

#### Lecture 8: Scalable k-means Clustering

COM6012: Scalable ML with Robert Loftin

Slides courtesy of Haiping Lu

#### Week 8 Contents / Objectives

Introduction to Cluster Analysis

• k-means Clustering

Scalable k-means

• *k*-means in Spark & Limitations

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• k-means Clustering

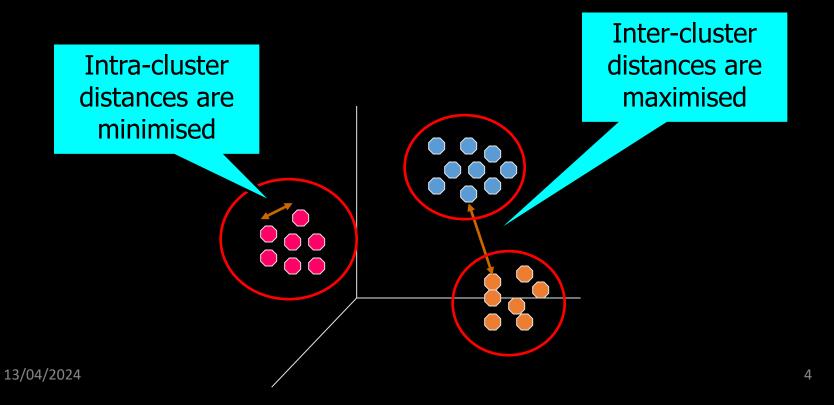
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# Cluster Analysis

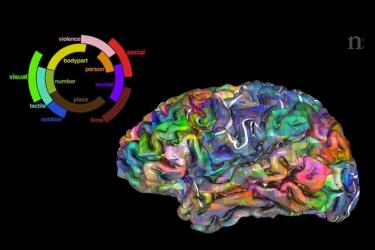
Output	Supervised	Unsupervised
Discrete	Classification	Clustering
Continuous	Regression	Dimensionality Reduction

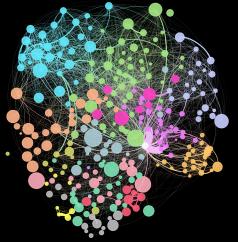
- "Classification without labelled data"
- "Dimensionality reduction to 1 (cluster index)"



## Cluster Analysis

- Divide data into clusters that are meaningful/useful
- Clusters: pseudo-classes
- Key for exploratory data analysis 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

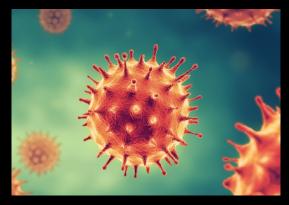
- Anomalies: data points very different from others
- Applications: credit card fraud, network intrusion, unknow virus, eco disturbance, unusual symptom, ...



<u>financial protection credit card fraud img1</u> <u>.jpg (512×347) (traderdefenseadvisory.com)</u>



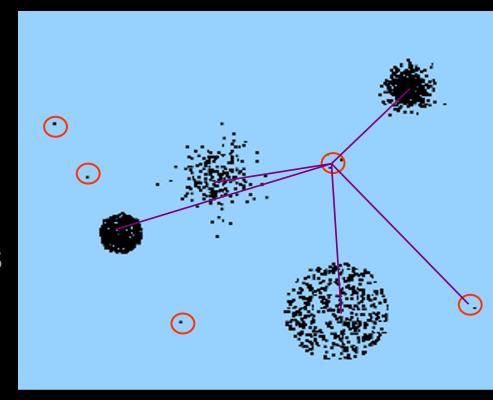
<u>Network-Intrusion-Detection-and-</u> Prevention-.jpg (1600×962) (wp.com)



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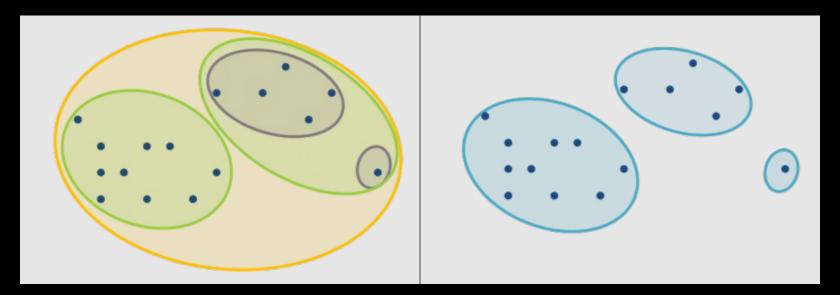
# Clustering-based Anomaly Detection

