

Lecture 4: Scalable k-means Clustering

COM6012: Scalable ML by Haiping Lu

YouTube Playlist: https://www.youtube.com/c/HaipingLu/

Week 4 Contents / Objectives

Introduction to Cluster Analysis

• k-means Clustering

• Scalable *k*-means

• k-means in Spark & Limitations

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Introduction to Cluster Analysis

• k-means Clustering

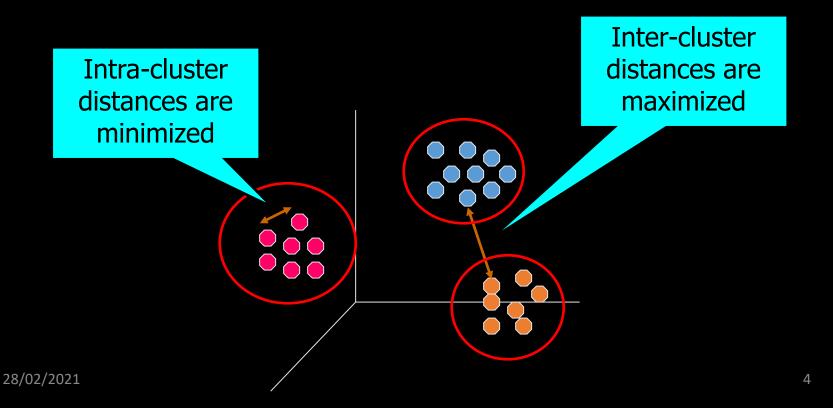
• Scalable *k*-means

k-means in Spark & Limitations

Cluster Analysis

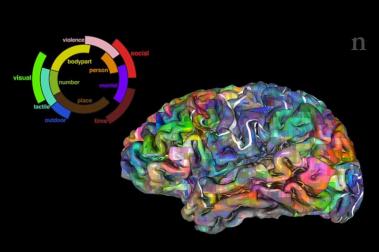
Output	Supervised	Unsupervised
Discrete	Classification	Clustering
Continuous	Regression	Dimensionality Reduction

- "Classification without labelled data"
- "Dimensionality reduction to 1 (cluster index)"



Cluster Analysis

- Divide data into clusters that are meaningful/useful
- Clusters: pseudo-classes
- Key for <u>exploratory data analysis</u> in many areas
 - Brain parcellation, social network analysis, customer segmentation, patient stratification, drug discovery, ...





brain-video2.jpg (1014×570) (wp.com)

https://miro.medium.com/max/114 5/1*YjvoSe-hGaPpbWi5fDqV2w.png segmentation3.png (586×236) (visionedgemarketing.com)

Anomaly/Outlier Detection

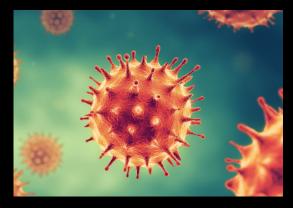
- Anomalies: data points very different from others
- Applications: credit card fraud, network intrusion, unknow virus, eco disturbance, unusual symptom, ...



financial protection credit card fraud img1.jpg (512×347) (traderdefenseadvisory.com)



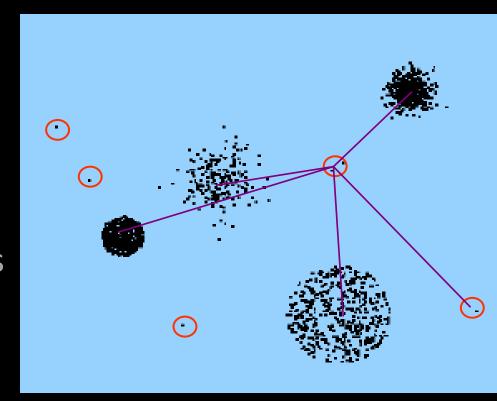
<u>Network-Intrusion-Detection-and-</u> Prevention-.jpg (1600×962) (wp.com)



newseventsimage 1613554577167 mainne ws2012 x1.jpg (700×484) (imperial.ac.uk)

