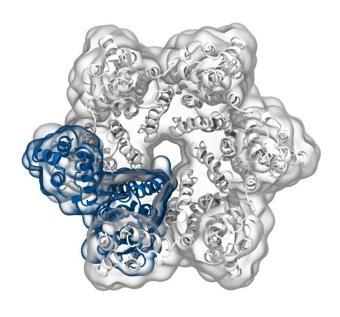
## Simulation of Biomolecules



### Clustering



Dr Matteo Degiacomi

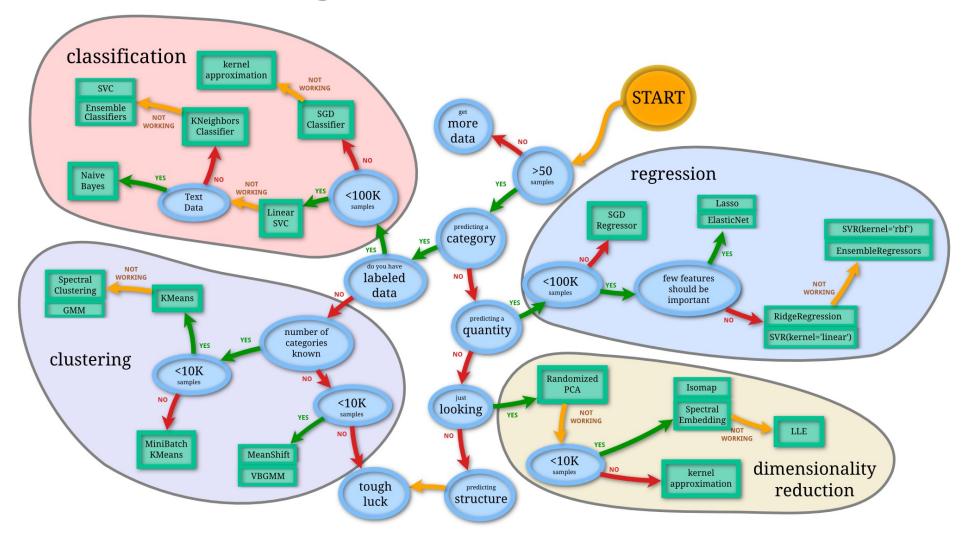
Durham University

matteo.t.degiacomi@durham.ac.uk

Dr Antonia Mey University of Edinburgh

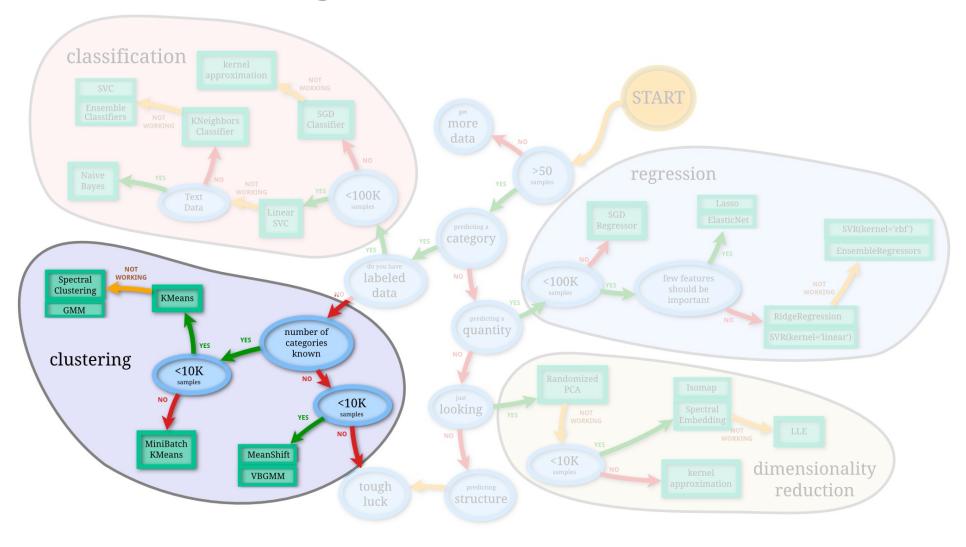
antonia.mey@ed.ac.uk

### The Data Mining world



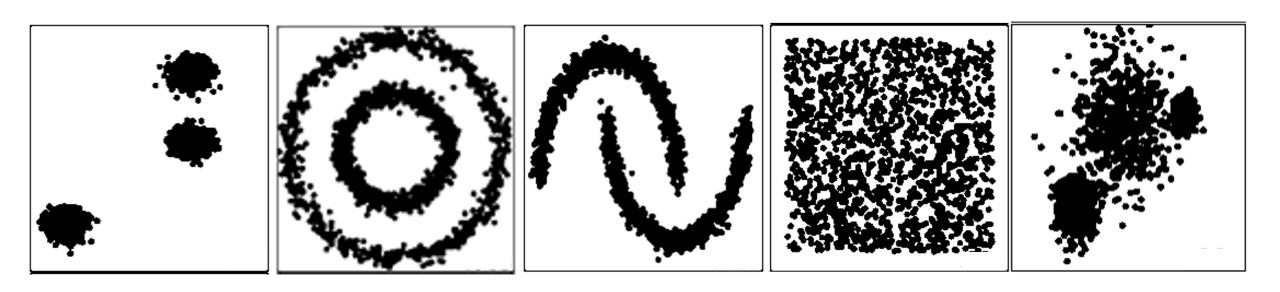
From scikit-learn.org 2

### The Data Mining world



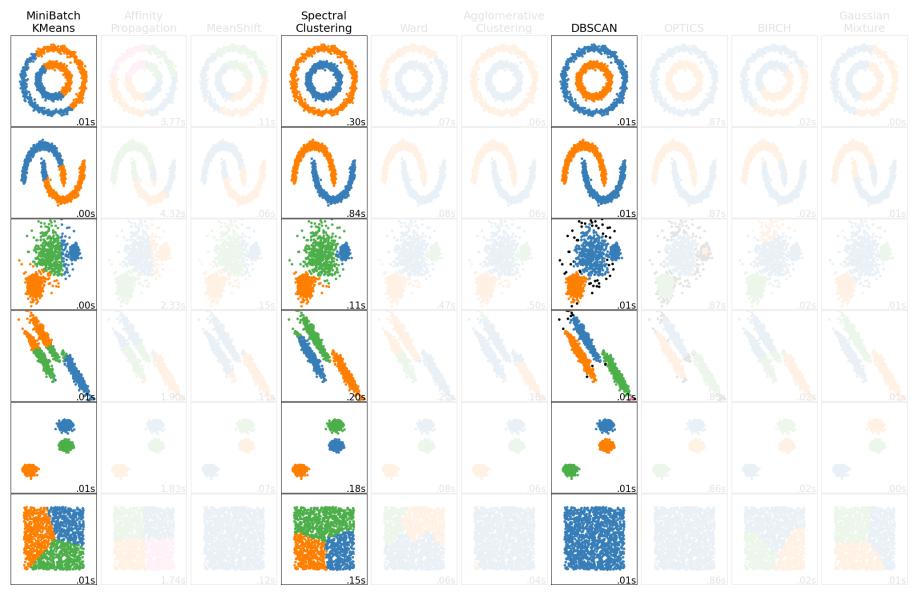
From scikit-learn.org

### Clustering (i.e., unsupervised learning)



Known number of clusters? Flat geometry? Even cluster size? Outliers? Centroids needed?

## Clustering algorithms



### How does k-means work?

**Input**: K, set of points x<sub>1</sub>...x<sub>n</sub> (can be in N-dimensional)

Place centroids, c1...,cn at random locations

Repeat until convergence:

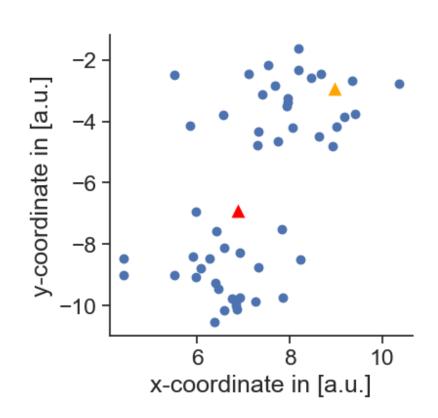
For each point xi:

Find nearest centroid  $c_j = \arg\min_j D(x_i, c_j)$ Assign the point  $x_i$  to cluster j

For each cluster j = 1...K:

Compute the centroid mean for all points in one cluster and update the centroid

$$c_j(a) = \frac{1}{n_j} \sum_{x_i \to c_j} x_i(a)$$



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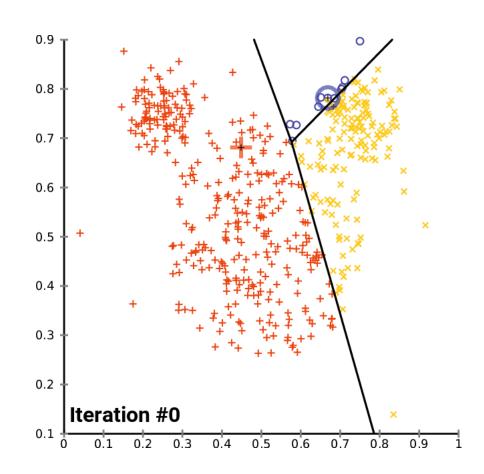
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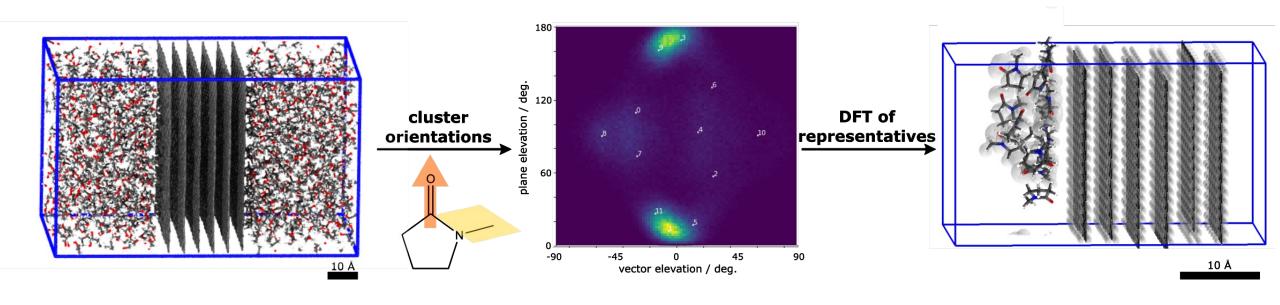
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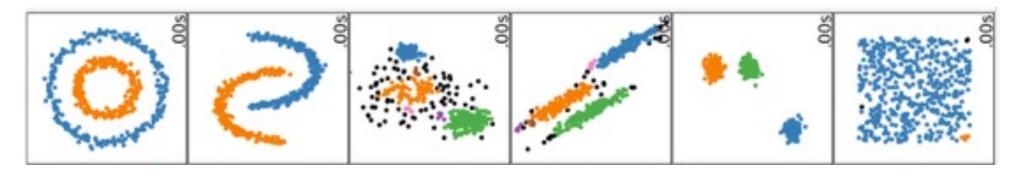
## [Example] k-means vs solvent-graphite interactions

- Molecular Dynamics simulation of graphite immersed in solvents.
- >100k individual solvent-graphene interactions

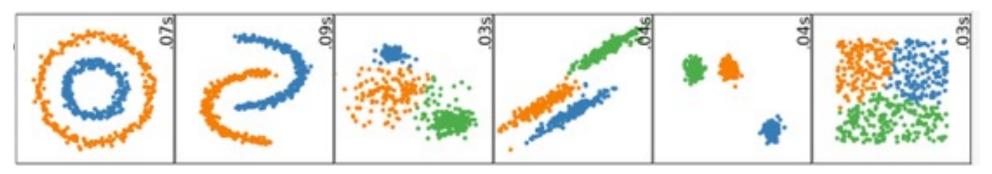


### Density-based and spectral clustering

#### **DBSCAN**

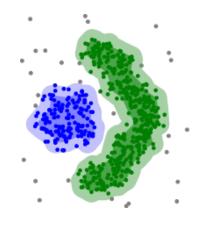


### **Spectral Clustering**



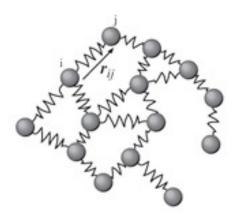
### Density-based and spectral clustering

#### **DBSCAN**



- Find the points in the  $\varepsilon$  neighbourhood of every point, identify core points with more than n neighbours.
- Find the connected components of core points on the neighbour graph, ignoring all non-core points.
- Assign each non-core point to a nearby cluster if the cluster is an ε neighbour, otherwise assign to noise otherwise.

## Spectral Clustering



- Calculate the Laplacian
- Calculate the first k eigenvectors
- Consider the matrix formed by the first k-eigenvectors
- Cluster the graph nodes based on these features (e.g. k-means)

# [Example] DBSCAN for noise detection in EM maps

- Load electron density map and chose a threshold t
- Place pseudoatoms where intensity > t
- Cluster beads and delete small clusters (<1% of total beads)</li>