- 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

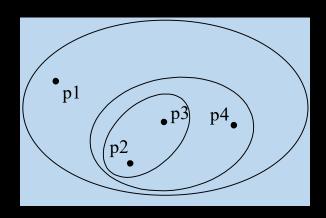


#### Hierarchical vs Partitional Clustering

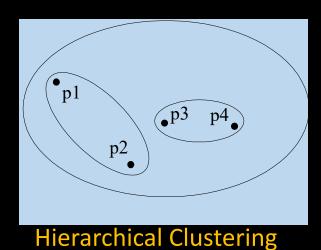
- Hierarchical: nested clusters as a hierarchical tree
  - Each node (cluster) in the tree (except for the leaf nodes) is the union of its children (subclusters)
  - The root of the tree → the cluster containing all data points
- Partitional: non-overlapping clusters
  - Each data point is in exactly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



#### Hierarchical Clustering

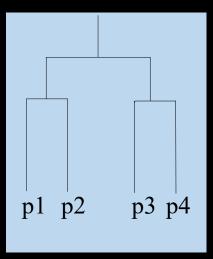


**Hierarchical Clustering** 



p1 p2 p3 p4

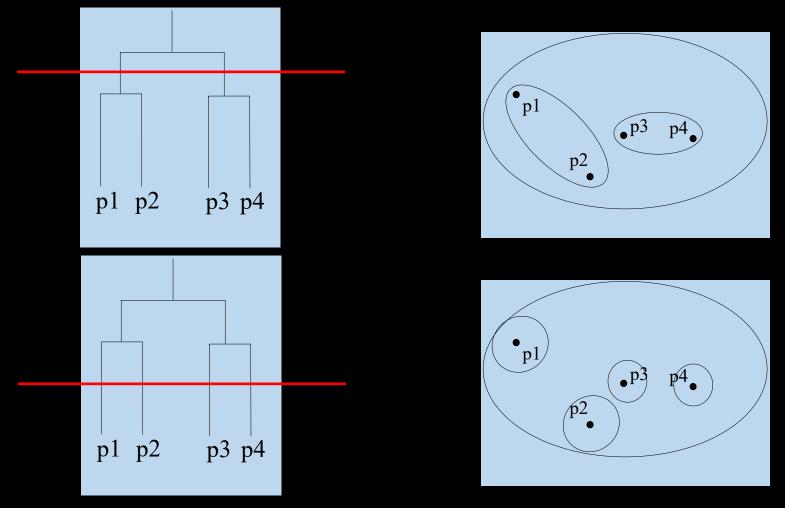
Dendrogram



Dendrogram

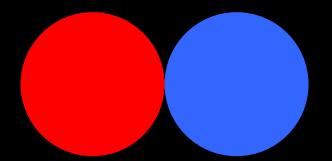
#### Hierarchical -> Partitional

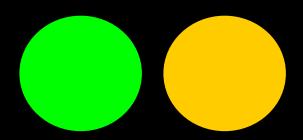
 Hierarchical = a sequence of partitional clustering cutting the hierarchical tree at a particular level



## 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)

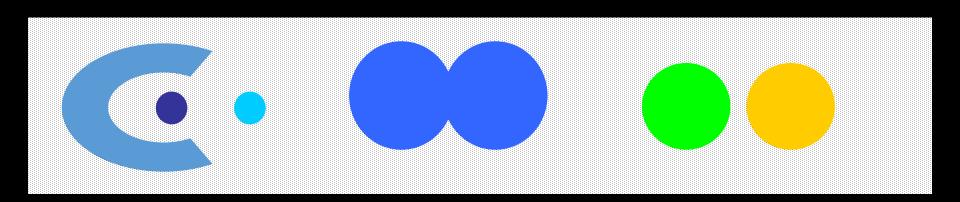




4 centre-based clusters

#### 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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## 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 $\rightarrow$ Stuck

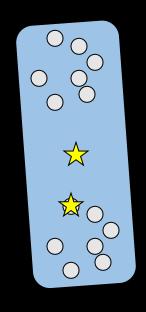






Figure credited to David Arthur

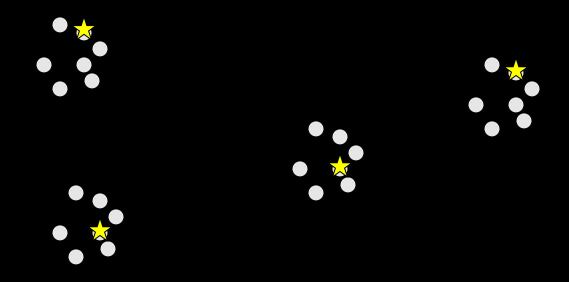
## k-means++ [Arthur et al. '07]

