

Multivariate Data Analytics

Clustering Analysis

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Clustering

- **Objective:** Dividing records (rows) in a multivariate dataset into “natural” clusters (groups), where the records in each group are similar to one another.
- **Applications**
 - Cluster customers into different segments
 - Cluster products according to their attributes
 - Cluster businesses according to their location
- **Methods:** K-means, Model-based Clustering (Gaussian Mixture Model), Hierarchical Clustering, DBSCAN...
- **Distance Measure**
 - Euclidean Distance (most common): square-root of the sum of the squared differences between each variable.

$$d(\mathbf{X}_i, \mathbf{X}_j) = \sqrt{\sum_{k=1}^p (X_{ik} - X_{jk})^2}$$

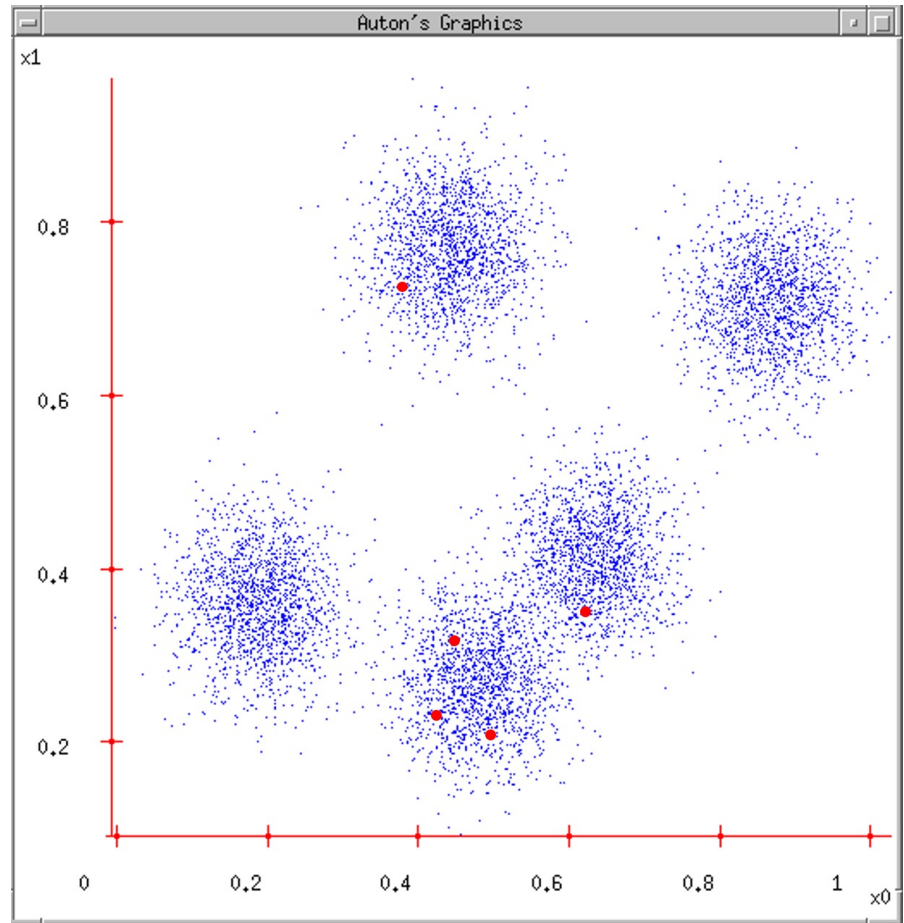
- Other measures may be more appropriate for a specific dataset