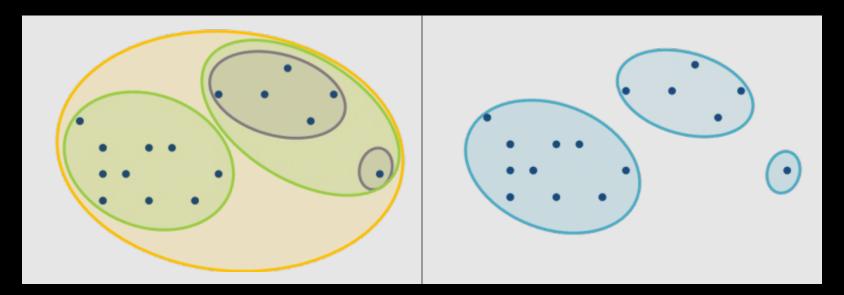
Clustering-based Anomaly Detection

- Cluster the data points
- Those in small clusters
 → candidate outliers
- Compute the distance between candidate points and non-candidate clusters
- Candidate points far from all other non-candidate points → outliers

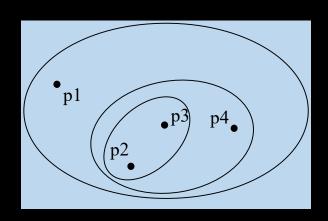


Hierarchical vs Partitional Clustering

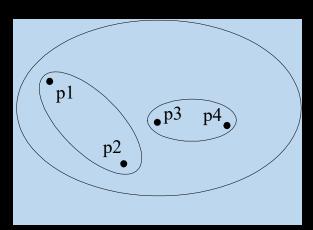
- Hierarchical: nested clusters as a hierarchical tree
 - Each node (cluster) in the tree (except for the leaf nodes) is the union of its children (subclusters)
 - The root of the tree → the cluster containing all data points
- Partitional: non-overlapping clusters
 - Each data point is in exactly one cluster



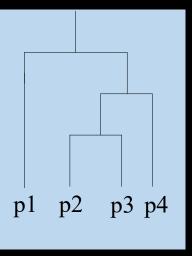
Hierarchical Clustering



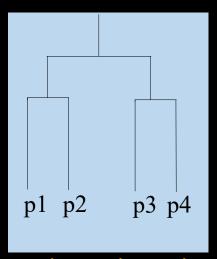
Traditional Hierarchical Clustering



Non-traditional Hierarchical Clustering



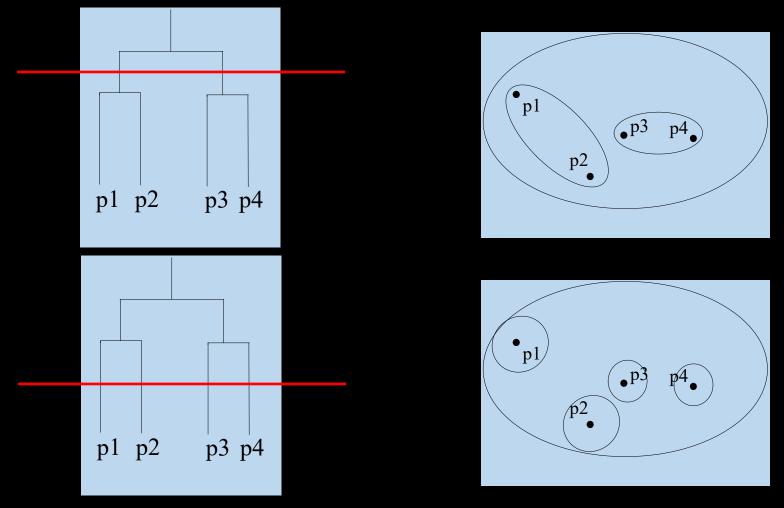
Traditional Dendrogram



Non-traditional Dendrogram

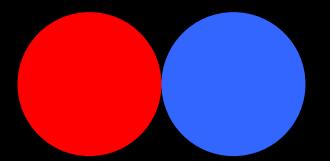
Hierarchical -> Partitional

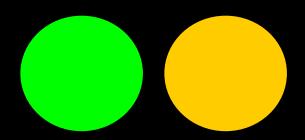
 Hierarchical = a sequence of partitional clustering cutting the hierarchical tree at a particular level



Centre/Prototype-based Clusters

- Data points in a cluster are closer (more similar) to the centre of that cluster than to the centre of any others
- Centre
 - Centroid: the average of all the points in a cluster.
 - Medoid: the most representative point of a cluster (e.g., categorical)

