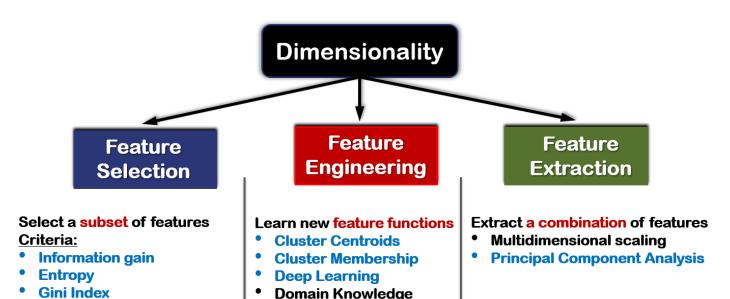
#### NCSU Python Exploratory Data Analysis

## **PCA: Dimension Reduction**



- Dimensionality and underdetermined problems
- Feature selection vs. feature extraction vs. feature engineering
- Dimension reduction (DR): unsupervised vs. supervised; linear vs. non-linear; orthogonal vs. non-orthogonal
- DR: linear, orthogonal with Principal Component Analysis (PCA)
- Eigenvalue and eigenvectors

#### **Prof. Nagiza F. Samatova**

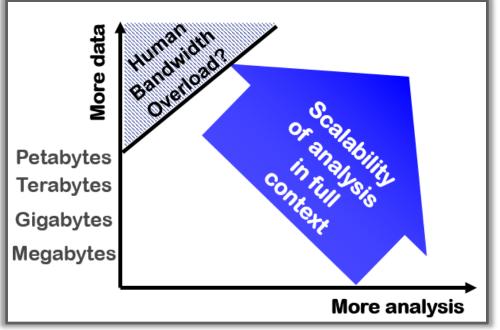
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Department of Computer Science North Carolina State University

Correlation

**Domain Knowledge** 

# Dimension Reduction MOTIVATION GAMMAN GAMMA



## What Analysis Methods to Use?



#### Analysis methods fail for a few gigabytes.

#### **Method Complexity:**

Calculate means O(n)

Calculate Histogram  $O(n \log(n))$ 

Calculate PCA  $O(n \cdot d)$ 

Clustering algorithms  $O(n^2)$ 

If n = 10GB, then what is O(n) or  $O(n^2)$  on a teraflop computers?

 $1GB = 10^9$  bytes  $1Tflop = 10^{12}$  op/sec

O(...)- in the order of

n – number of rows

d - number of columns

Data size n	Algorithm Complexity				
	n	nlog(n)	n <sup>2</sup>		
100B	10 <sup>-10</sup> sec.	10 <sup>-10</sup> sec.	10 <sup>-8</sup> sec.		
10KB	10 <sup>-8</sup> sec.	10 <sup>-8</sup> sec.	10 <sup>-4</sup> sec.		
1MB	10 <sup>-6</sup> sec.	10 <sup>-5</sup> sec.	1 sec.		
100MB	10 <sup>-4</sup> sec.	10 <sup>-3</sup> sec.	3 hrs		
10GB	10 <sup>-2</sup> sec.	0.1 sec.	3 yrs.		

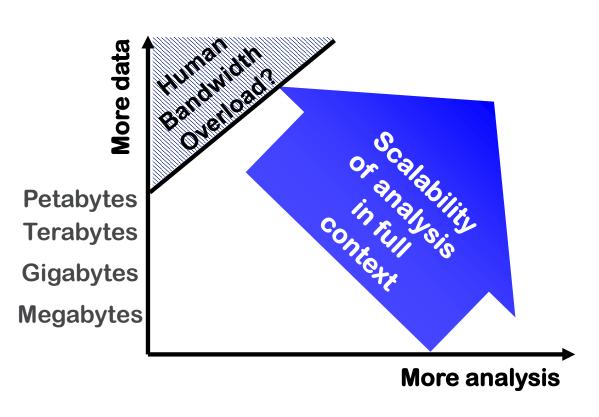
For illustration chart assumes 10<sup>-12</sup> sec. (1Tflop/sec) calculation time per data point

#### **How to Make Sense of Data?**





Not humanly possible to browse a petabyte of data. Analysis must reduce data to quantities of interest.



