Unsupervised learning

Inteligencia Artificial en los Sistemas de Control Autónomo Máster en Ciencia y Tecnología desde el Espacio

Departamento de Automática





Objectives

I. TODO

Bibliography

- TODO Bishop, Christopher M. Pattern Recognition and Machine Learning. 2nd edition. Springer-Verlag. 2011
- TODO Müller, Andreas C., Guido, Sarah. Introduction to Machine Learning with Python. O'Reilly. 2016

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Clustering

K-means, agglomerative clustering, DBSCAN and GMM

Clustering

Applications

Set of unsupervised techniques that identify groups of data (named clusters)

• No universal definition of cluster: Centroid, medoid, dense regions, etc

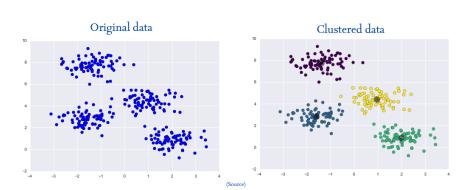
Applications

- Customer segmentation
- Data analysis
- Dimensionality reduction
- Anomaly detection
- Semi-supervised learning
- Search engines
- Image segmentation

Main algorithms

• K-means, DBScan, GMM, hierarchical clustering, EM, ...





In k-means, clusters are identified by a centroid



K-means algorithm (I)

K-means algorithm

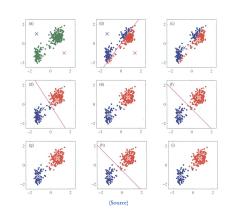
1. Set k random centroids

K-means 00000000

- 2. Assign each data point to its closest centroid
- 3. Recompute centroids
- 4. Go to 2 until no point reassignment

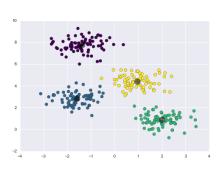
k is an hyperparameter

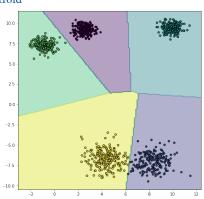
Number of clusters



K-means algorithm (II)

New data points are assigned to its closest centroid

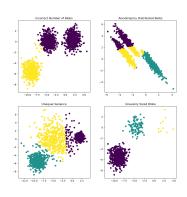




K-means limitations

K-means can fail in several conditions

- Incorrect number of clusters
- Different clusters variance
- Non-spheric clusters \Rightarrow normalization



(Source)



K-means

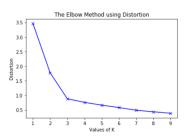
Elbow's method

Election of k

- Not a problem when domain information is available
- ... that is rarely the case

Elbow's method

- I. Select K = 1, ..., n
- 2. Visualize performance for each k
- 3. Choose K where metric stabilizes



Performance measures

- Inertia: mean squared error between each instance and its closest centroid
- Silhouette: $(b a)/\max(a, b)$, where a mean intra-cluster distance, and b is the mean nearest-cluster distance



K-means

Application: Image segmentation





(Source)



K-means

Application: Clustering for semi-supervised learning

Semi-supervised learning: Only a subset of the dataset is labeled

- Supervised and unsupervised learning
- Quite common in real-world applications (labels use to be expensive)

f1	f_2		fn	Υ
$\mathfrak{a}_{1,1}$	$\mathfrak{a}_{2,1}$	• • •	$\mathfrak{a}_{\mathfrak{n},1}$	γ1
$\mathfrak{a}_{1,2}$	$\mathfrak{a}_{2,2}$	• • •	$\mathfrak{a}_{\mathfrak{n},2}$	
$\mathfrak{a}_{1,3}$	$\mathfrak{a}_{2,3}$	• • •	$\mathfrak{a}_{\mathfrak{n},3}$	
$\mathfrak{a}_{1,4}$	$\mathfrak{a}_{2,4}$	• • •	$\mathfrak{a}_{\mathfrak{n},4}$	γ4
$\mathfrak{a}_{1,5}$	$\mathfrak{a}_{2,5}$	• • •	$\mathfrak{a}_{n,5}$	

Label propagation

- 1. Obtain k clusters
- Get a representative instance of each cluster (medoid) measuring the distance to the centroid
- 3. Label the members of each cluster with its medoid's label

K-means: Summary

Hyperparameters	Advantages	Disadvantages
	Fast	Simple shapes
k	Few hyperparameters	Determine k
	Scalable	Random initialization

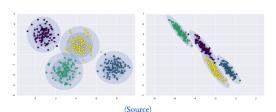
Gaussian Mixure Model (GMM) (I)

GMM is a generative clustering algorithm

Assumes data coming from a set of multidimensional gaussian distributions

GMM fits a set $\{(\phi_i, \mu_i, \sigma_i)\}_{i=1,...,k}$

- ϕ is a weight
- μ is a multidimensional mean
- σ is a covariance matrix
- k is the number of clusters (hyperparameter)





Gaussian Mixure Model (GMM) (II)

Gaussian parameters are fit with the Expectation-Maximization (E-M) algorithm

• E-M is a generalization of K-means

Expectation-Maximization algorithm

- 1. Init parameters randomly
- 2. Expectation step: Assign each instance to a cluster
 - Assignment is probabilistic
- 3. Maximization step: Update cluster parameters
 - Each cluster is updated using all the data
 - Instances contribution to a cluster parameters is weighted by the probability that it belongs to it
- 4. Go to 2

