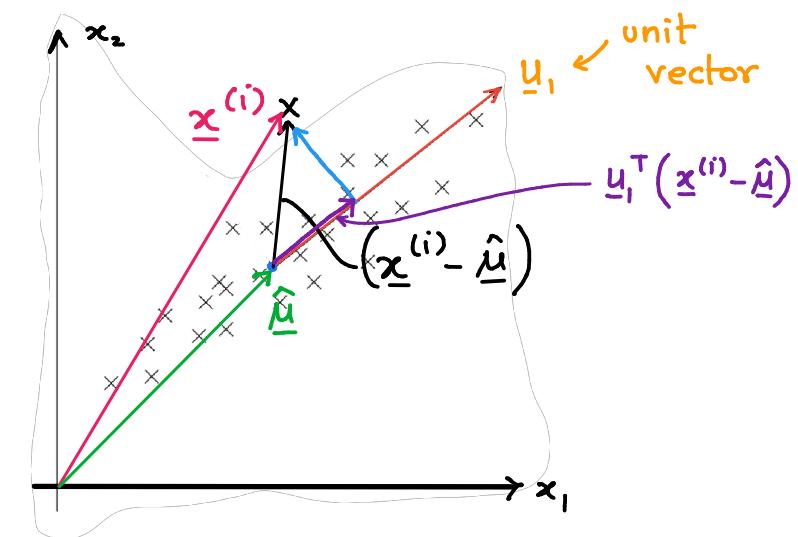


Derivation of PCA

- 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

- Let's maximize the projected variance

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



- So the optimization task is:

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

Lagrangian: $L(\underline{u}, \lambda) = \underline{u}^\top \hat{\Sigma} \underline{u} - \lambda (\underline{u}^\top \underline{u} - 1)$

Take gradient and set to zero:

$$\hat{\Sigma} \underline{u} = \lambda \underline{u} \quad \begin{matrix} \text{eigenvalue} \\ \leftarrow \underline{u} \text{ eigenvector} \end{matrix}$$

\therefore 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

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

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

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

$$= \underline{\boldsymbol{U}}^T \underline{\boldsymbol{Q}} \triangleq \underline{\boldsymbol{Q}}^T \underline{\boldsymbol{U}}$$

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

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

Spectral decomposition

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

\uparrow eigenvector matrix \uparrow eigenvalues matrix

(orthonormal)

$$\underline{\boldsymbol{Q}}_{P \times P} = \begin{bmatrix} \underline{\boldsymbol{U}} & | & \underline{\boldsymbol{U}}_L \\ \hline \underline{\boldsymbol{U}}_{P \times M} & | & \underline{\boldsymbol{U}}_{P \times (P-M)} \end{bmatrix}$$

- Covariance of feature $\underline{\boldsymbol{x}}$ 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