K-Means

K-means

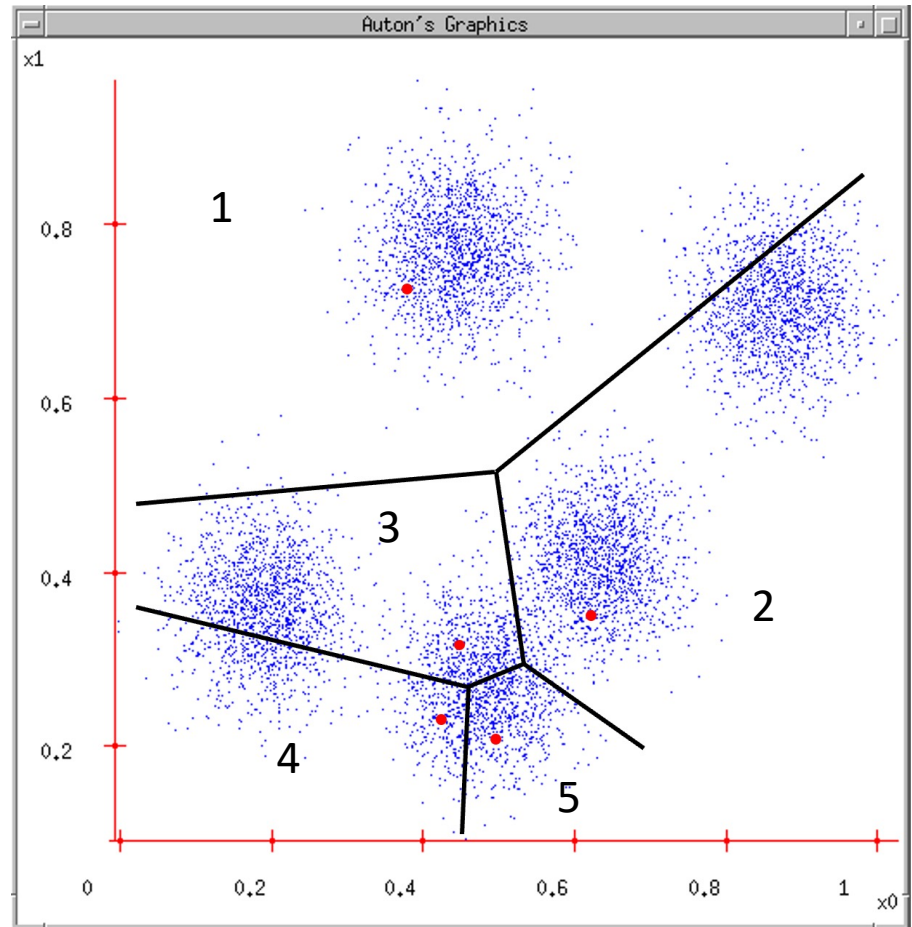
1. Ask user how many clusters they'd like. ($k = 5$)
2. Randomly guess k cluster centers