#### **Computations:**

Must be smart about which probe combinations to see!

#### **Physical Experiments:**

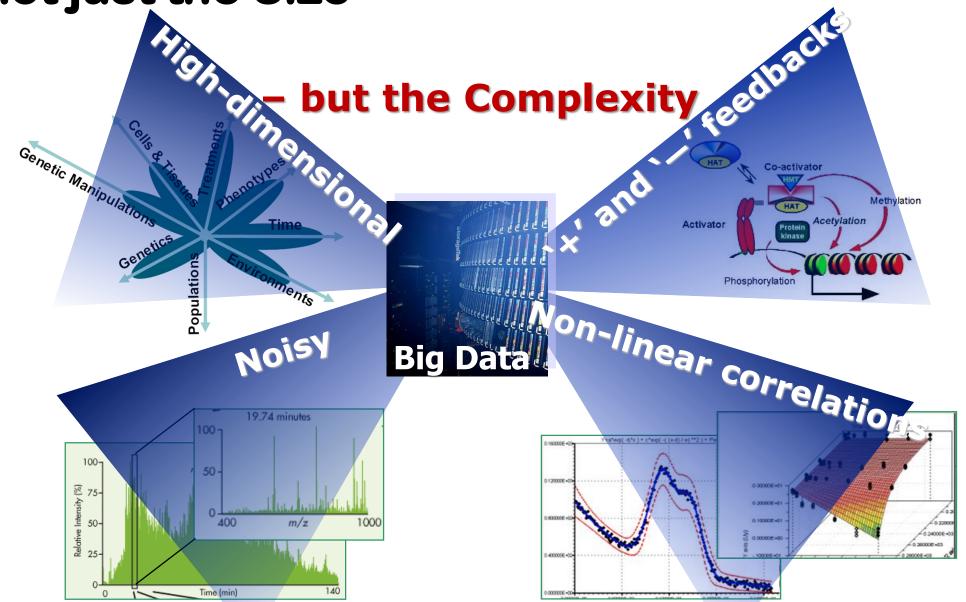
Must be smart about probe placement!

To see 1 percent of a petabyte at 10 megabytes per second takes:

35 8-hour days!

It is not just the Size





## **High-dimensional Data**

#### Text Data

- Record/Row: Document ID
- Dimension/Column: Each word in a collection of text documents

#### Image Data

- Record/Row: Image ID
- Dimension/Column: Each pixel

#### Audio Data

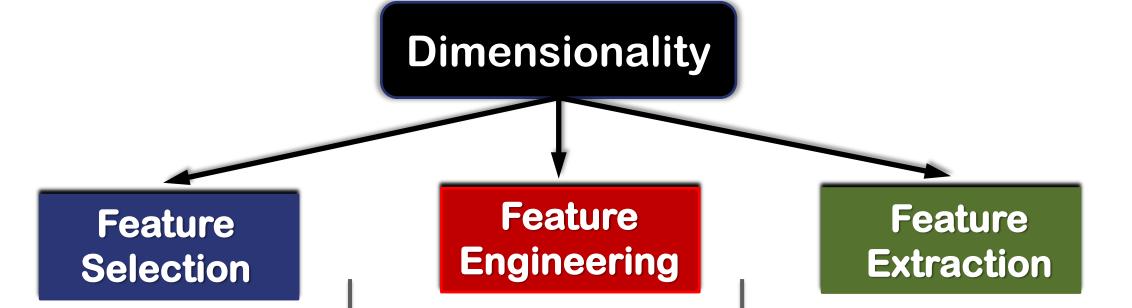
- Record/Row: Audio record ID
- Dimension/Column: Frequency of an audio signal OR a word if audio → text conversion

