



Université Constantine 2
جامعة قسنطينة 2

Module : Machine Learning (ML – SDSI)

– Course 3 –

Chapter 3 : Unsupervised Learning

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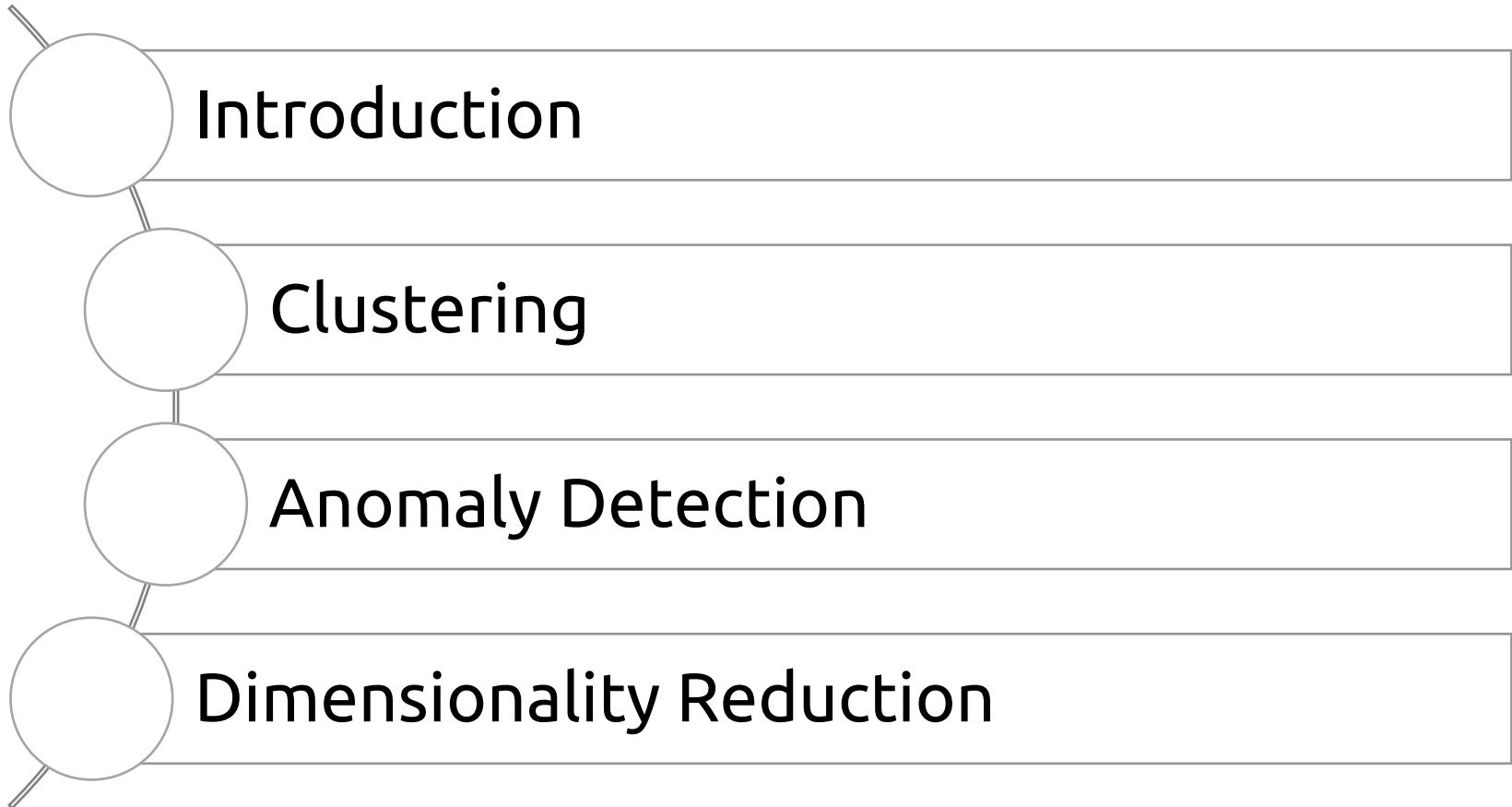
Etudiants concernés

Faculté/Institut	Département	Niveau	Spécialité
NTIC	TLSI	M1	SDSI

Goals of the Chapter

- Grasp Unsupervised Learning Basics
- Learn Key Clustering Algorithms such as **K-Means**, **Hierarchical Clustering**, and **DBSCAN**
- Assess Clustering Quality like **Silhouette Score**, **Davies-Bouldin Index**, and **Adjusted Rand Index**
- Explore Dimensionality Reduction , such as **PCA** and **t-SNE**
- Apply to Real-World Use Cases (e.g., segmentation, anomaly detection)

Main Titles



Introduction

Introduction

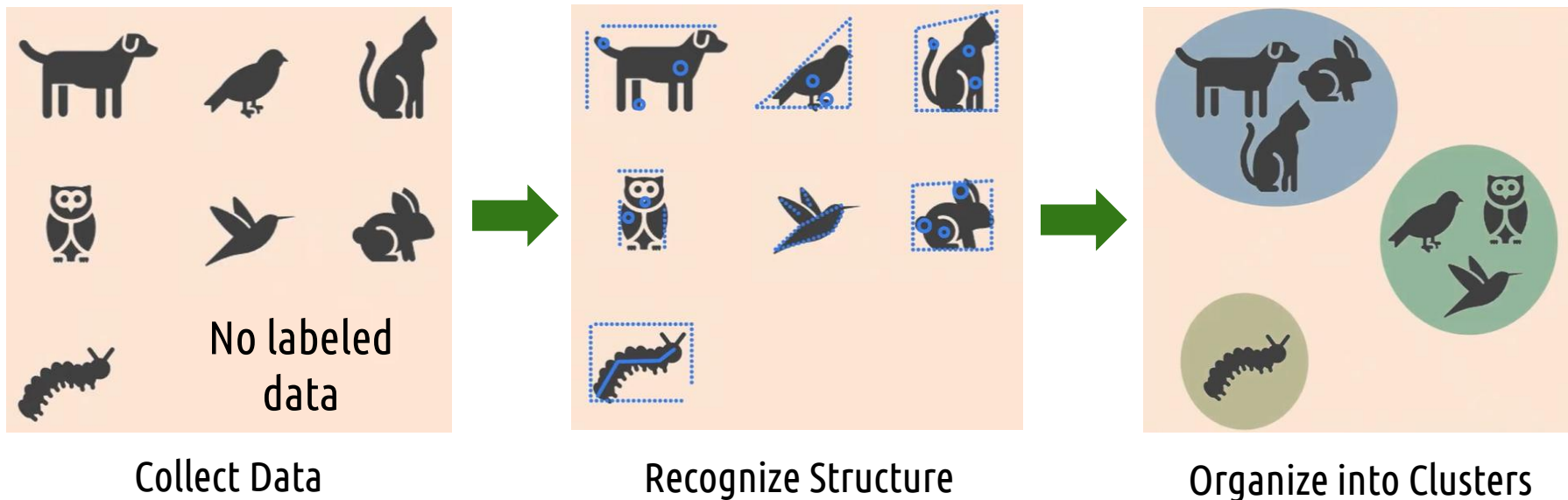
What is Unsupervised Learning?

- Unsupervised learning deals with datasets that have no labels. The goal is to discover the inherent structure in the data.
- Key Tasks:
 - **Clustering**: Group similar instances.
 - **Dimensionality Reduction**: Reduce the number of features while retaining key structure.
 - **Anomaly Detection**: Identify rare or abnormal observations.

Introduction

Clustering

- Clustering is a method to organize unlabeled data by finding natural groups based on similarities.
- Example:



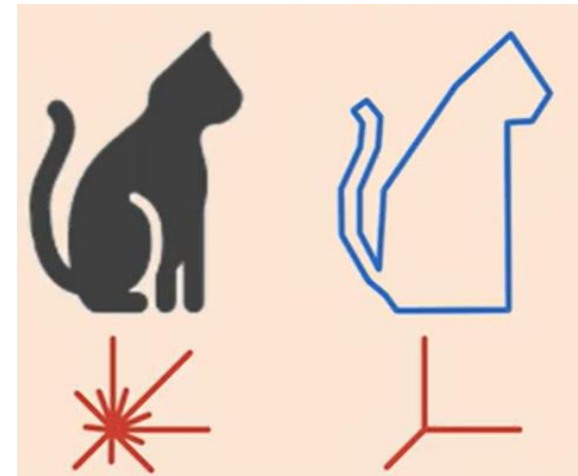
Introduction

Dimensionality Reduction

- Transform high-dimensional data into a lower-dimensional representation while retaining its meaningful structure.