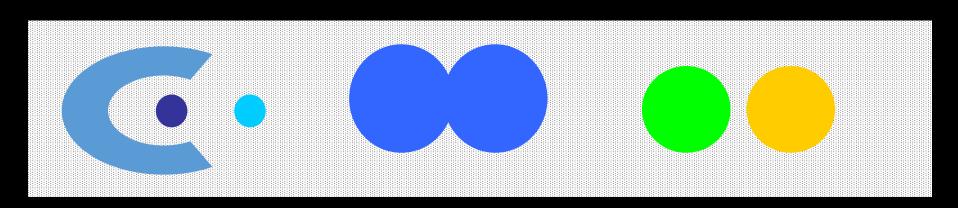




4 centre-based clusters

Density-based Clusters

- Cluster: a dense region of points separated by lowdensity regions from other regions of high density
- Used when the clusters are irregular/intertwined, and when noise and outliers are present



6 density-based clusters

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Scalable k-means

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k-means Clustering

- A centre-based, partitional clustering approach
- Input: a set of n data points $X = \{x_1, x_2, ..., x_n\}$ and the number of clusters k
- For a set $C=\{c_1, c_2, ..., c_k\}$ of cluster centres, define the Sum of Squared Error (SSE) as:

$$SSE_X(C) = \sum_{x \in X} d(x, C)^2$$

d(x,C): distance from x to the closest centre in C

Goal: find C centres minimising SSE_X(C)

Lloyd Algorithm for k-means

- Start with k centres $\{c_1, c_2, ..., c_k\}$ chosen uniformly at random from data points
- Assign clusters and compute centroids till convergence
 - Alternating optimisation again, similar to ALS
- Limitations
 - Many iterations to converge
 - Sensitive to initialisation
 - Random initialisation can get two centres in the same cluster >> stuck in a local optimum (example on next slide)

Initialisation > Stuck

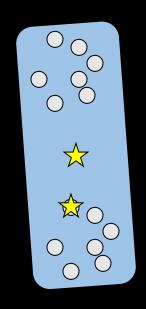






Figure credited to David Arthur

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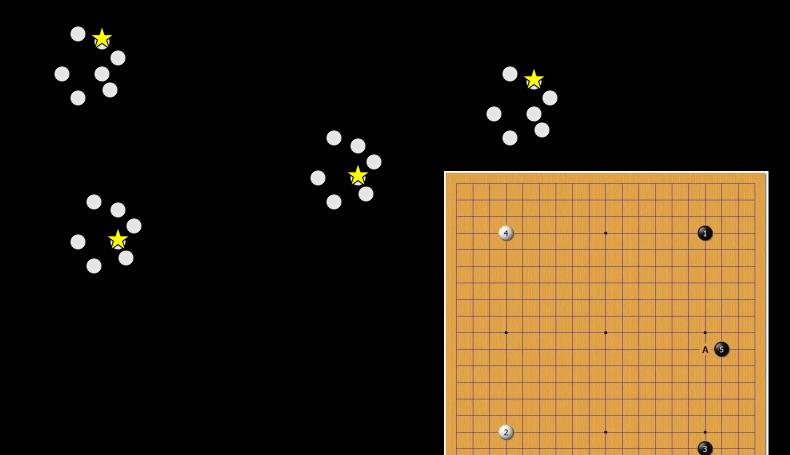
k-means++ [Arthur et al. '07]

- Key idea: spread out the centres
- Choose the first centre c_1 uniformly at random
- Repeat for $2 \le i \le k$:
 - Choose c_i to be equal to a data point x_0 sampled from the distribution:

$$\frac{d(x_0, C)^2}{SSE_X(C)} \propto d(x_0, C)^2$$

- Reminder: d(x,C) = distance from x to the closest centre in C
- Theorem: $O(\log k)$ -approximation to the optimum

k-means++ Initialisation



k-means++ $\rightarrow k$ -means||

- *k*-means++ limitations
 - Needs k passes over the data for initilisation
 - In big data applications, k is typically large (e.g., 1000) \rightarrow not scalable!
- A promising solution
 - k -means++ samples just one point per iteration
 - What if we oversample by sampling each point independently with a larger probability?
 - Equivalent to updating the distribution less frequently
 - Coarser sampling $\rightarrow k$ -means | | [Bahmani et al. '12]