#### Sales Data

- Record/Row: Transaction ID
- Dimension/Column: Each Product ID

# Taxonomy: Dimensionality-aware Methods FEATURE SELECTION, FEATURE EXTRACTION, FEATURE ENGINEERING

## Dimensionality Challenge: Strategies to Cope With



#### Select a subset of features Criteria:

- Information gain
- Entropy
- Gini Index
- Correlation
- Domain Knowledge

#### **Learn new feature functions**

- Cluster Centroids
- Cluster Membership
- Deep Learning
- Domain Knowledge

#### Extract a combination of features

- Multidimensional scaling
- Principal Component Analysis

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## **Motivation: Why to Cope with Dimensionality Issue?**

- Not all the measured variables are important for understanding the underlying "interesting" phenomena:
  - could complicate the process of data analysis
- Decrease the computational cost for other data mining tasks:
  - Proximity measure calculations:  $O(a) \rightarrow O(k)$ , d>>k
- Reduce the noise in the data:
  - improve signal to noise ratio
- Improve the accuracy of predictive models
  - better designed/engineered features capable of capturing non-linear signals in the data
- Reduce collinearity among variables/features
  - Critical for regression models

## What is Dimensionality Reduction (DR)?

Objective: To transform data from a high-dimensional representation to a low-dimensional representation, while "best" preserving the information.

#### Given dataset with *n* objects:

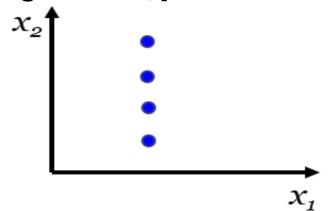
$$X \in \mathbb{R}^{n \times m}$$
  $\longrightarrow$   $X' \in \mathbb{R}^{n \times p}$   $M \in \mathbb{R}^{n \times p}$   $M \in \mathbb{R}^{n \times p}$  where  $p << m$ 

#### **Text Mining Example:**

•  $m = 50,000 \text{ words} \rightarrow p \approx 100 \text{ extracted features}$ 

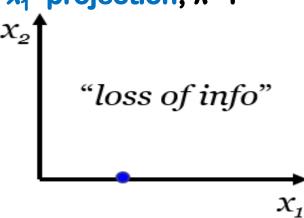
## **Example 1: p=2→d=1**

#### Original data, p=2

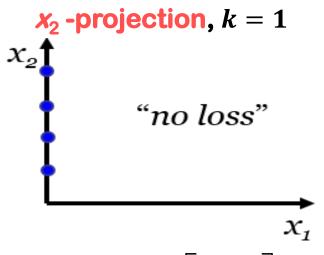


$$X_{4..2}$$

#### $x_1$ -projection, k=1



$$X'_{4\times 2} = X_{4\times 2} \cdot \begin{bmatrix} 1 & 0 \\ 0 & 0 \end{bmatrix}$$



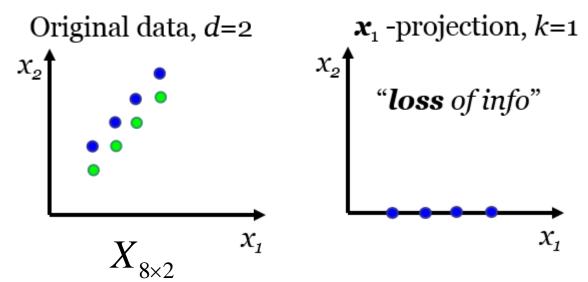
$$X_{4\times 2}' = X_{4\times 2} \cdot \begin{bmatrix} 0 & 0 \\ 0 & 1 \end{bmatrix}$$

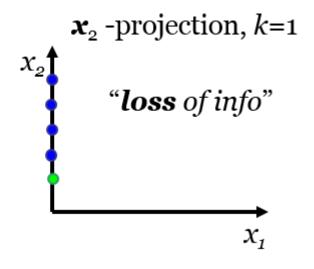
$$X'_{m \times d} = X_{m \times d} \cdot P_{d \times d}$$

$$\boldsymbol{P}_{d\times d} = \begin{bmatrix} 1 & 0 & \dots & 0 \\ 0 & 0 & \dots & 0 \\ \dots & \dots & \dots & \dots \\ 0 & 0 & 0 & 1 \end{bmatrix} - \text{ projection matrix; some diagonal elements are } 0$$