Example by Andrew W. Moore

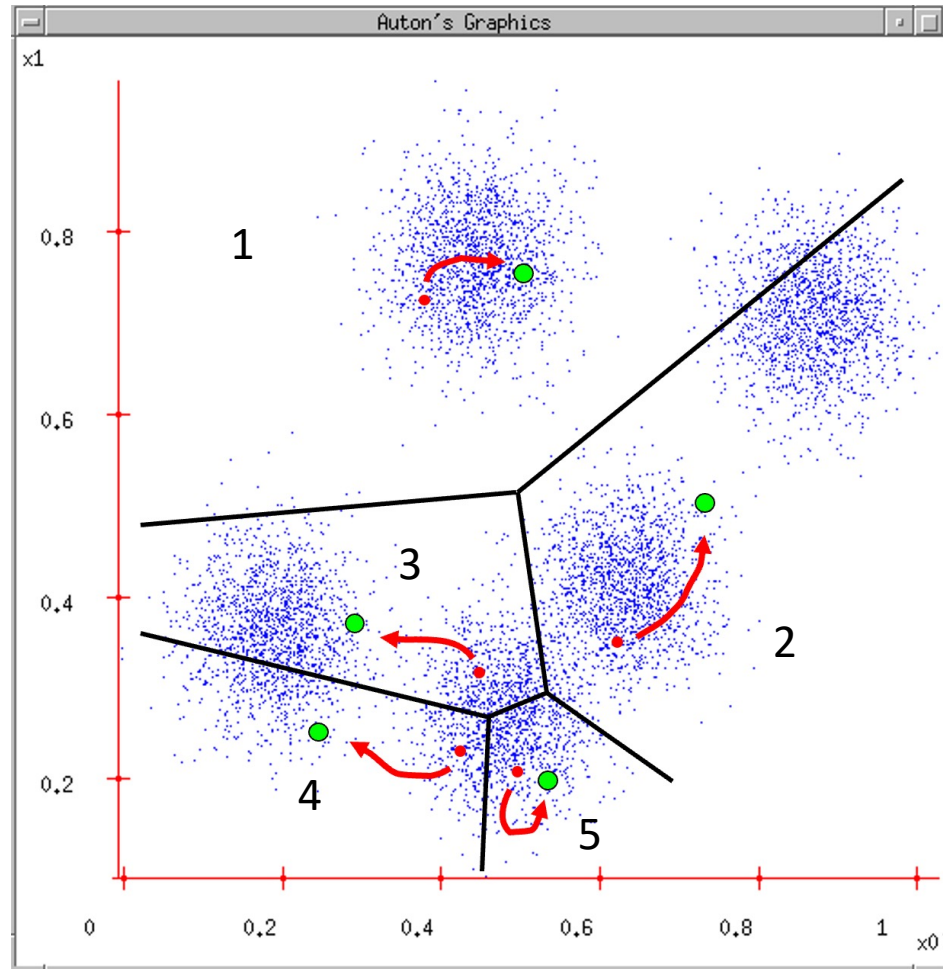
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