Unsupervised learning

Aprendizaje Automático para la Robótica Máster Universitario en Ingeniería Industrial

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

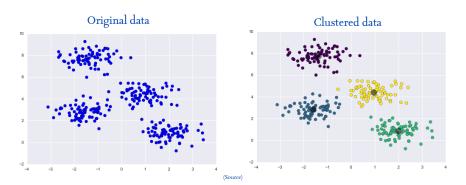
- Customer segmentation
- Data analysis
- Dimensionality reduction
- Anomaly detection
- Semi-supervised learning
- Search engines
- Image segmentation

Main algorithms

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



Overview



In k-means, clusters are identified by a centroid

K-means algorithm (I)

K-means algorithm

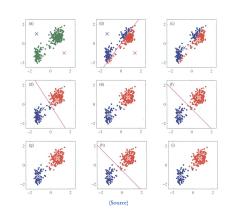
1. Set k random centroids

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

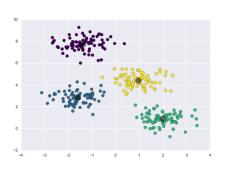
• Number of clusters

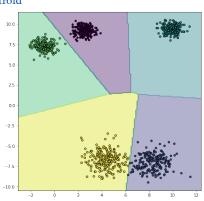


K-means algorithm (II)

K-means 000000000

New data points are assigned to its closest centroid

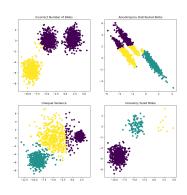




K-means limitations

K-means can fail in several conditions

- Incorrect number of clusters
- Different clusters variance
- Non-spheric clusters ⇒ normalization



(Source)



Elbow's method

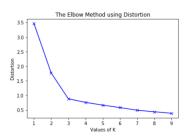
Election of k

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

K-means

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



Application: Image segmentation





(Source)



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: Scikit-learn

TODO: SCikit-Learn



K-means: Summary

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



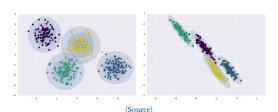
Gaussian Mixure Model (GMM) (I)

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
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GMM can be seen as a fuzzy clustering algorithm

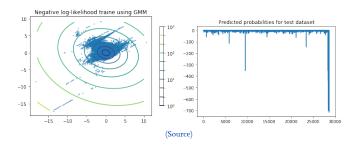


Gaussian Mixure Model (GMM) (IV)

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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





DBSCAN: Scikit-learn

TODO: Scikit-Learn



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

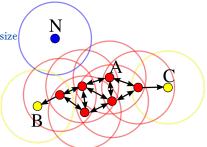
- Identifies high density regions (dense regions) in feature space
- Asumtion: Clusters form dense regions separated by empty areas

Hyperparameters

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

Type of points

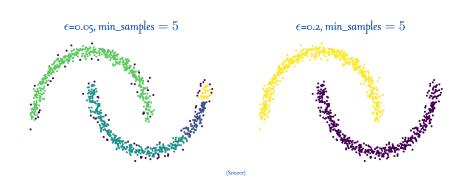
- Core instance
- Outliers



(Source)



Other clustering algorithms DBSCAN (II)





DBSCAN: Scikit-learn

TODO: Scikit-Learn



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

- I. 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









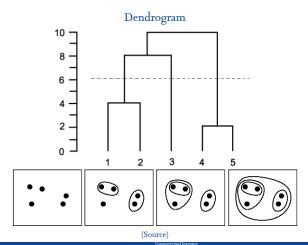
We need a similarity measure between two clusters

- 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: Scikit-Learn

TODO: SCikit-Learn



Agglomerative clustering: Summary

Hyperparameters	Advantages	Disadvantages
	Complex shapes Hierarchical clustering	
	Therarchical clustering	



Anomaly detection

Anomaly detection

TODO



PCA and manifold learning



Main approaches for dimensionality reduction

Dimensionality reduction transforms data into more convenient representations

- Reduce data dimensionality
- Visualize multidimensional data

Projection

TODO

Manyfold learning

TODO



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 (I)

Dimensionality reduction transforms data into more convenient representations

- Reduce data dimensionality
- Visualize multidimensional data

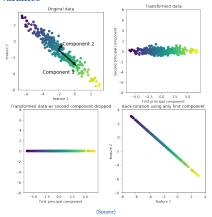
Main algorithms

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



Principal Components Analysis (II)

PCA maximizes data variance





Dimensionality reduction

Principal Components Analysis (III)

Example: Hand-written digits recognition

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