k-means | | Initialisation

k=4, oversampling factor L=3

Cluster the intermediate centres







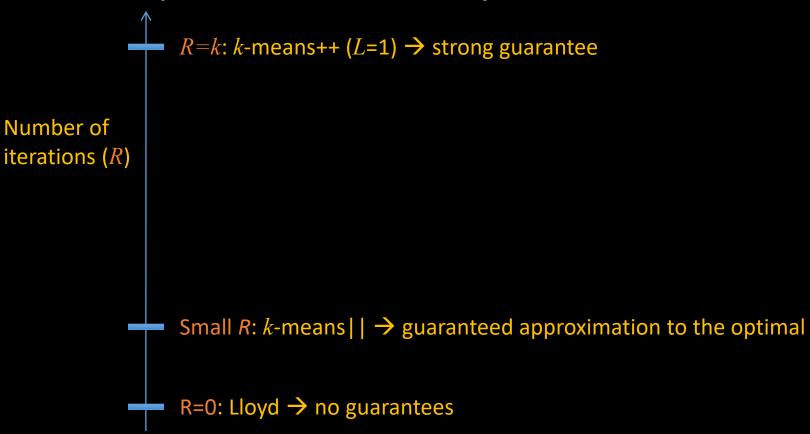


k-means | | [Bahmani et al. '12]

- Choose the oversampling factor L>1
- Initialise C to an arbitrary set of points
- For R iterations do:
 - Sample each point x in X independently with probability $p_x = Ld(x,C)^2/SSE_X(C)$.
 - Add all the sampled points to C
- ullet Cluster the intermediate centres in C using k-means++
- Benefits over k-means++
 - Less susceptible to noisy outliers
 - More reduction in the number of Lloyd iterations

k-means | | : Relationships

An interpolation between Lloyd and k-means++



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k-means in Spark

- Implemented in the RDD API Mllib Kmeans
- Scala ml code: org/apache/spark/ml/clustering/KMeans.scala

```
import org.apache.spark.ml.util._
import org.apache.spark.ml.util.Instrumentation.instrumented

import org.apache.spark.mllib.clustering.{DistanceMeasure, KMeans => MLlibKMeans, KMeansModel => MLlibKMeansModel}

import org.apache.spark.mllib.linalg.{Vector => OldVector, Vectors => OldVectors}

import org.apache.spark.mllib.linalg.VectorImplicits._
```

• Scala mllib: org/apache/spark/mllib/clustering/KMeans.scala

```
/**
34  * K-means clustering with a k-means++ like initialization mode
35  * (the k-means|| algorithm by Bahmani et al).
36  *
37  * This is an iterative algorithm that will make multiple passes over the data, so any RDDs given
38  * to it should be cached by the user.
39  */
40  @Since("0.8.0")
41  class KMeans private (
42  private var k: Int,
```

k-means API

- k: the number of desired clusters
- maxIter: the maximum number of iterations
- initMode: random or via k-means | (default) initialization
- initSteps: the number of steps in the k-means | | algorithm (default=2, oversampling factor $L = 2 \times k$)
- tol: the distance threshold for checking convergence
- seed: the random seed
- distanceMeasure: Euclidean (default) or cosine dist measure
- weightCol: optional weighting of data points

```
// On each step, sample 2 * k points on average with probability proportional
// to their squared distance from the centers. Note that only distances between points
// and new centers are computed in each iteration.

var step = 0

val bcNewCentersList = ArrayBuffer[Broadcast[_]]()
while (step < initializationSteps) {</pre>
```