- 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



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

- k-means++ limitations
  - Needs k passes over the data for initialisation
  - 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 | | [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 | | Initialisation

k=4, oversampling factor L=3

Cluster the intermediate centres



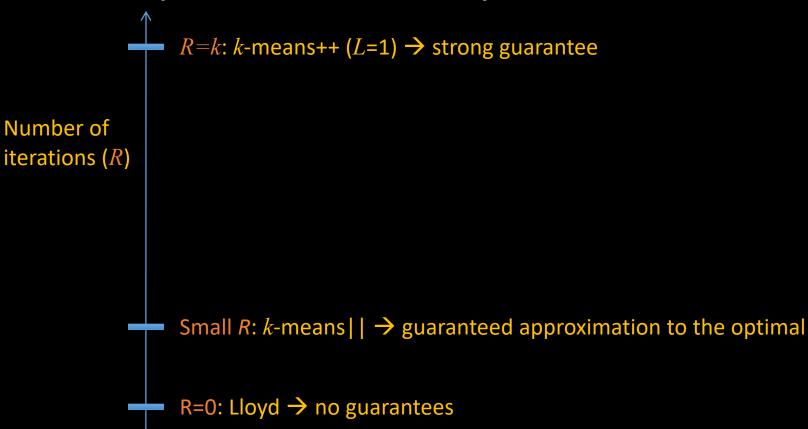






# k-means $|\cdot|$ : Relationships

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

```
/**

* K-means clustering with a k-means++ like initialization mode

* (the k-means|| algorithm by Bahmani et al).

* 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 (
    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>
```

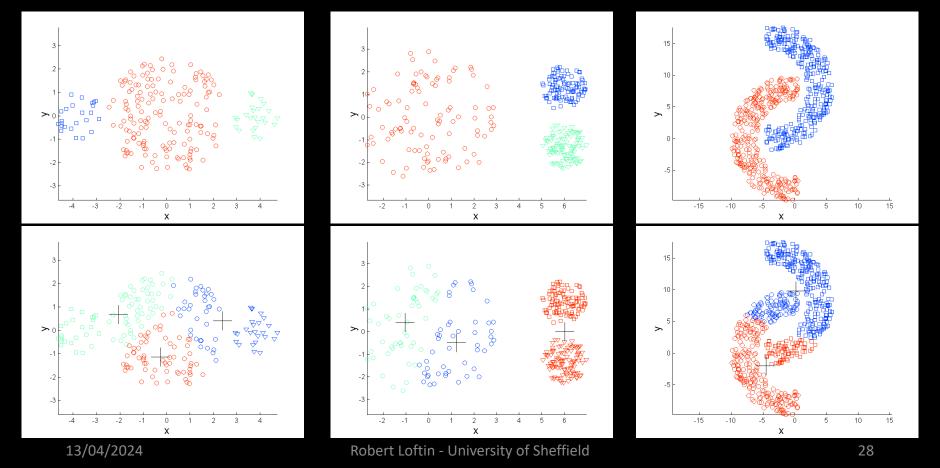
#### Remember to Cache

- 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
212
          val instances = data.map(point => (point, 1.0))
          val handlePersistence = data.getStorageLevel == StorageLevel.NONE
213
```

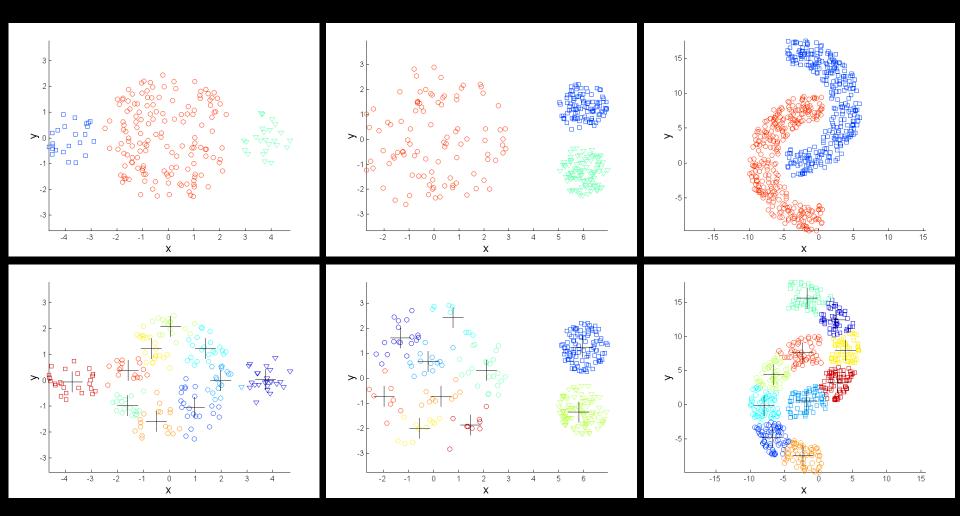
#### Limitations of k-means

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



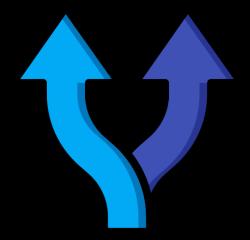
#### Overcoming k-means Limitations

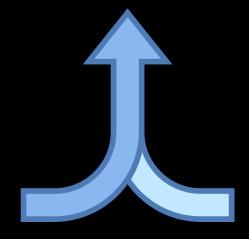
Use many clusters to find parts of clusters → put together



#### 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>