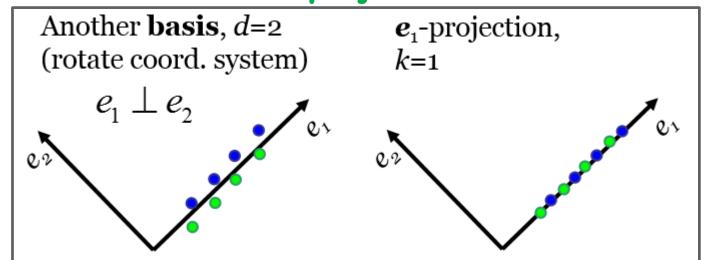
#### Which projection is "better"?

## **Example 2: Linear, Orthogonal Projection**





#### Is there a "better" projection?



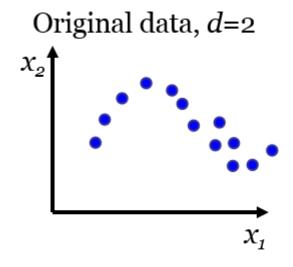
#### **Projection:**

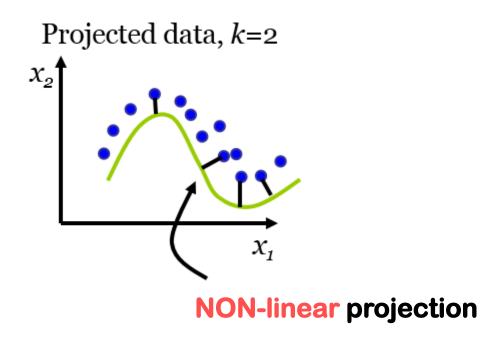
- Linear, e<sub>1</sub> line
- Orthogonal

$$e_1 \perp e_2$$

 $e_1$ ,  $e_2$  – eigenvectors of X

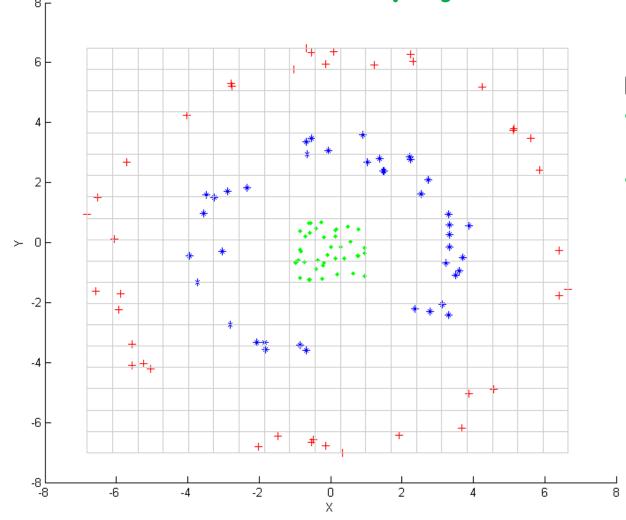
## **Example 3: Non-linear projection**





## **Example 4: Projection for Labeled Data**

#### What is a "better" projection?



#### Ideal for machine learning (ML) algorithm:

- Points within the same group come from a Gaussian distribution (spherical shapes)
- Points from different groups are linearly separable

#### "better" means:

- it bests, ideally, linearly separates different groups of data, i.e.
- points from the same group are closer to each other and are farther away from the points in different groups

## **Summary: Taxonomy of Dimension Reduction**

- Linear vs. NON-linear
- Orthogonal vs. non-orthogonal
- Unsupervised (unlabeled data) vs. supervised (labeled data)

#### **Linear dimension reduction:**

- Interpretable in original space
- Preserves non-linearity for visualization
- Orthogonal projections
- Non-orthogonal projections
- Favored for structure discovery

#### **Non-linear dimension reduction:**

- Lower-dimensional representation
- Interpretable w.r.t. Non-linear transformation
- 1 to 3 orders of magnitude more computation
- Favored for prediction or classification