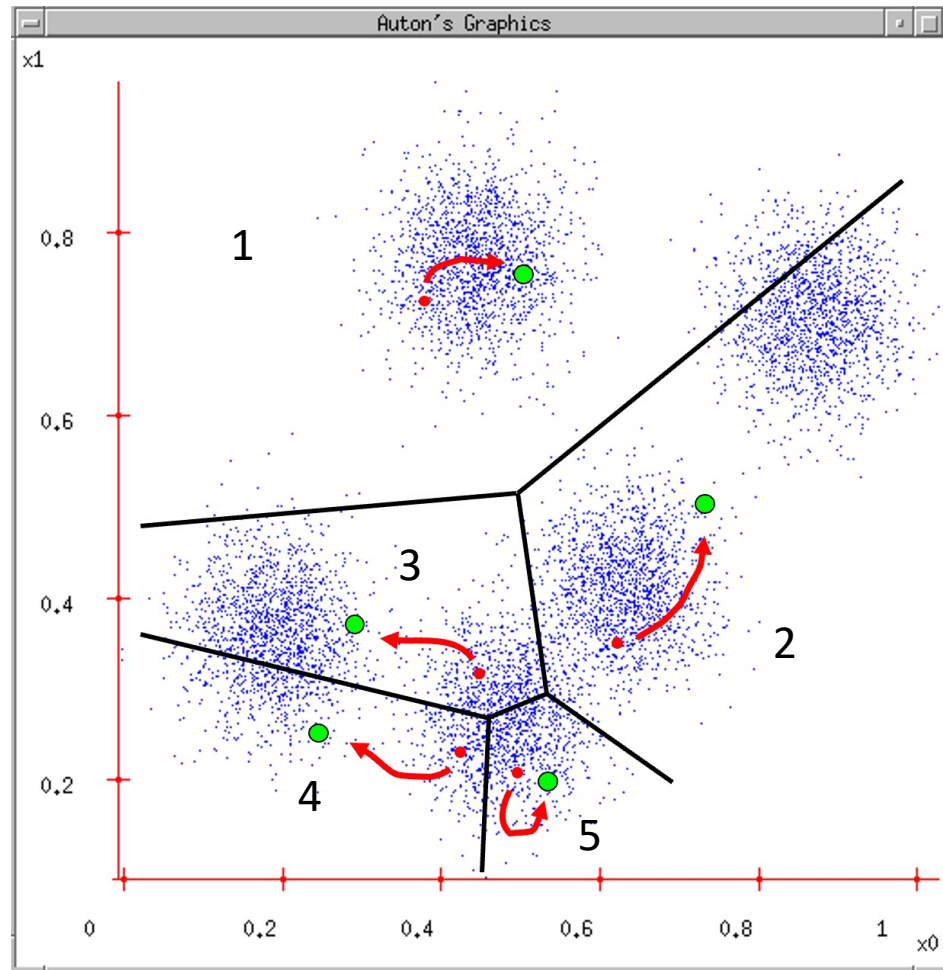
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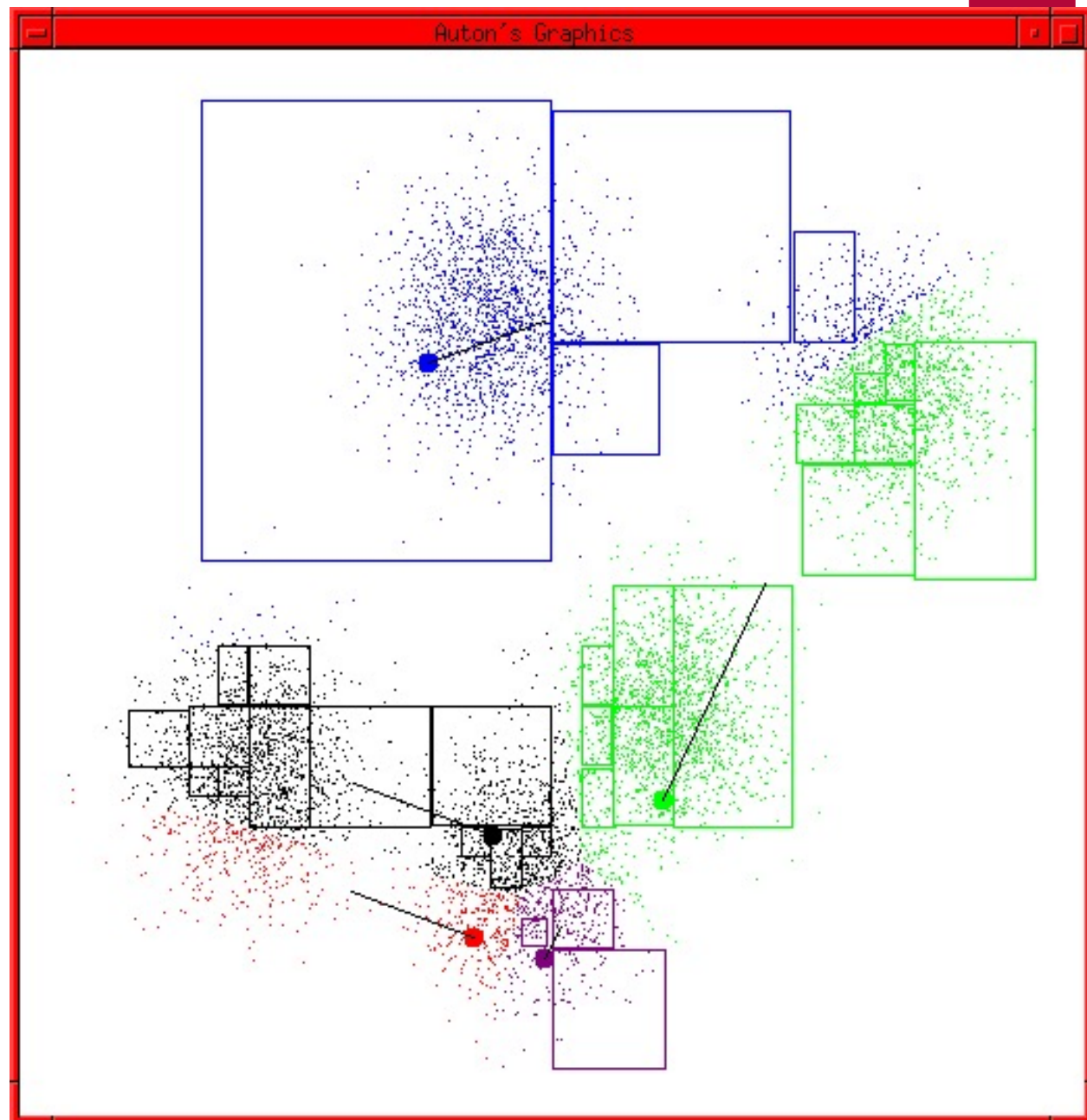