Goals:

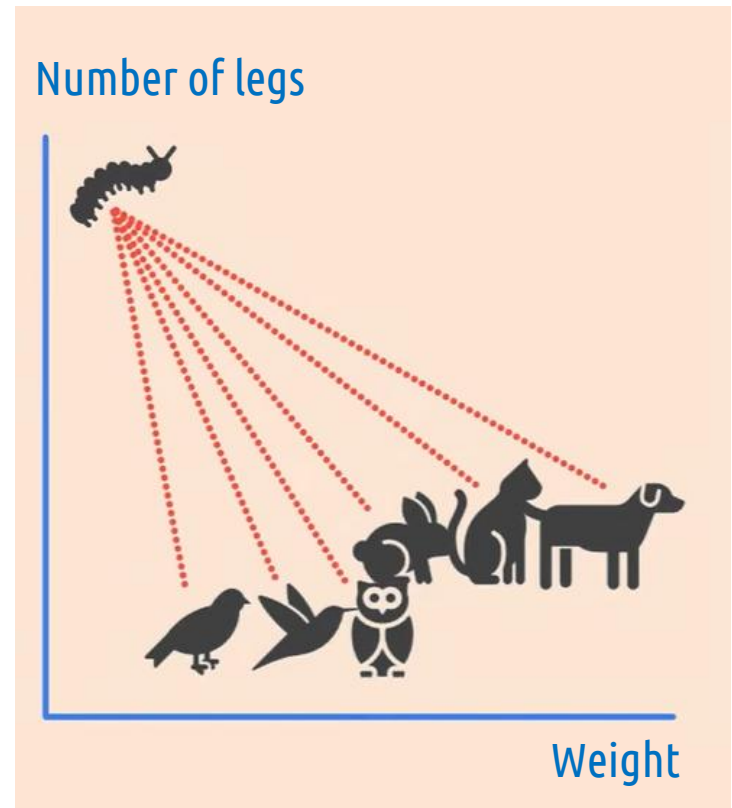
- Reducing noise (removing irrelevant features).
- Preserving critical patterns (keeping the most useful information).
- Enabling efficiency (faster computation, easier visualization).



Introduction

Anomaly Detection

- The process of identifying rare, unusual, or suspicious data points that deviate significantly from the majority of the data

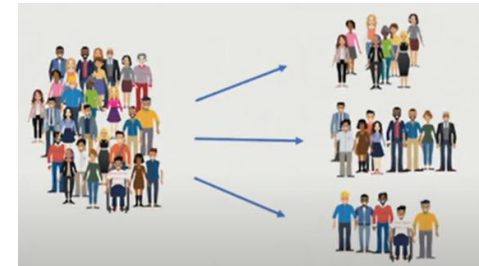


Introduction

What is Unsupervised Learning?

Applications:

- Market Segmentation,
- Image Compression
- Social Network Analysis
- Fraud Detection



100%



53%



Clustering

Introduction

What is Clustering?

- Group similar data points into clusters based on feature similarity
- No predefined labels; goal is to reveal hidden structures
- Distance/similarity metrics play a central role (e.g., Euclidean distance)
- Applications:
 - Customer segmentation
 - Document classification
 - Image grouping

Introduction

Clustering – Principal Algorithms

- **K-Means:** partitions data into K clusters based on centroid distance
- **DBSCAN:** groups data based on density; handles noise and irregular shapes
- **Hierarchical Clustering:** merges or splits clusters iteratively
- **Spectral Clustering:** uses graph Laplacian and eigen decomposition

Introduction

K-Means Clustering (Overview)

- Objective: partition n observations into K clusters
- Each observation belongs to the cluster with the nearest mean
- Assumes spherical, equally sized clusters

Introduction

Clustering – K-means

Example

- Plot the data

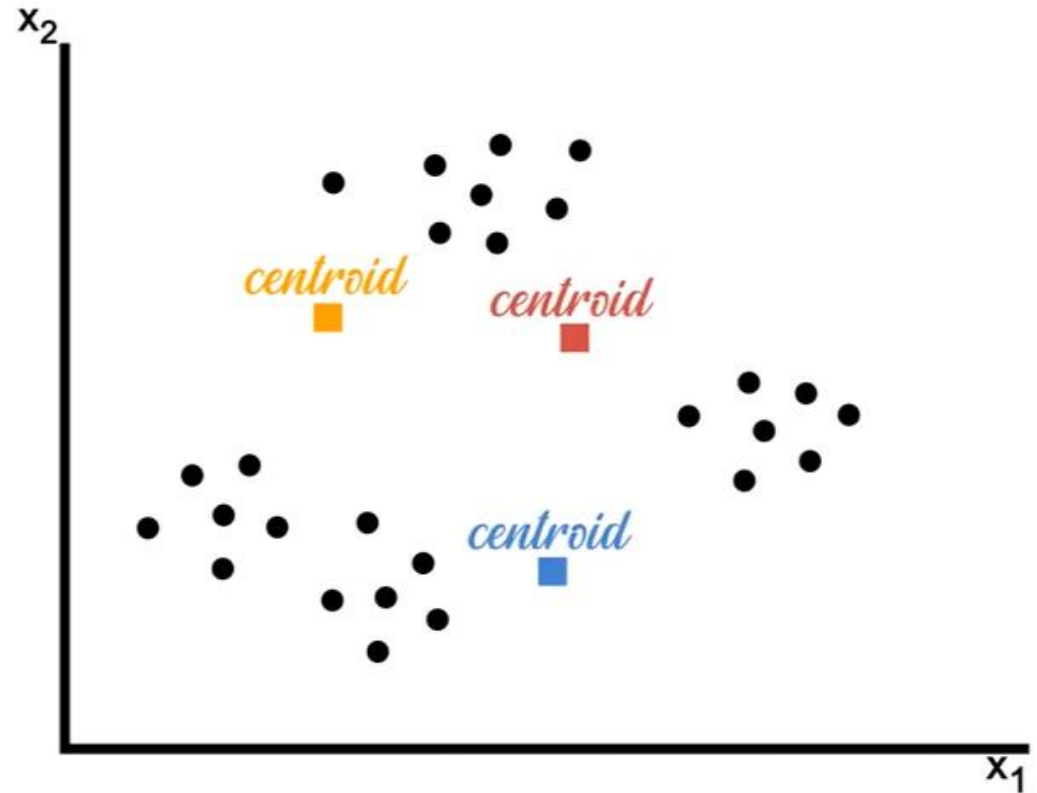


Introduction

Clustering – K-means

Example:

- Initialize 3 centroids randomly

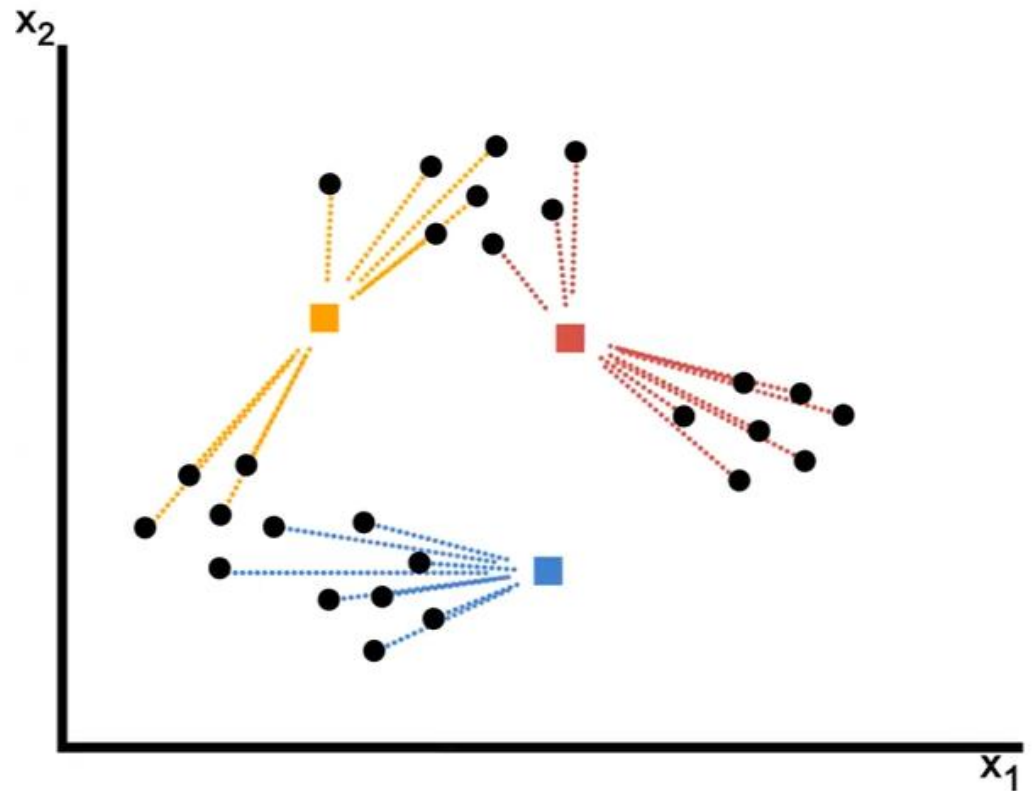


Introduction

Clustering – K-means

Example:

- Assign each point to nearest centroid

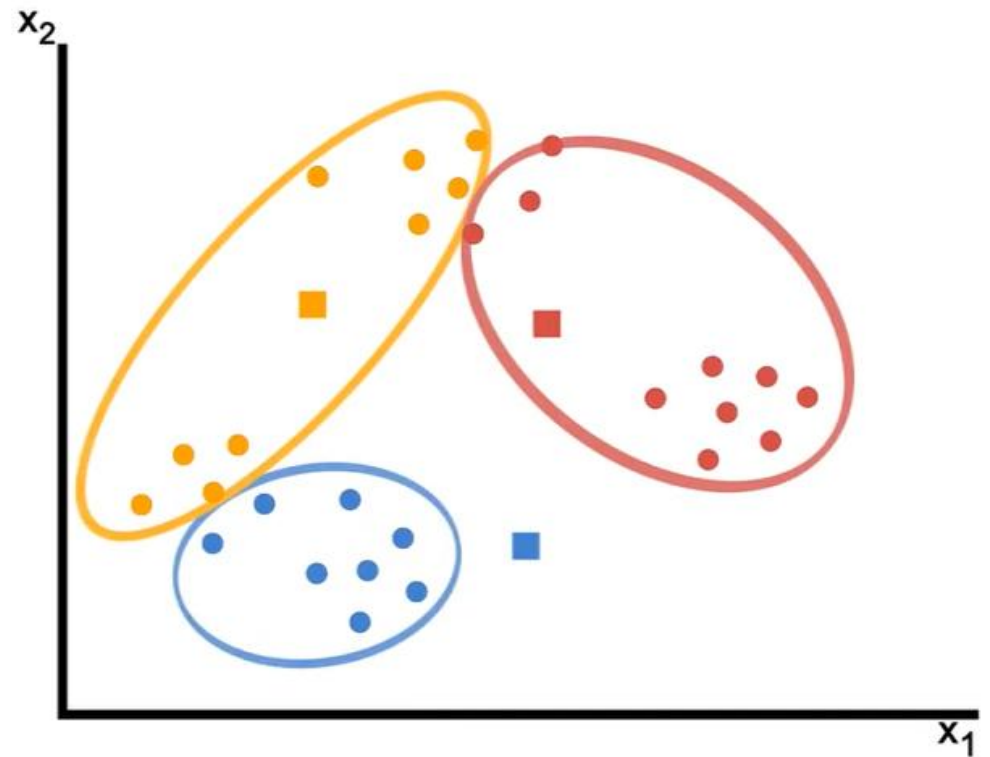


Introduction

Clustering – K-means

Example:

- Form the clusters

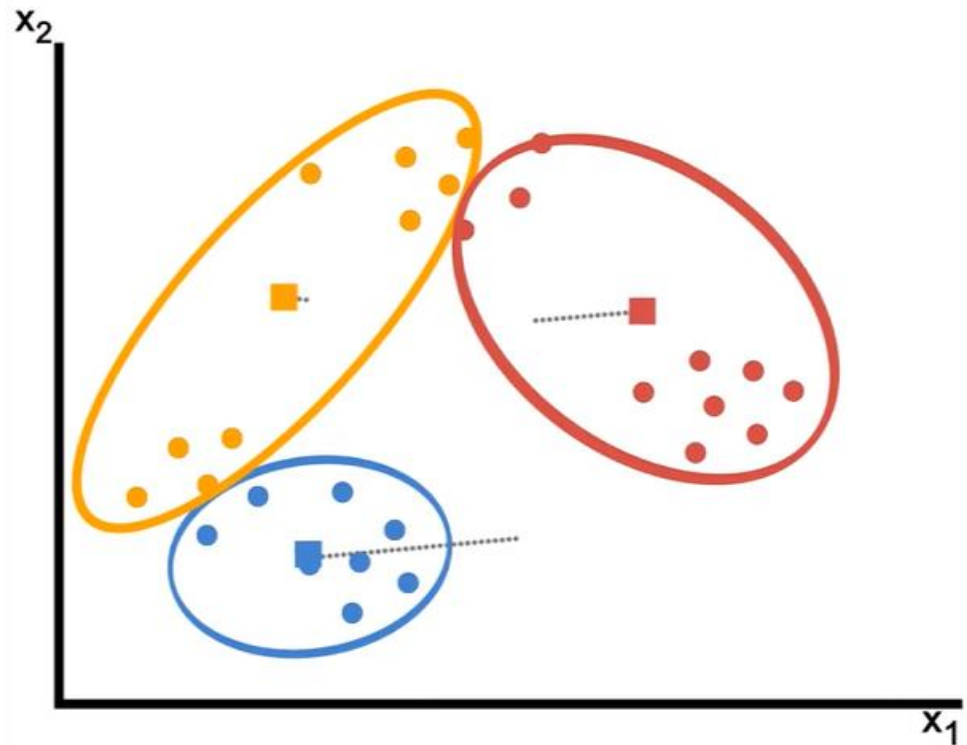


Introduction

Clustering – K-means

Example:

- Update centroids as mean of assigned points

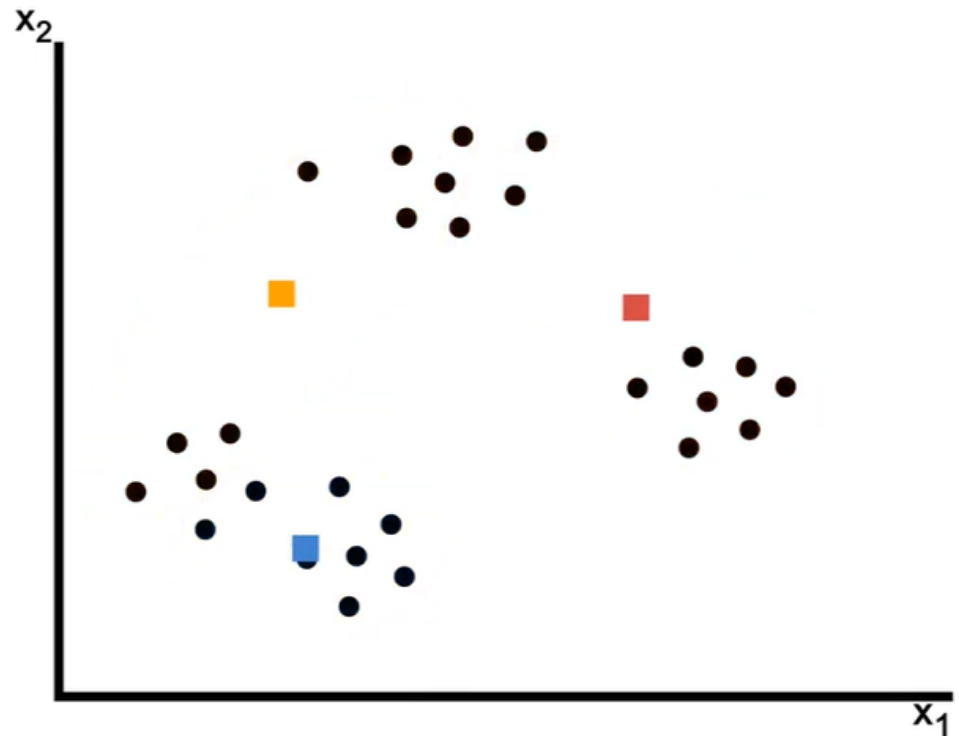


Introduction

Clustering – K-means

Example:

- Repeat the process until convergence

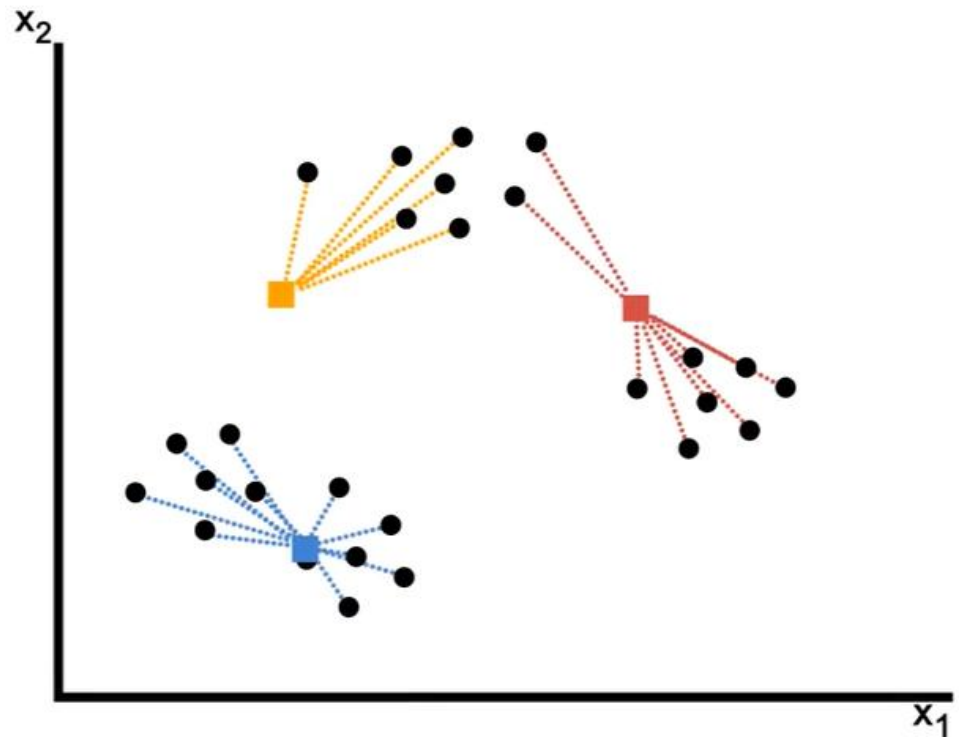


Introduction

Clustering – K-means

Example:

- Repeat the process until convergence

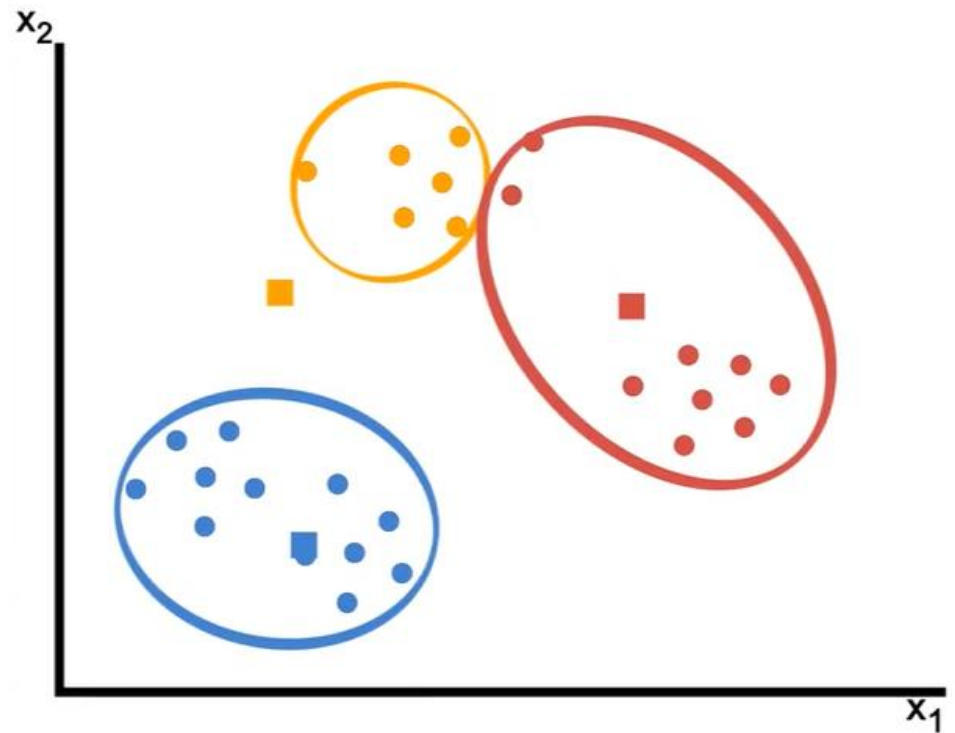


Introduction

Clustering – K-means

Example:

- Repeat the process until convergence

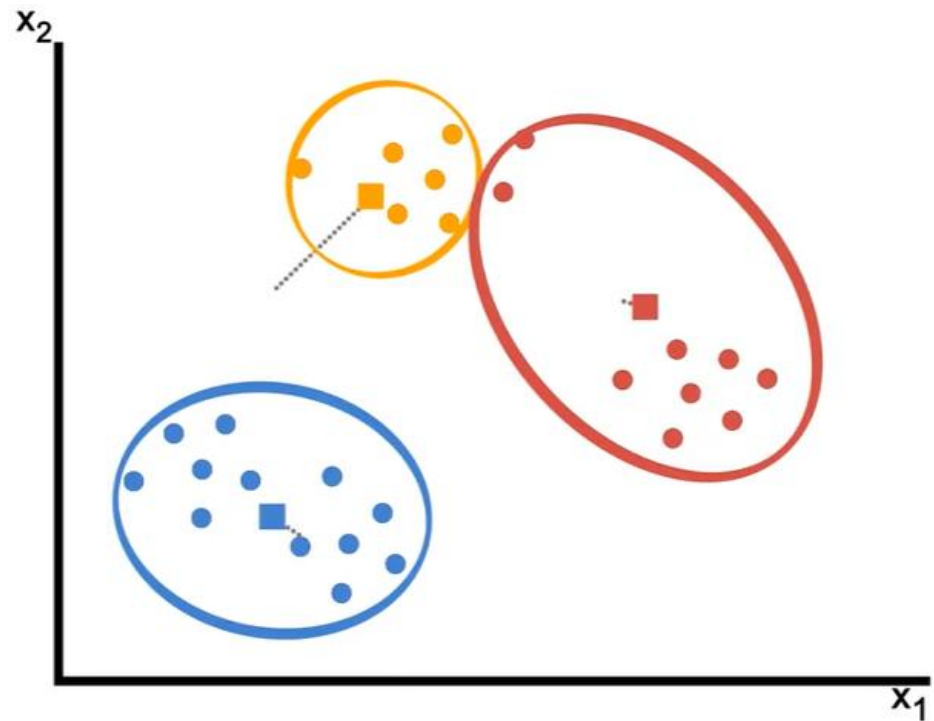


Introduction

Clustering – K-means

Example:

- Repeat the process until convergence



Introduction

K-Means (Algorithm Details)

- Input: dataset X , number of clusters K
- Steps:
 1. Initialize K centroids randomly
 2. Assign each point to nearest centroid
 3. Update centroids as mean of assigned points
 4. Repeat steps 2-3 until convergence

Introduction

K-Means (Mathematical Formulation)

- Minimize within-cluster variance:

$$J = \sum_{i=1}^n \sum_{k=1}^K r_{ik} \|x_i - \mu_k\|^2$$

Where:

- $r_{ik} = 1$ if point x_i belongs to cluster k , 0 otherwise
- μ_k : centroid of cluster k

Introduction

Evaluating Clustering (Inertia & Silhouette)

Inertia:

- sum of squared distances of samples to their closest cluster center
- Measures compactness of clusters
- Lower inertia indicates tighter clusters

Silhouette Score:

- Measures cohesion vs. separation
- Values range from -1 to 1
- High value means points are well matched within their cluster and poorly matched to others

Introduction

Choosing the Number of Clusters (K)

- Choosing an appropriate K is crucial for optimal clustering.
- Common methods:
 - Elbow Method
 - Silhouette Score
 - Gap Statistic

Introduction

Choosing the Number of Clusters (K)

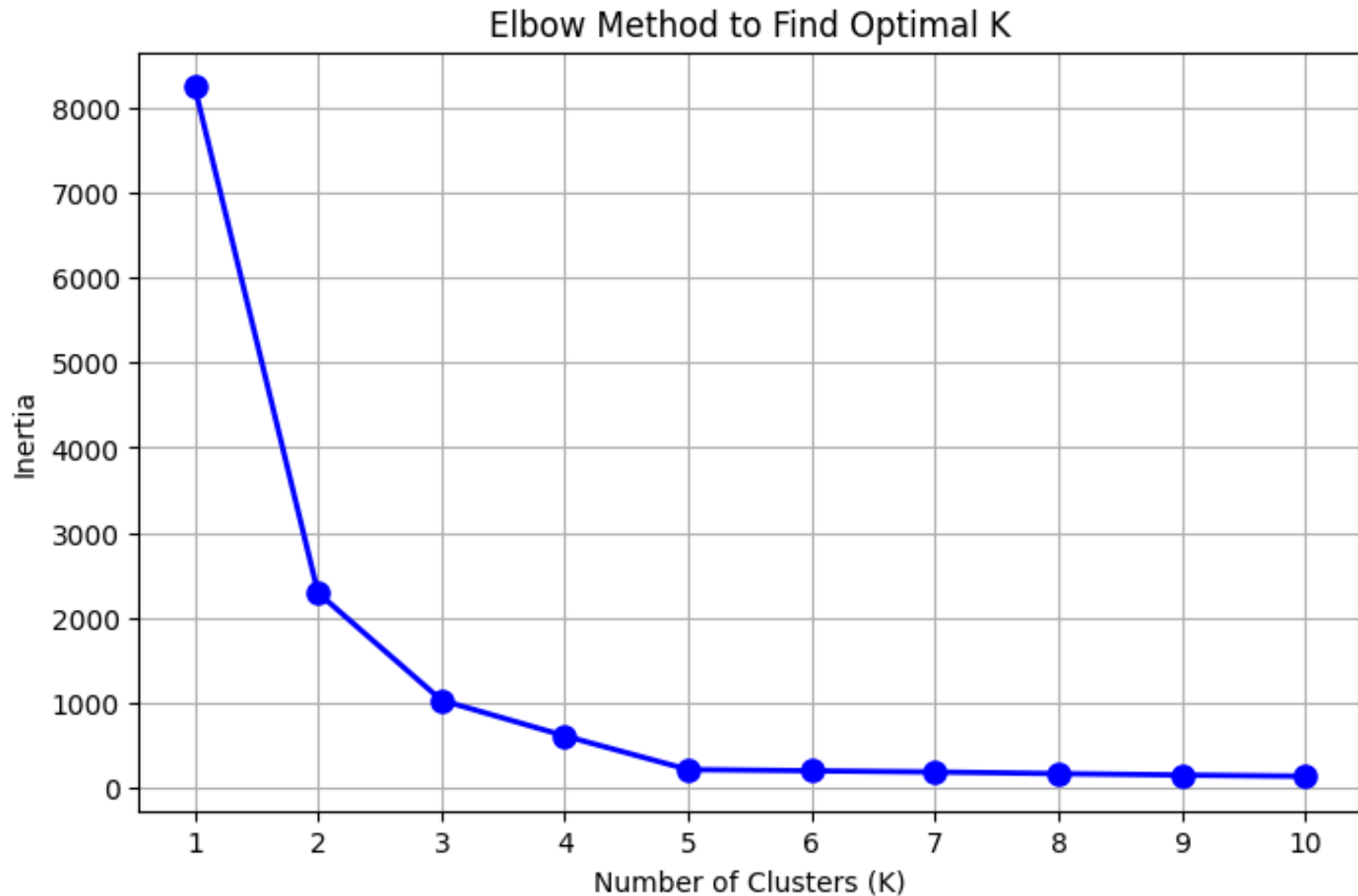
Elbow Method:

- Plot inertia vs. number of clusters.
- Look for a "knee" point where inertia starts decreasing more slowly.
- Suggests optimal trade-off between number of clusters and compactness.