GMM can be seen as a fuzzy clustering algorithm



Gaussian Mixure Model (GMM) (III)

Gaussian parameters are fit with the Expectation-Maximization (E-M) algorithm

• E-M is a generalization of K-means

Expectation-Maximization algorithm

- 1. Init parameters randomly
- 2. Expectation step: Assign each instance to a cluster
 - Assignment is probabilistic
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 - Each cluster is updated using all the data
 - Instances contribution to a cluster parameters is weighted by the probability that it belongs to it
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GMM can be seen as a fuzzy clustering algorithm

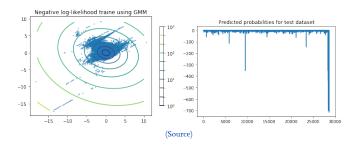


Gaussian Mixure Model (GMM) (IV)

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GMM provides a probability of an instance to belong to a cluster

- This can be used to detect anomalies
- Just assign a probability threshold





GMM: Summary

Hyperparameters	Advantages	Disadvantages
Number of clusters	Probabilistic clustering	Number of clusters
Covariance matrix type	Generative model	Gaussian data
	Anomaly detection	Sensitive to outliers



Other clustering algorithms DBSCAN (I)

DBSCAN: Density-Based Spatial Clustering of Applications with Noise

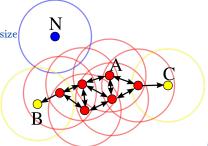
- Identifies high density regions (dense regions) in feature space
- Asumtion: Clusters form dense regions separated by empty areas

Hyperparameters

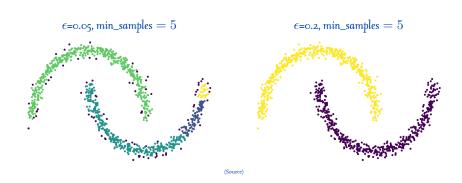
- ullet ϵ : Radius of a neighborhood
- min_samples: Minumun cluster size

Type of points

- Core instance
- Outliers



(Source)





DBSCAN: Summary

Hyperparameters	Advantages	Disadvantages
ϵ	No explicit number of clusters	Slower than K-means
min_samples	Scales relatively well	Clusters with different densities
	Almost deterministic	
	Robust to outliers	
	Anomaly detection	



Agglomerative clustering (I)

Agglomerative clustering

- 1. Initially, each instance forms a cluster
- 2. Merge the two most similar clusters according to a metric
- 3. Repeat 2 until a stop criterion is satisfied









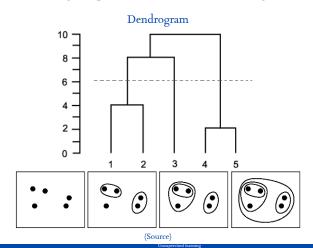
We need a similarity measure between two clusters

- Ward: Minimizes variance within merged clusters. Leads to equally sized clusters
- Average: Minimizes average distances between their points
- Complete: Minimizes maximun distance between their points



Agglomerative clustering (II)

Agglomerative clustering is a special case of hierarchical clustering





Agglomerative clustering: Summary

Hyperparameters	Advantages	Disadvantages
	Complex shapes Hierarchical clustering	
	Therarchical clustering	



Anomaly detection

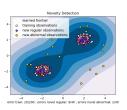
Two related concepts

Outlayer detection and novelty detection

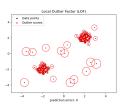
Adaptation of clustering and classification algorithms

• PCA, GMM, autoencoders, etc

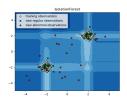
One-Class SVM



LOF



Isolation Forest



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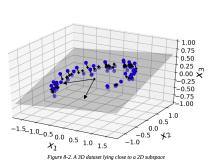
PCA and manifold learning



Main approaches for dimensionality reduction (I)

Two main approaches to dimensionality reduction: Projection and manifold learning

Projection



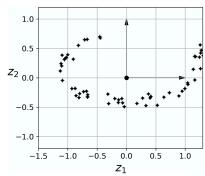


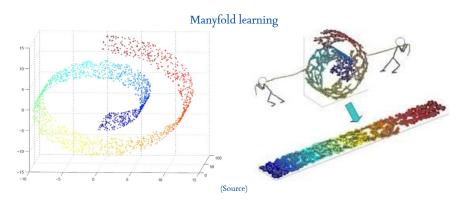
Figure 8-3. The new 2D dataset after projection

(Source)





Main approaches for dimensionality reduction (II)



Manifold learning algorithms

 Isomap, T-distributed Stochastic Neighbor Embedding (t-SNE), Multi-dimensional Scaling (MDS), Locally Linear Embedding (LLE), ...



Principal Components Analysis (I)

Dimensionality reduction transforms data into more convenient representations

- Reduce data dimensionality
- Visualize multidimensional data

Main algorithms



Principal Components Analysis (I)

Dimensionality reduction transforms data into more convenient representations

- Reduce data dimensionality
- Visualize multidimensional data

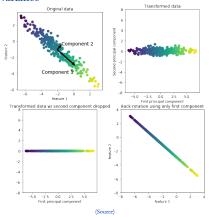
Main algorithms

- Isomap
- T-distributed Stochastic Neighbor Embedding (t-SNE)
- Principal Components Analysis (PCA)



Principal Components Analysis (II)

PCA maximizes data variance





Principal Components Analysis (III)

Example: Hand-written digits recognition

- Images of hand-written digits
- 8x8 images (64 dimensions)
- 10 digits
- Classification problem