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5. Jump to Step 3. Repeat until terminated.

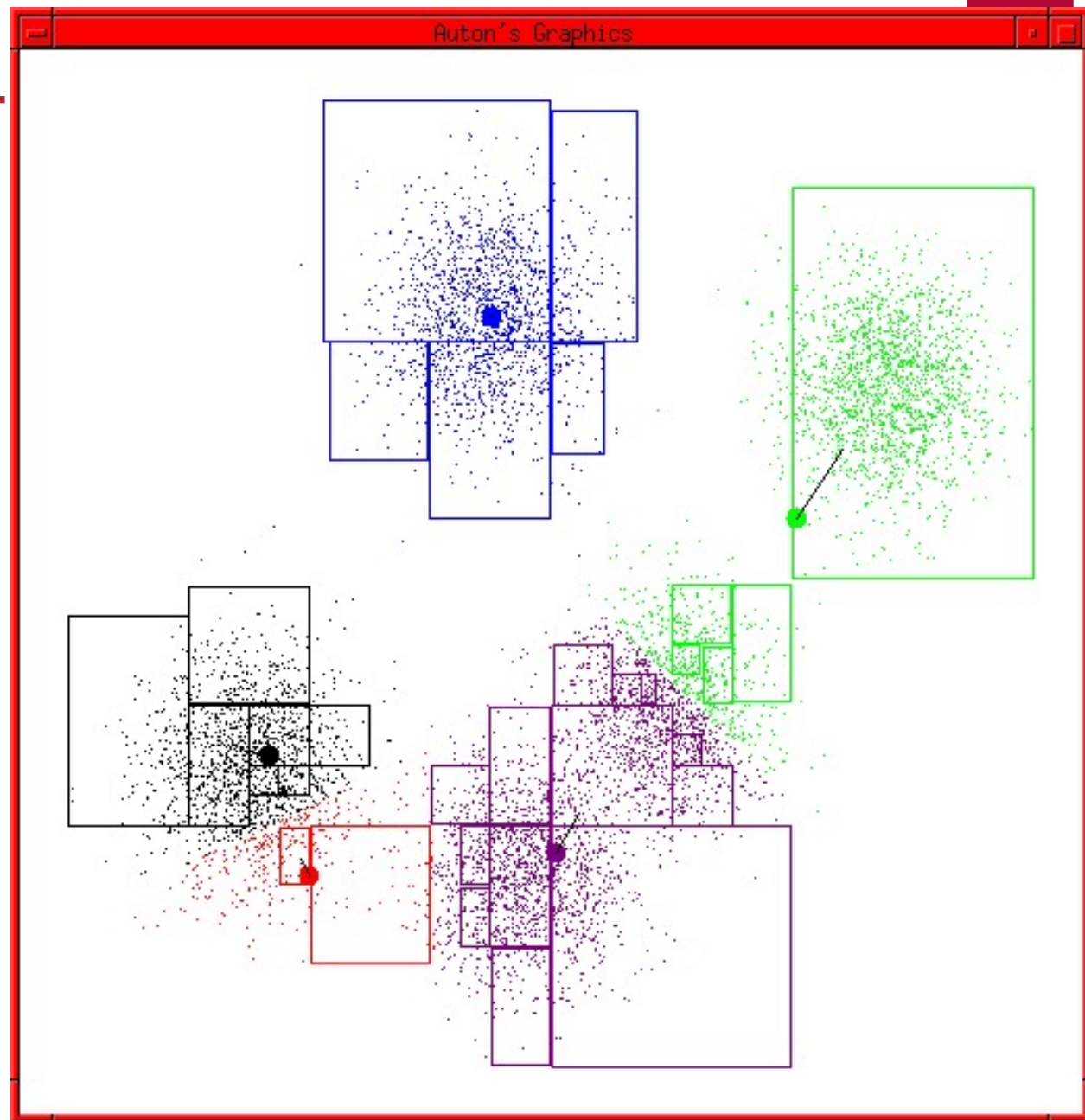


K-means starts

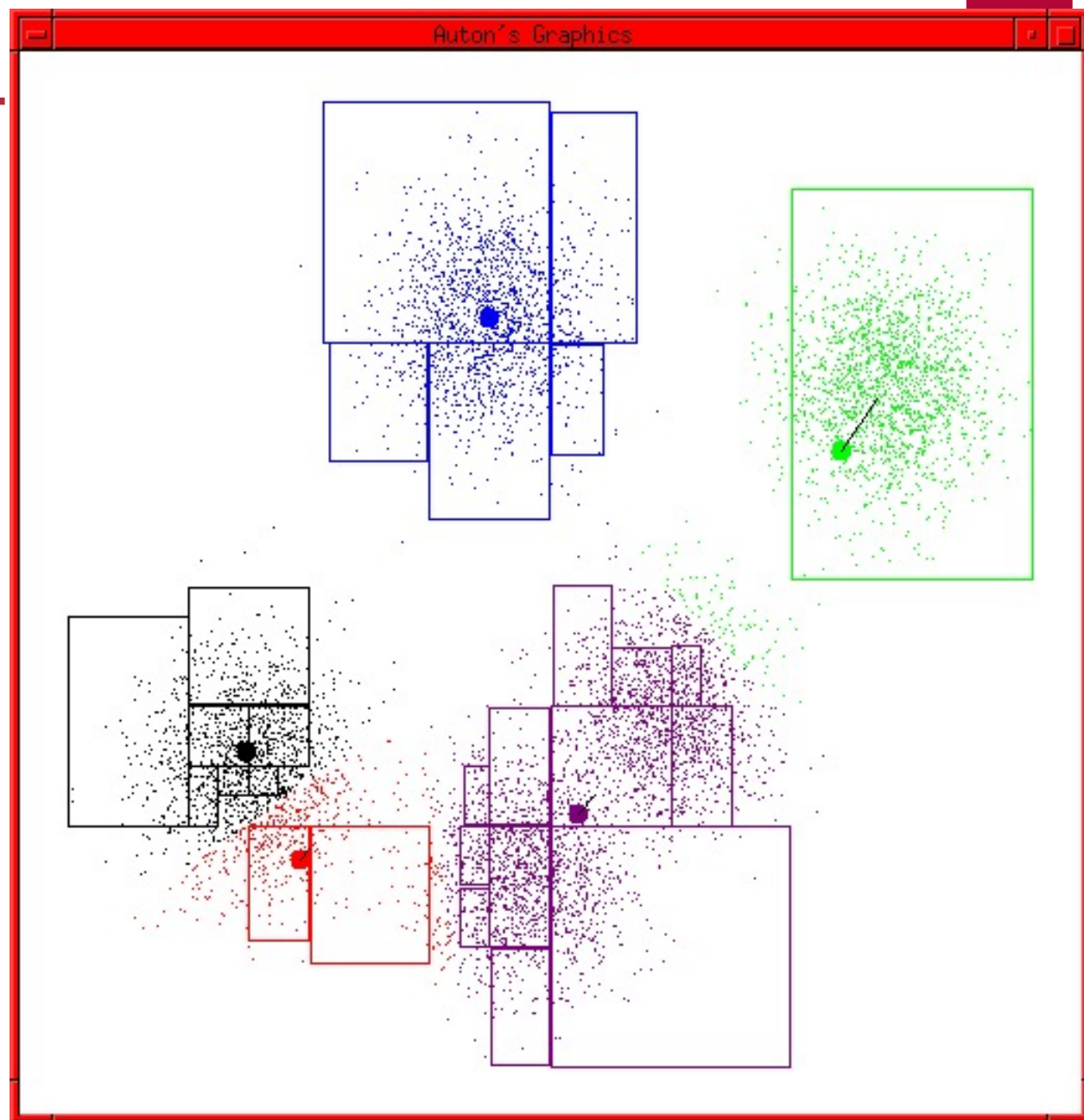