Introduction

Choosing the Number of Clusters (K)

Elbow Method (Example)



Anomaly Detection

Anomaly Detection

What is it?

- Anomaly detection is the process of identifying rare, unusual, or suspicious data points that deviate significantly from the majority of a dataset. These anomalies (also called outliers) can indicate:
 - Fraud (e.g., credit card transactions)
 - Defects (e.g., faulty sensors in manufacturing)
 - Cyberattacks (e.g., unusual network traffic)
 - Biological abnormalities (e.g., rare diseases in medical data)

Anomaly Detection

Anomaly Detection Techniques

Several approaches exist, each suited to different scenarios:

- Isolation Forest → Fast, tree-based outlier isolation
- Statistical Methods (Z-Score) → Simple threshold-based detection
- Density-Based (LOF, DBSCAN) → Neighborhood density analysis
- Boundary-Based (One-Class SVM) → Learned normal-data boundaries
- ...

Anomaly Detection

Isolation Forest

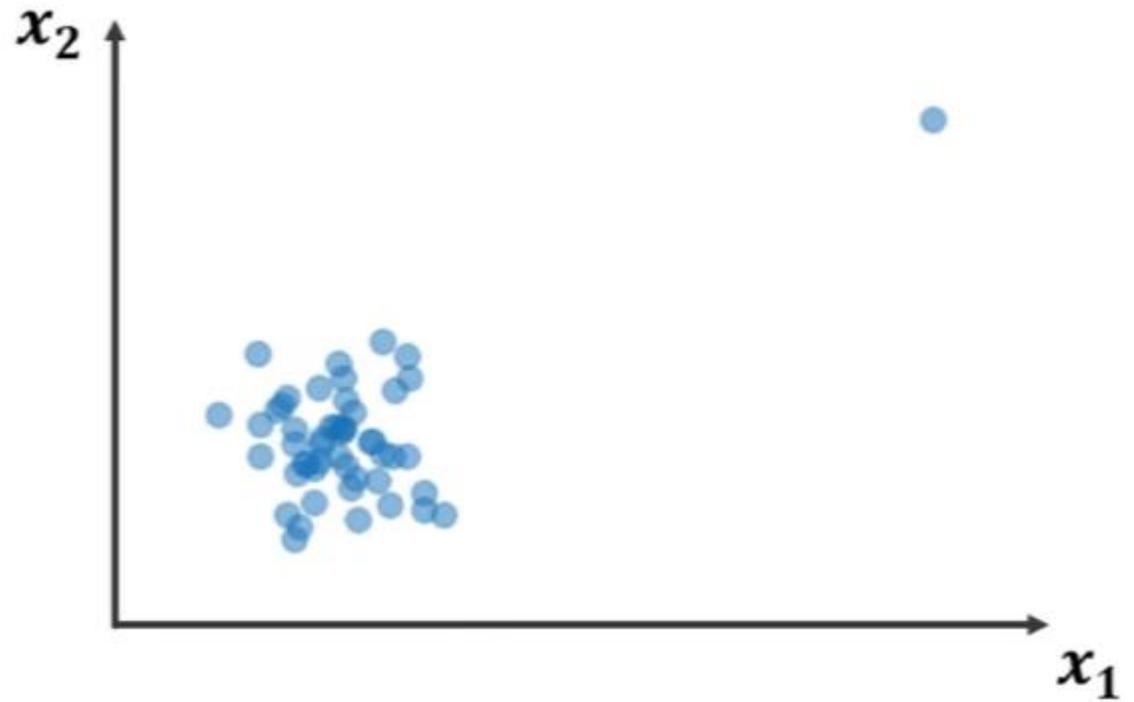
How it works:

- Builds random decision trees to "isolate" points
- Anomalies require fewer splits to separate
- Best when: You need speed and have high-dimensional data
- Limitation: May miss grouped anomalies