- A lower-dimensional representation that contains the essence of the high dimensional data
- Blessing of dependence/correlation that saves us from the curse of dimensionality

## Linear Orthogonal Dimension Reduction PCA: PRINCIPAL COMPONENT ANALYSIS

## Dimensionality Challenge: Strategies to Cope With

Dimensionality

## Feature Selection

Select a subset of features Criteria:

- Information gain
- Entropy
- Gini Index
- Correlation
  - **Domain Knowledge**

Feature Engineering

#### **Learn new feature functions**

- Cluster Centroids
- Cluster Membership
- Deep Learning
- Domain Knowledge

Feature Extraction

**Extract a combination of features** 

- Multidimensional scaling
- Principal Component Analysis

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## What is "better" projection: $d \rightarrow k$ ( k < d)?

- Many definitions are possible
- Definition 1:
  - Projection that maximizes the VARIANCE of the original d-dimensional data upon its projection onto the target k-dimensions (k < d)

## Python Code Example: Iris Data

#### original 4-d data in 2-d

```
pca = decomposition.PCA(n components=3)
pca.fit(X)
X rot = pca.transform(X)
# Plot rotated 4-D iris data in 2D: the first two PCs
mglearn.discrete scatter(X rot[:, 0], X rot[:, 1], y)
plt.title("PCA Rotated: Iris Data Set")
plt.xlabel("PC-1")
plt.ylabel("PC-2")
plt.show()
                            PCA Rotated: Iris Data Set
                 1.5
                 1.0
                 0.5
                0.0
                -1.0
```

PCA-transformed data in 2-d

```
Iris Data Set
                                 2.0
                                                                    7.0
                                                                         7.5
iris = datasets.load iris()
                                                     sepal length (cm)
```

```
# Plot original 4-D iris data in 2D
mglearn.discrete scatter(X[:, 0], X[:, 1], y)
plt.title("Iris Data Set")
plt.xlabel("sepal length (cm)")
plt.ylabel("sepal width (cm)")
plt.show()
```

X = iris.data

print(X[0:5])

y = iris.target

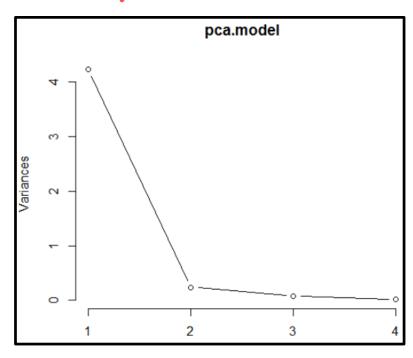
print(iris.feature names)

print (iris.target names)

print (y[[0, 51, 101]])