Source: www.autonlab.org

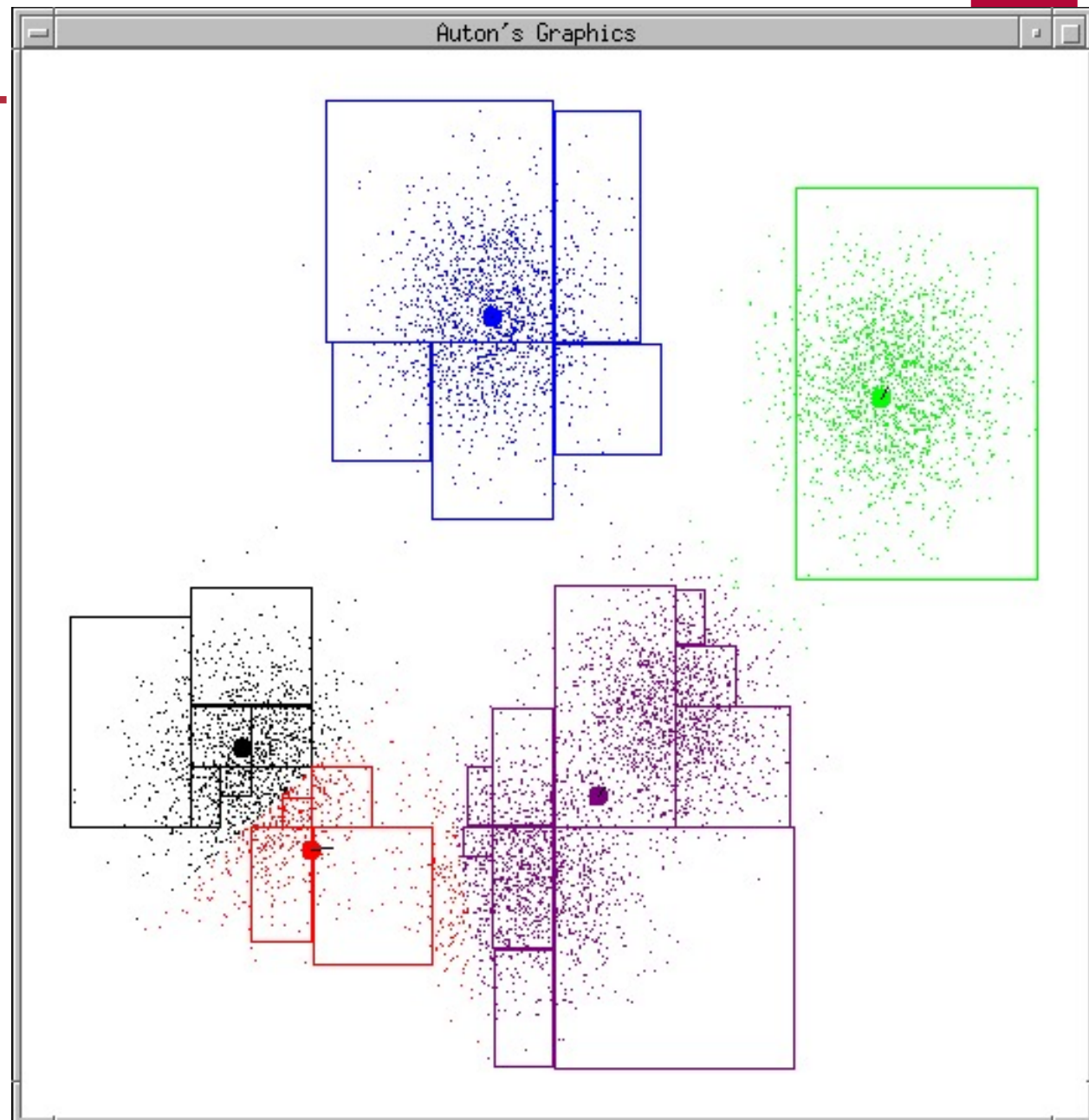
K-means continues.