Anomaly Detection

Isolation Forest

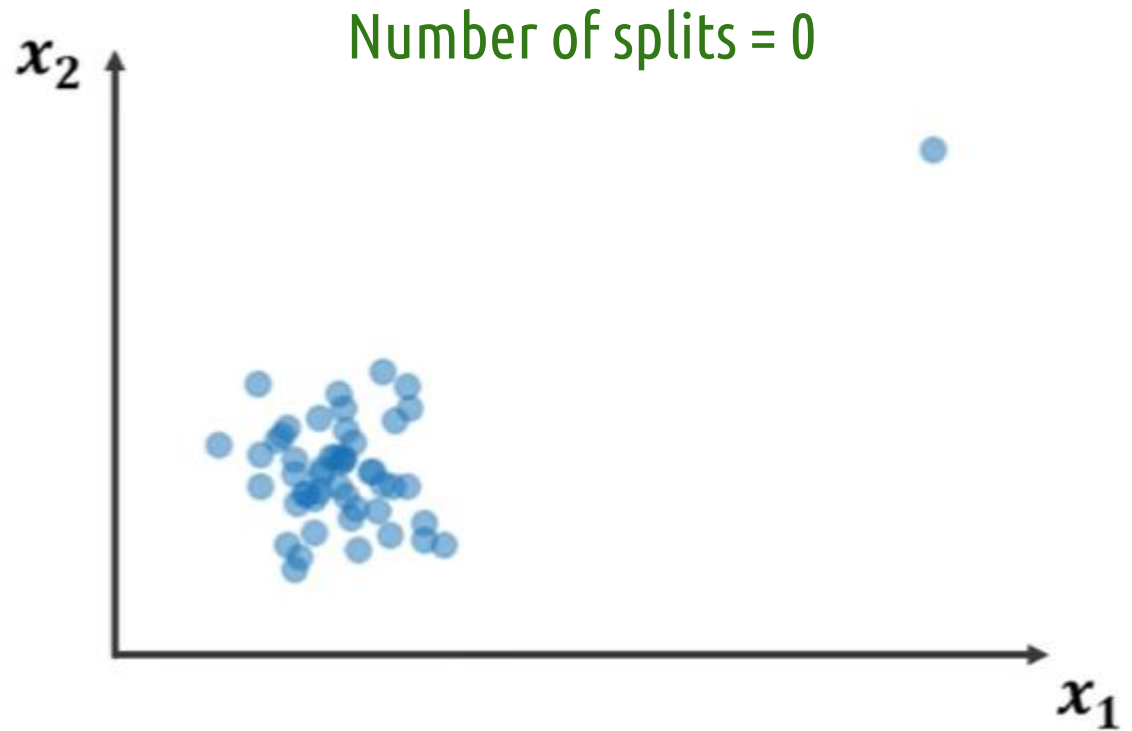
Example



Anomaly Detection

Isolation Forest

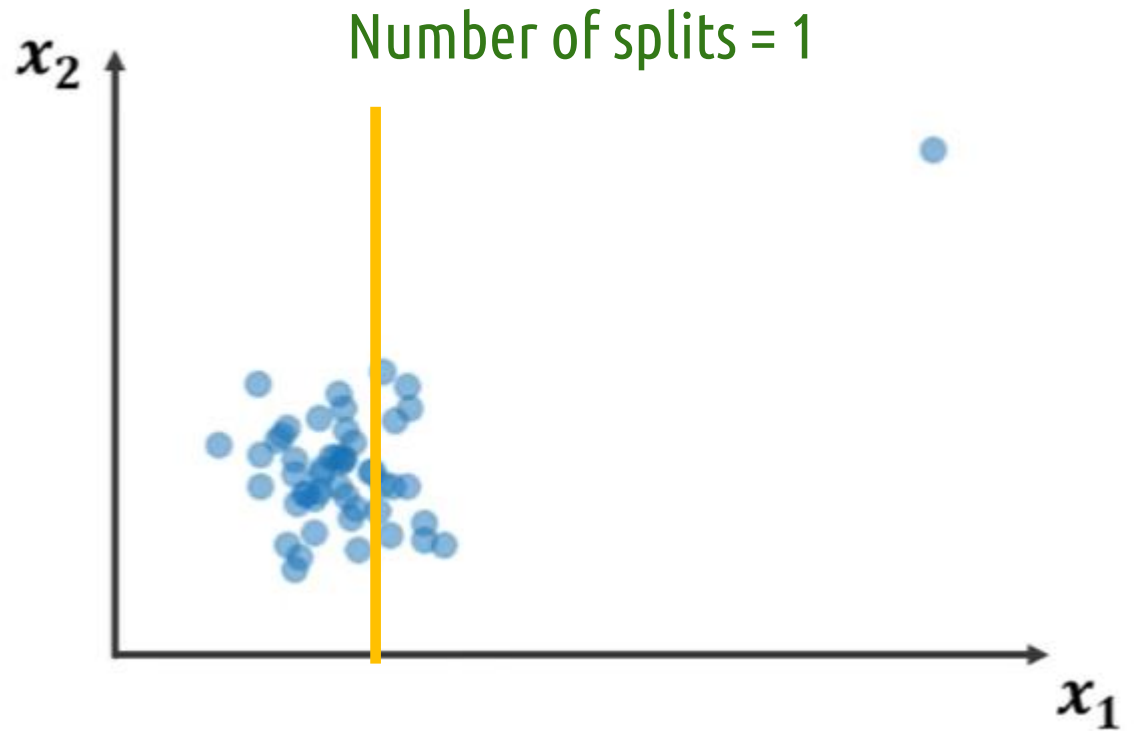
Example



Anomaly Detection

Isolation Forest

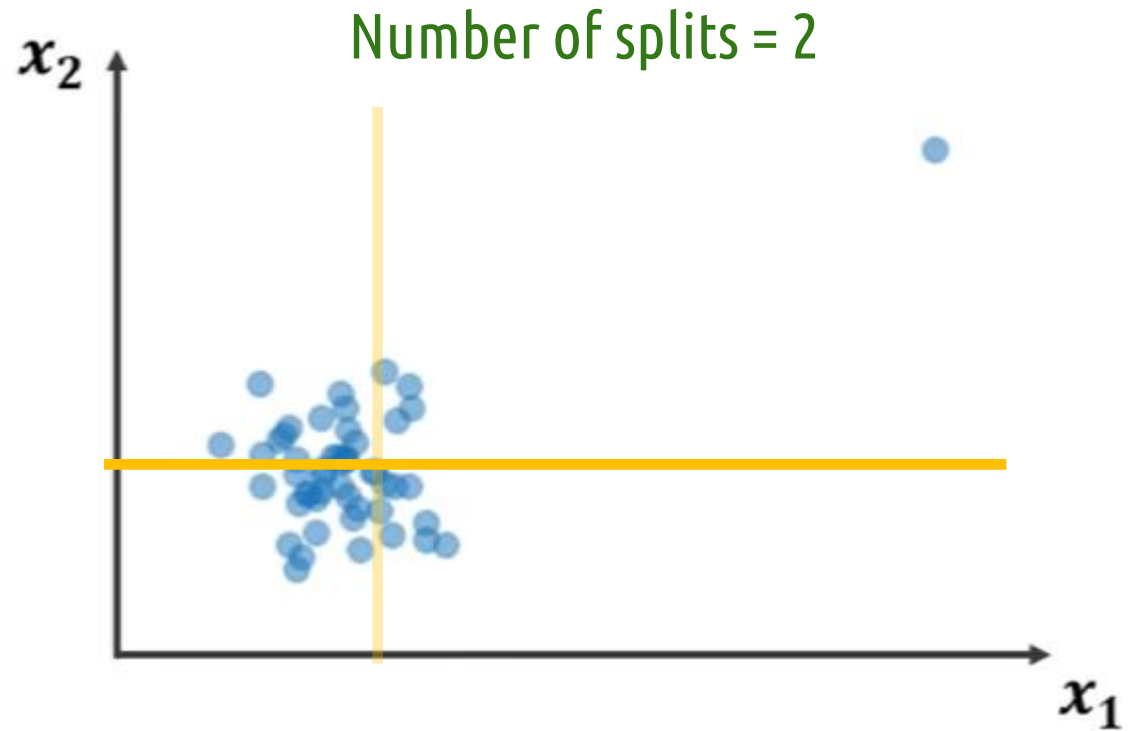
Example



Anomaly Detection

Isolation Forest

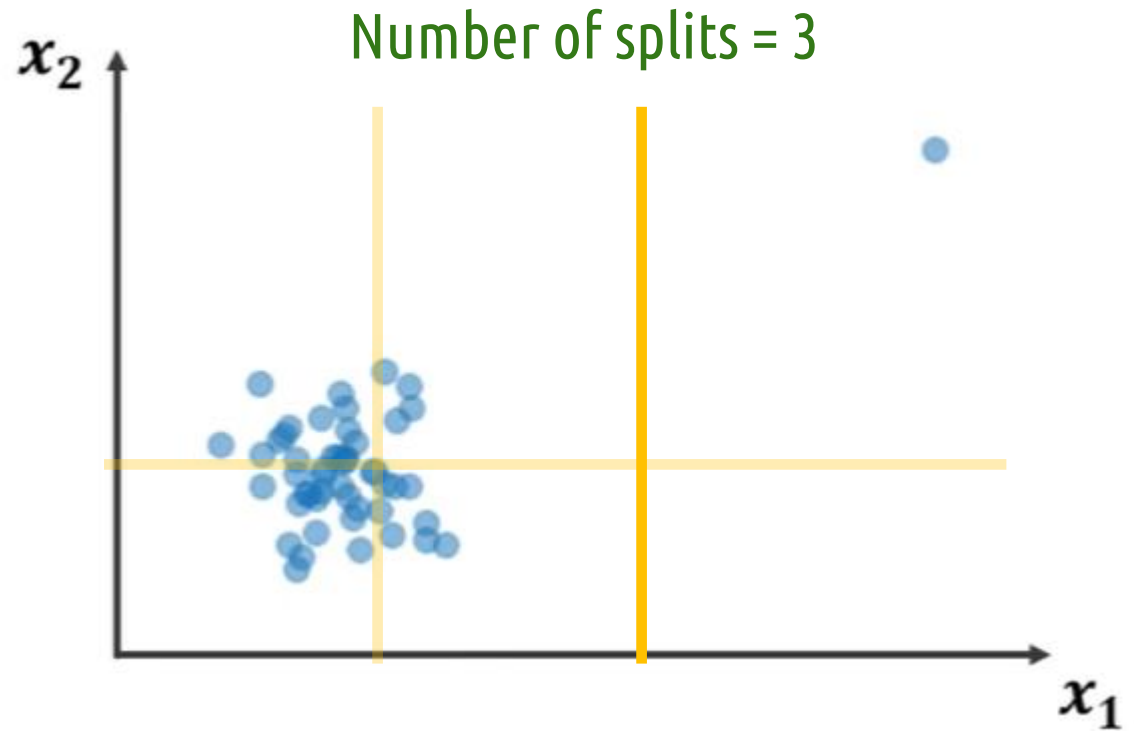
Example



Anomaly Detection

Isolation Forest

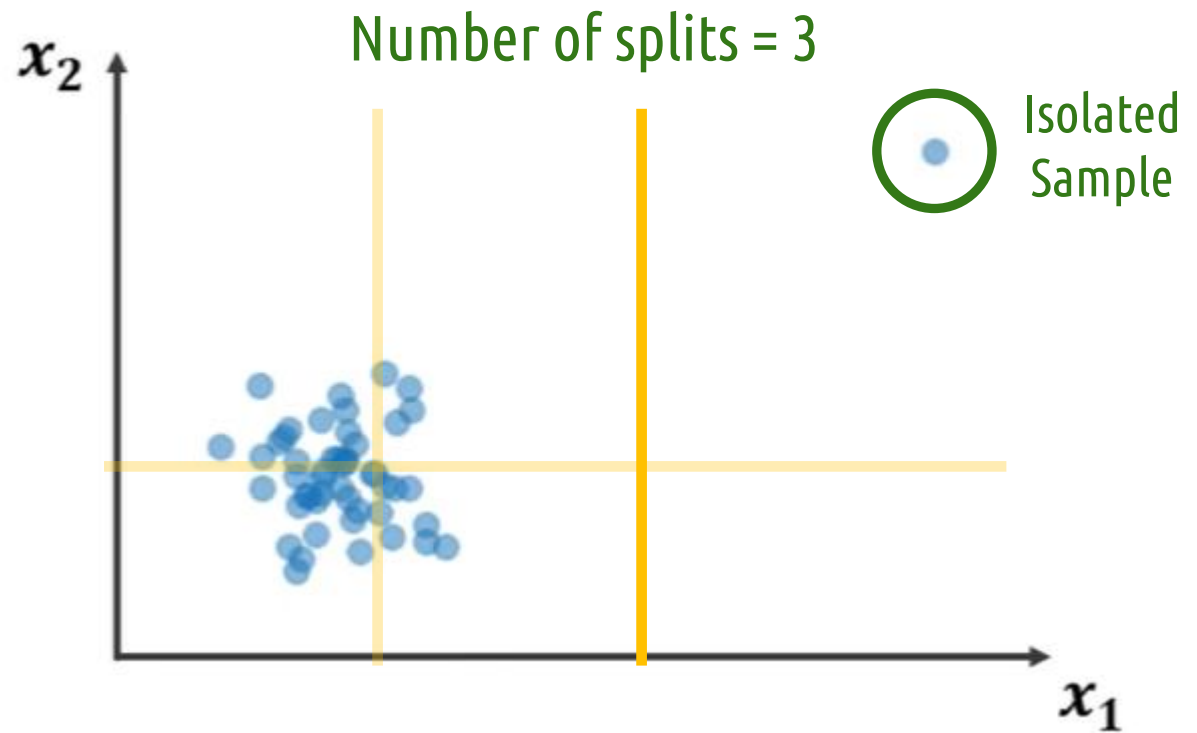
Example



Anomaly Detection

Isolation Forest

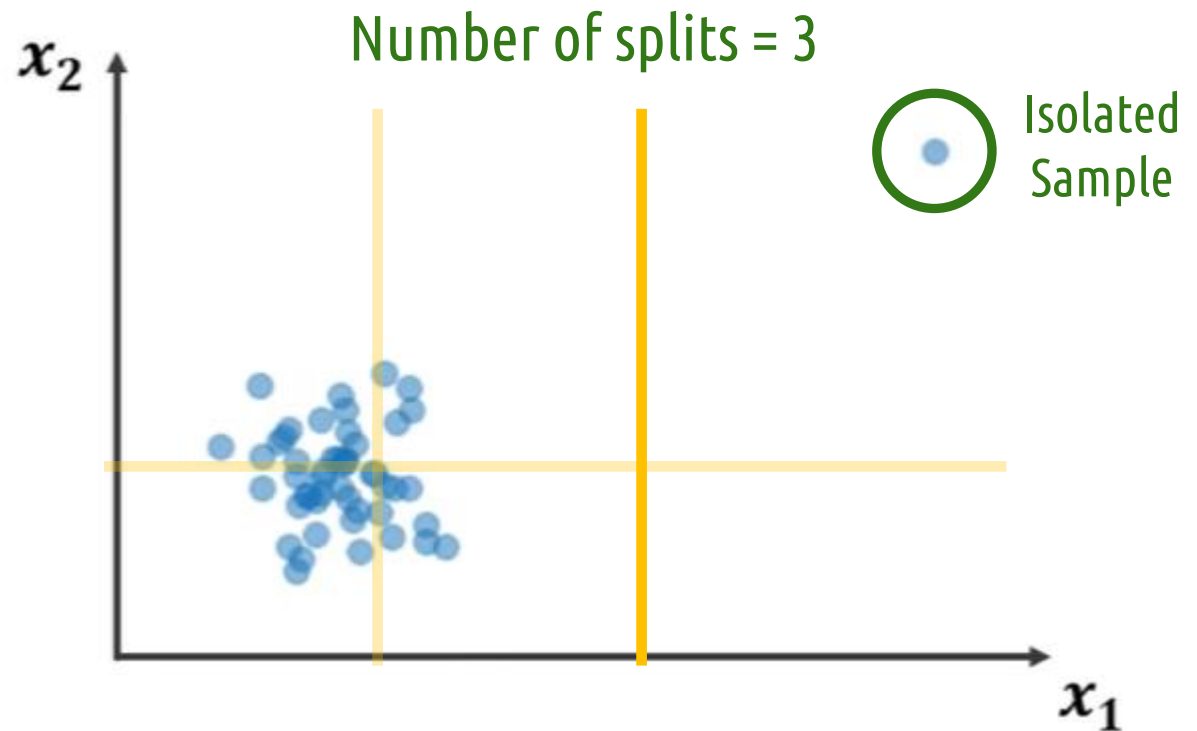
Example



Anomaly Detection

Isolation Forest

Example



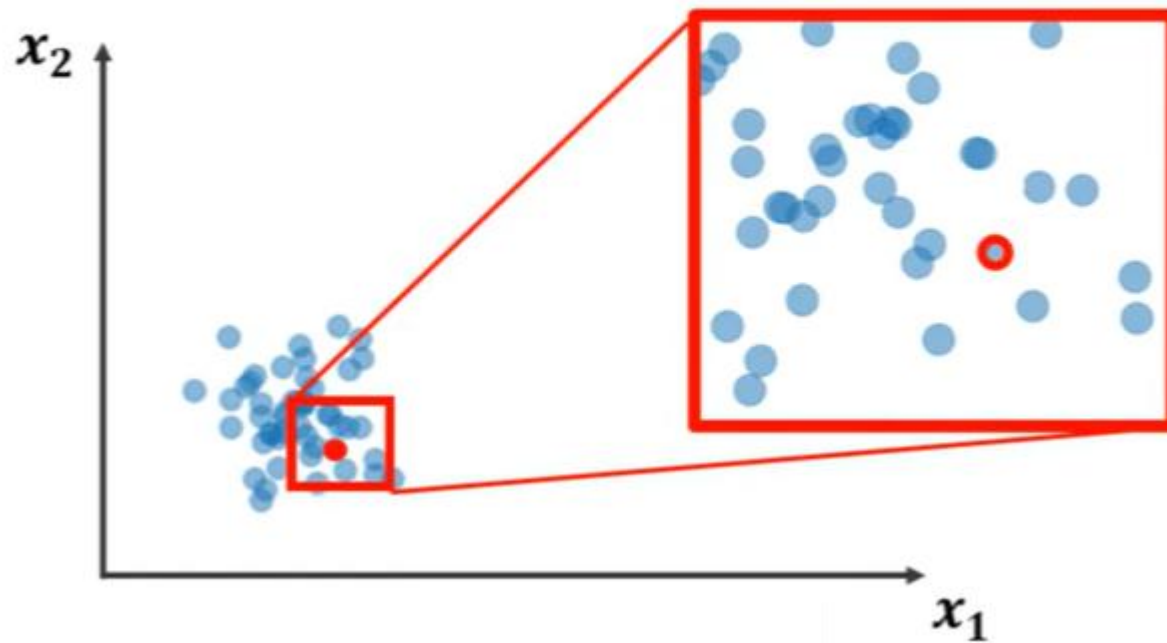
Low number of splits = High probability of anomaly

Anomaly Detection

Isolation Forest

Example

By comparison, isolating a sample lost within the mass...



Requires very high number of splits

Dimensionality Reduction

Dimensionality Reduction

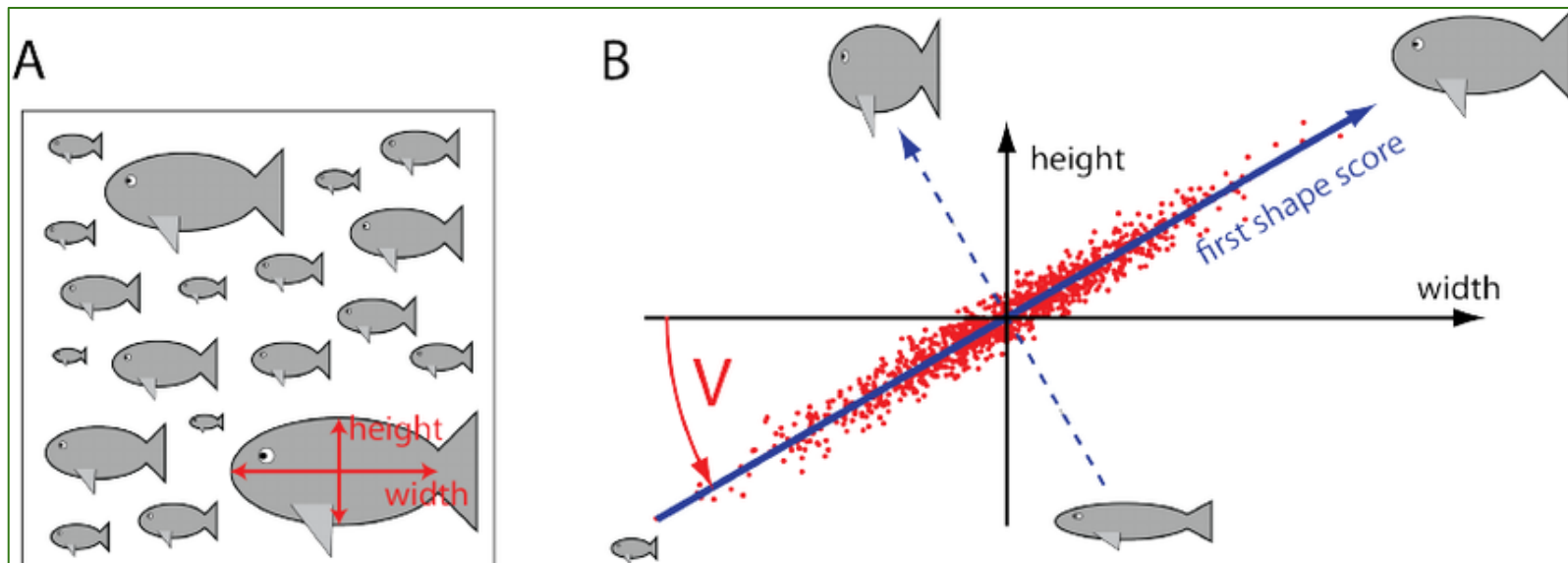
Principles

- Reduce number of features while preserving essential structure
- Removes redundancy and noise, improves visualization and learning
- **Main methods:**
 - PCA (Principal Component Analysis)
 - t-SNE, UMAP, Autoencoders (used for non-linear manifolds)

Dimensionality Reduction

PCA – Principal Component Analysis (Overview)

- Linear dimensionality reduction method
- Projects data onto orthogonal directions (principal components)
- Directions maximize variance of the data



Dimensionality Reduction

PCA – Principal Component Analysis (Overview)

Principal Steps of PCA:

- 1) Standardize the data (Optional but recommended: center and scale the features so they have mean 0 and variance 1)
- 2) Compute the covariance matrix

$$\mathbf{S} = \frac{1}{n-1} \mathbf{X}^T \mathbf{X}$$

- 3) Compute eigenvalues and eigenvectors of the covariance matrix. Solve:

$$\mathbf{S}\mathbf{w} = \lambda\mathbf{w}$$

Dimensionality Reduction

PCA – Principal Component Analysis (Overview)

Principal Steps of PCA:

- 4) Sort eigenvalues and eigenvectors
 - Sort the eigenvalues λ from largest to smallest.
 - Keep the corresponding eigenvectors w in the same order.
- 5) Select the top k eigenvectors (based on how much variance you want to keep).
- 6) Project the data onto the new space