## Visualizing PCA Results: Iris Data

#### Screeplot

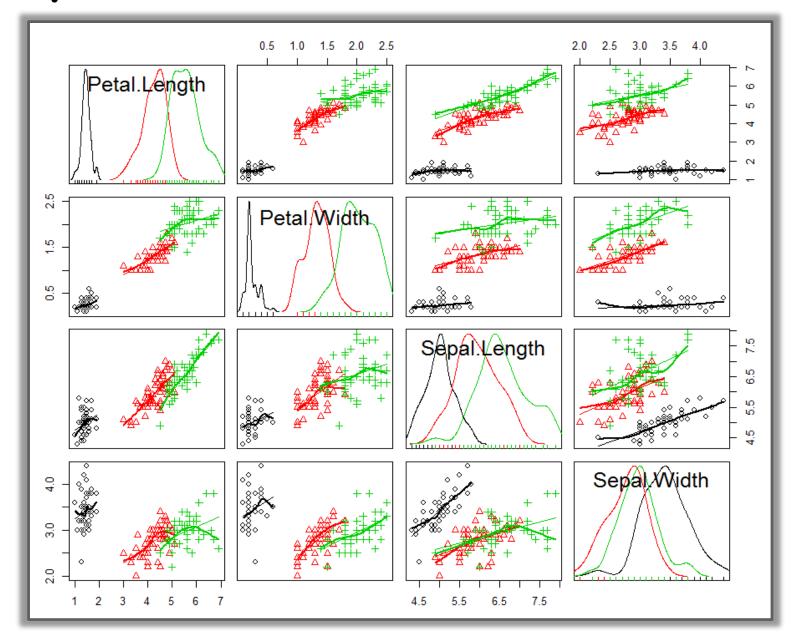


Component loadings: Comp.1 Comp.2 Comp.3 Comp.4 Petal.Length 0.5804131 0.02449161 0.1421264 0.8014492 Petal.Width 0.5648565 0.06694199 0.6342727 -0.5235971 Sepal.Length 0.5210659 0.37741762 -0.7195664 -0.2612863 Sepal.Width -0.2693474 0.92329566 0.2443818 Component variances: Comp.1 Comp.2 Comp.3 Comp.4 2.91849782 0.91403047 0.14675688 0.02071484 Importance of components: Comp.1 Comp.2 Comp.3 Comp. 4 Standard deviation 1.7083611 0.9560494 0.38308860 0.143926497 Proportion of Variance 0.7296245 0.2285076 0.03668922 0.005178709 Cumulative Proportion 0.7296245 0.9581321 0.99482129 1.000000000

Successive variance accounted by each component

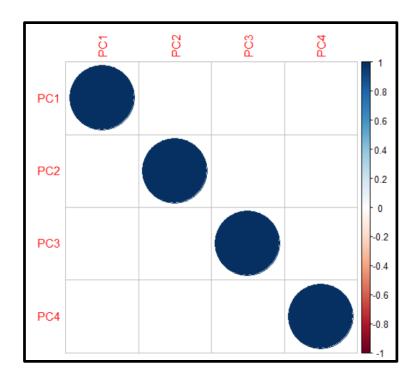
Original Relationships: Features are Correlated!

**Scatterplot Matrix** 



## **Extracted Principal Components (PCs) are Uncorrelated**

	PC1	PC2	PC3	PC4
PC1	1e+00	-1e-16	-1e-15	2e-15
PC2	-1e-16	1e+00	2e-16	-5e-16
PC3	-1e-15	2e-16	1e+00	-1e-15
PC4	2e-15	-5e-16	-1e-15	-1e-15 1e+00



The new extracted features (PCs) are UNCORRELATED!

#### PC is a weighted linear sum of original features

#### **Extracted Feature > PCA-based Feature Extraction**:

$$PC = w_1 * f_1 + w_2 * f_2 + \cdots + w_d * f_d$$

The magnitude of each weight indicates how important the corresponding feature is → it could be used as a feature selection technique!

	PC1	PC2	PC3	PC4
Sepal.Length	0.36	-0.66	0.58	0.3
Sepal.Width	-0.08	-0.73	-0.60	-0.3
Petal.Length	0.86	0.17	-0.08	-0.5
Petal.Width	0.36	0.08	-0.55	0.8

## Preserved Variability for Top-k PCs: d → k (k << d)

Percentage of variability preserved if the first k PCs are used for projection:

$$\frac{\sum_{i=1}^k \lambda_i}{\sum_{j=1}^d \lambda_j}$$

#### Standard deviation = sqrt (Variance)

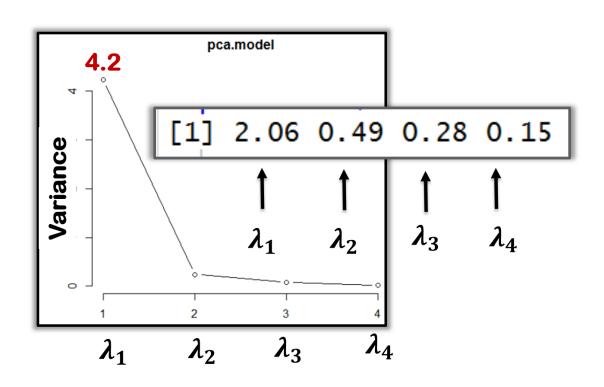
Standard deviations (1, .., p=4): [1] 2.06 0.49 0.28 0.15