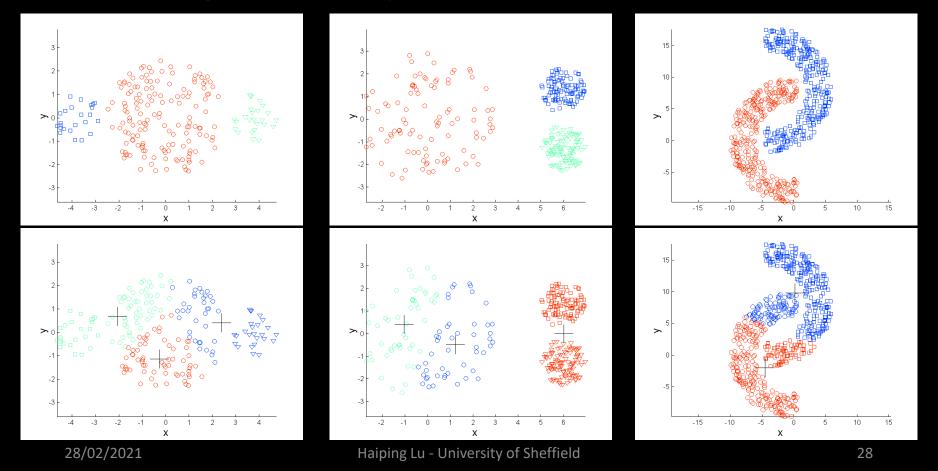
Remember to Cache

- Data need to be cached for high performance
- See from the source code

```
/**
     * K-means clustering with a k-means++ like initialization mode
     * (the k-means | algorithm by Bahmani et al).
36
     * This is an iterative algorithm that will make multiple passes over the data, so any RDDs given
     * to it should be cached by the user.
    @Since("0.8.0")
    class KMeans private (
41
         * Train a K-means model on the given set of points; `data` should be cached for high
207
         * performance, because this is an iterative algorithm.
208
         */
209
        @Since("0.8.0")
210
        def run(data: RDD[Vector]): KMeansModel = {
211
          val instances = data.map(point => (point, 1.0))
212
213
          val handlePersistence = data.getStorageLevel == StorageLevel.NONE
```

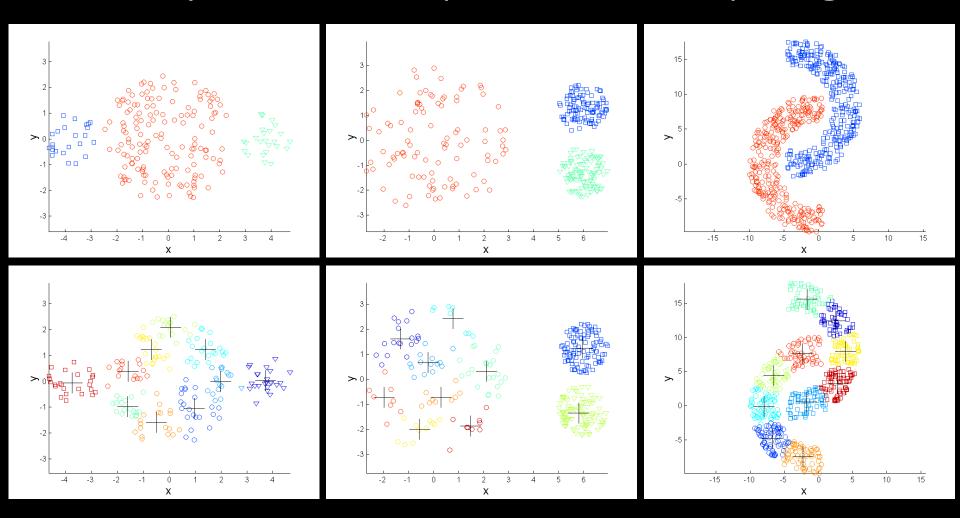
Limitations of *k*-means

 Problems when clusters are of differing sizes, densities, or non-globular shapes



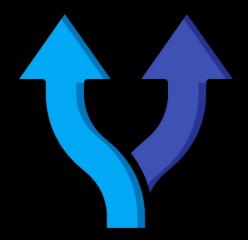
Overcoming k-means Limitations

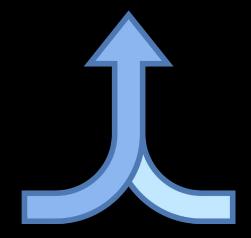
Use many clusters to find parts of clusters → put together



Pre-processing & Post-processing

- Pre-processing
 - Normalise the data (distance measure)
 - Eliminate outliers
- Post-processing
 - Eliminate small clusters that may represent outliers
 - Split loose clusters (with relatively high SSE)
 - Merge clusters that are close (with relatively low SSE)





Acknowledgement & References

Acknowledgement

 Some slides are adapted from 1) the <u>k-means|| slides</u> by Bahman Bahmani, Stanford University, 2012, and 2) Tan, Steinbach, Kumar's slides for the book <u>Introduction to</u> <u>Data Mining</u>

References

- Chapter on clustering from the book above (88 pages)
- *k*-means overview
- *k*-means ++ paper
- <u>k-means || paper and video</u>