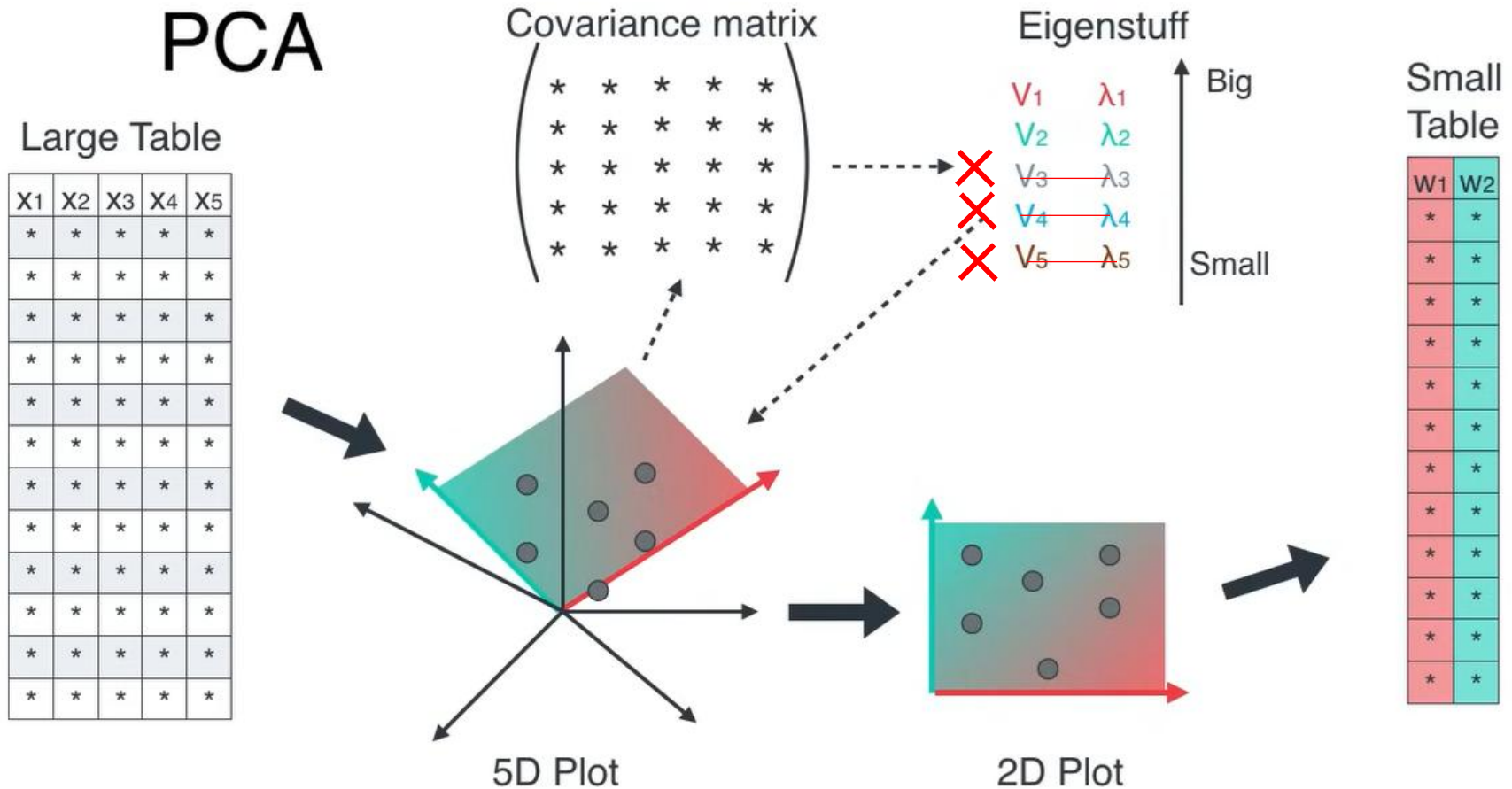
New data:

$$X_{\text{new}} = X \times W_k$$

where W_k is the matrix of the top k eigenvectors.

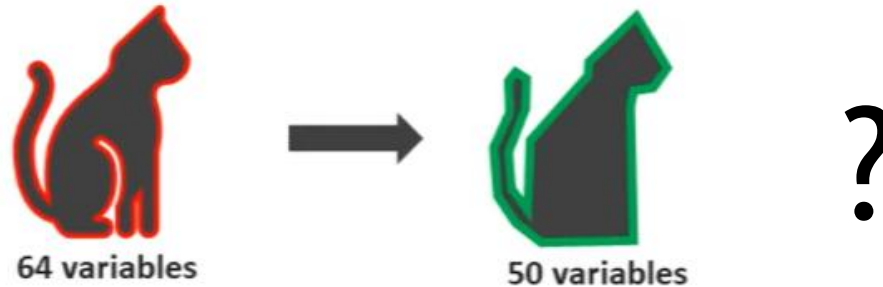
Dimensionality Reduction

PCA – Principal Component Analysis (Example)



Dimensionality Reduction

How to choose the number of components?



Variance explained threshold (most common way):

- Choose k so that the cumulative variance is above a threshold, like 95% or 99%.
- Formula:

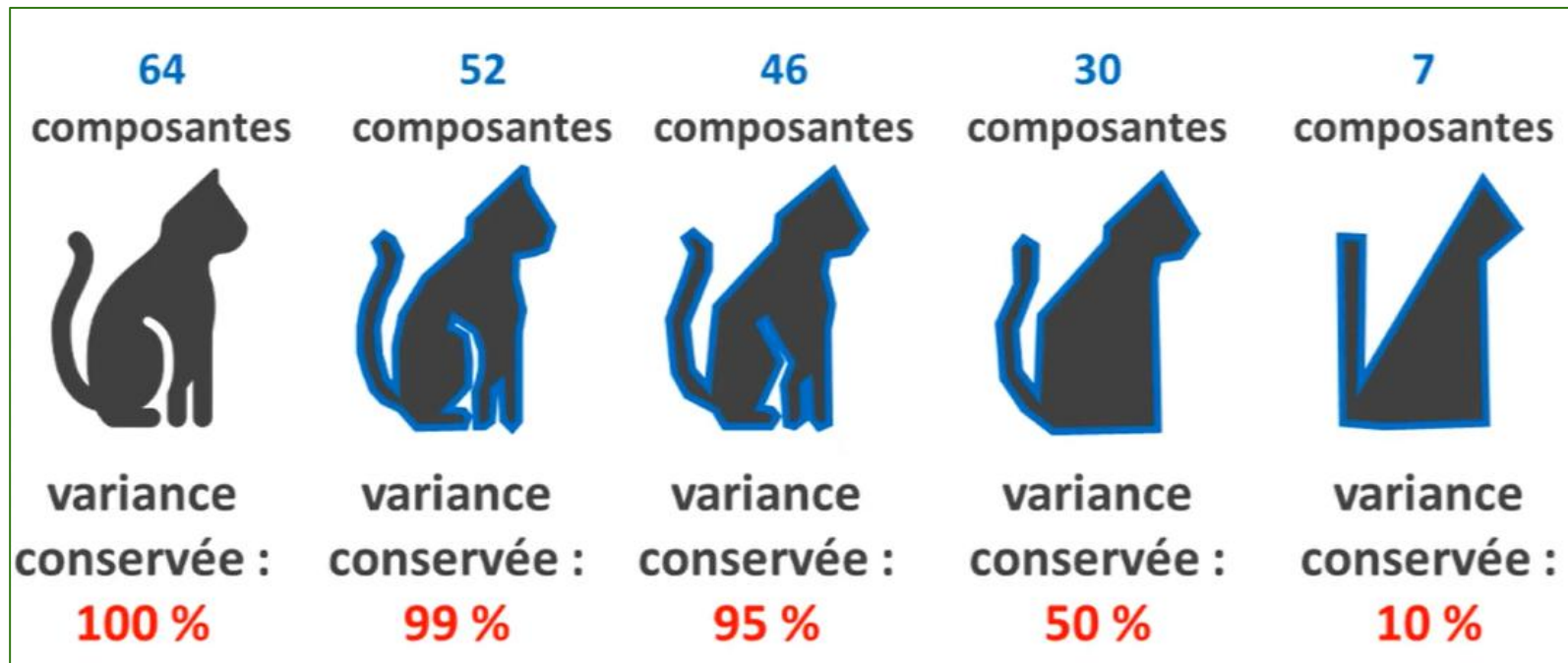
$$\text{Cumulative variance} = \frac{\sum_{i=1}^k \lambda_i}{\sum_{i=1}^n \lambda_i}$$

Here, λ_i are eigenvalues sorted decreasingly.

Dimensionality Reduction

How to choose the number of components?

- Example:



Dimensionality Reduction

How to choose the number of components?

Other methods:

- Scree plot (elbow method)
- Parallel Analysis
- Cross-validation
- ...

Reference

Reference

- Bishop, C. M. (2006). *Pattern Recognition and Machine Learning*. Springer. A comprehensive textbook with in-depth coverage of clustering and dimensionality reduction.
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- Saint-Cirgue, G. [Machine Learnia]. (2025). *Apprentissage non-supervisé avec Python* (24/30) [Video]. YouTube.
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Thank you for your attention...