#### Variance preserved by PC1:

$$2.06 \times 2.06 = 4.2$$

#### **SORTED Eigenvalues:** pca.model\$sdev

$$\lambda_1 \ge \lambda_2 \ge \dots \ge \lambda_k \ge \dots \ge \lambda_d \ge 0$$



## **Eigenvalues: Importance of Eigenvectors**

**SORTED:** 
$$\lambda_1 \geq \lambda_2 \geq ... \geq \lambda_k \geq ... \geq \lambda_d \geq 0$$

#### Importance of Principal Components (PC), or Eigenvectors, or Rotations

PC1 preserves more variance than PC2

PC2 preserves more variance than PC3

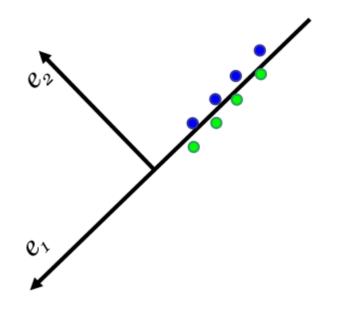
#### **Proportion of Variance Preserved if only k PCs are used:**

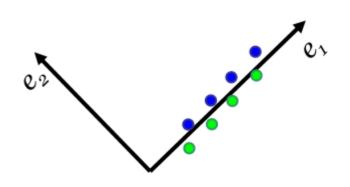
$$\frac{\sum_{i=1}^k \lambda_i}{\sum_{j=1}^d \lambda_j}$$

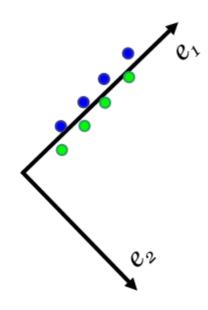
## Principal Components (eigenvectors) are NOT unique

#### **PCA: Linear Orthogonal Transformation:**

- PCs are simply a rotation of the original coordinate system
- Each PC is a line
- Each PC is perpendicular to the other PCs
  - PCs are Uncorrelated
  - The angle between any pair of PCs is 90°

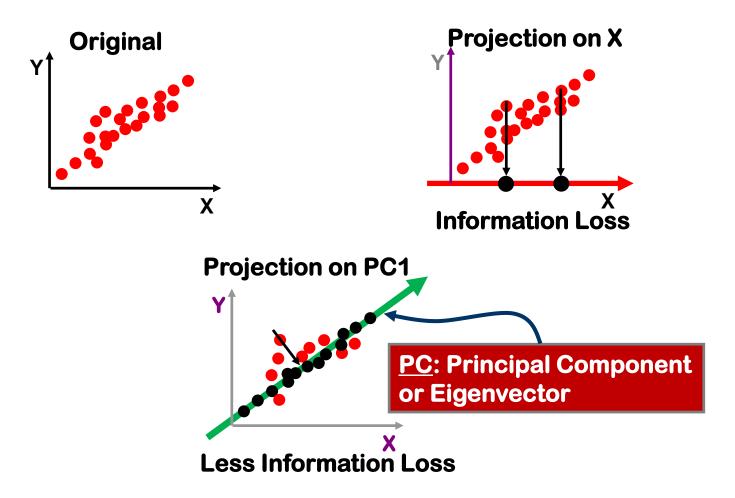






## PCA: Linear Orthogonal Dimension Reduction

Principal Component Analysis (PCA) finds intrinsic dimensionality and allows for low-dimensional representation of the data.



Principal Component Analysis is the Spectral Decomposition of the Covariance Matrix.

## **PCA: Covariance vs. Correlation Matrix**

- PCA results depend on the scales at which the variables are measured:
  - PCA should only be used with the raw data if all variables have the same units of measure
- PCA results depend on the variances of the variables: the ones with the highest sample variances will tend to be emphasized in the first few principal components:
  - Use PCA with covariance matrix only if you wish to give variables with higher variances more weight in the analysis
- If the variables either have different units of measurement (i.e., pounds, feet, gallons, etc), or if we wish each variable to receive equal weight in the analysis, then the variables should be standardized (Z-scores) before a principal components analysis is carried out.