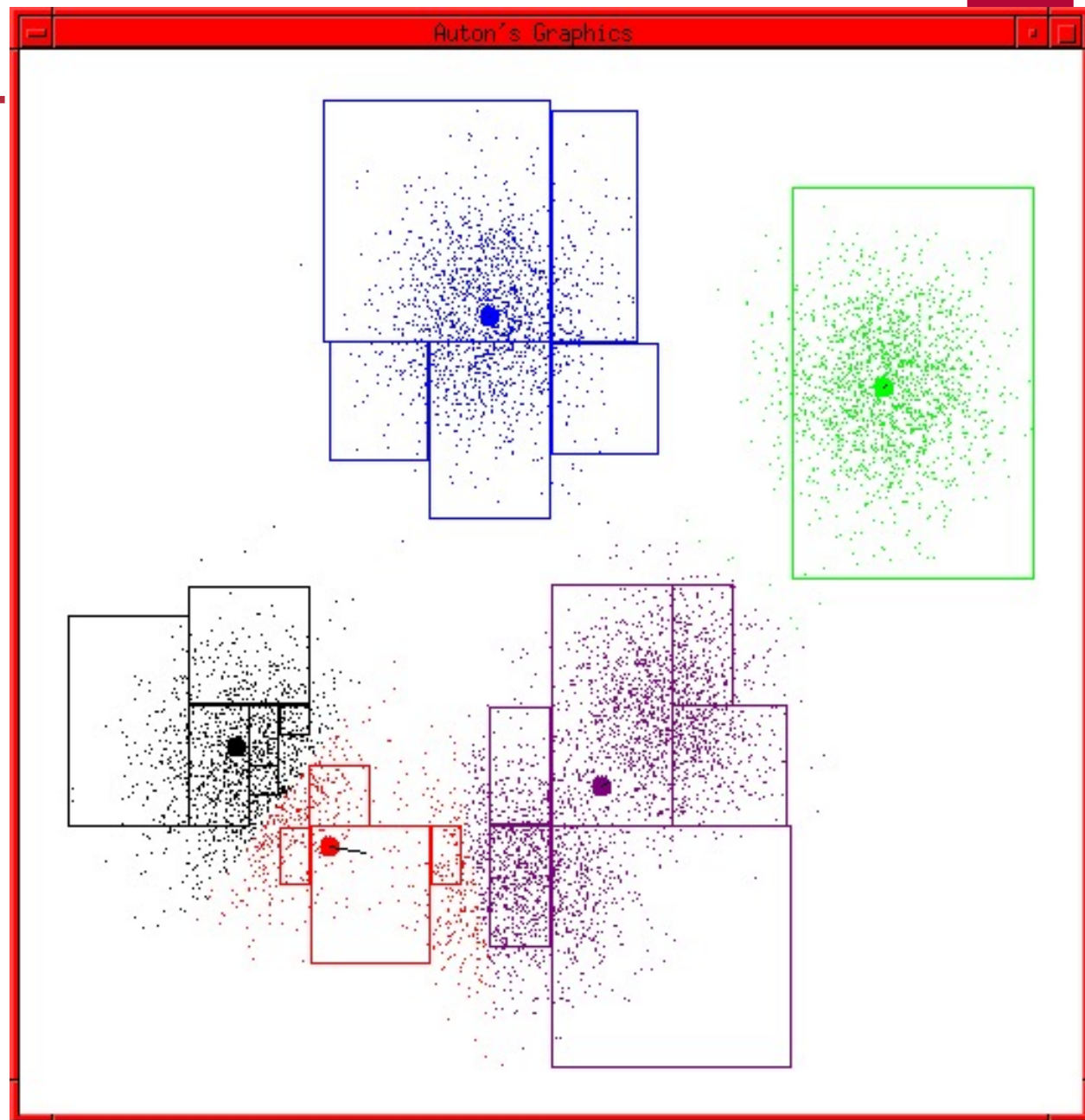
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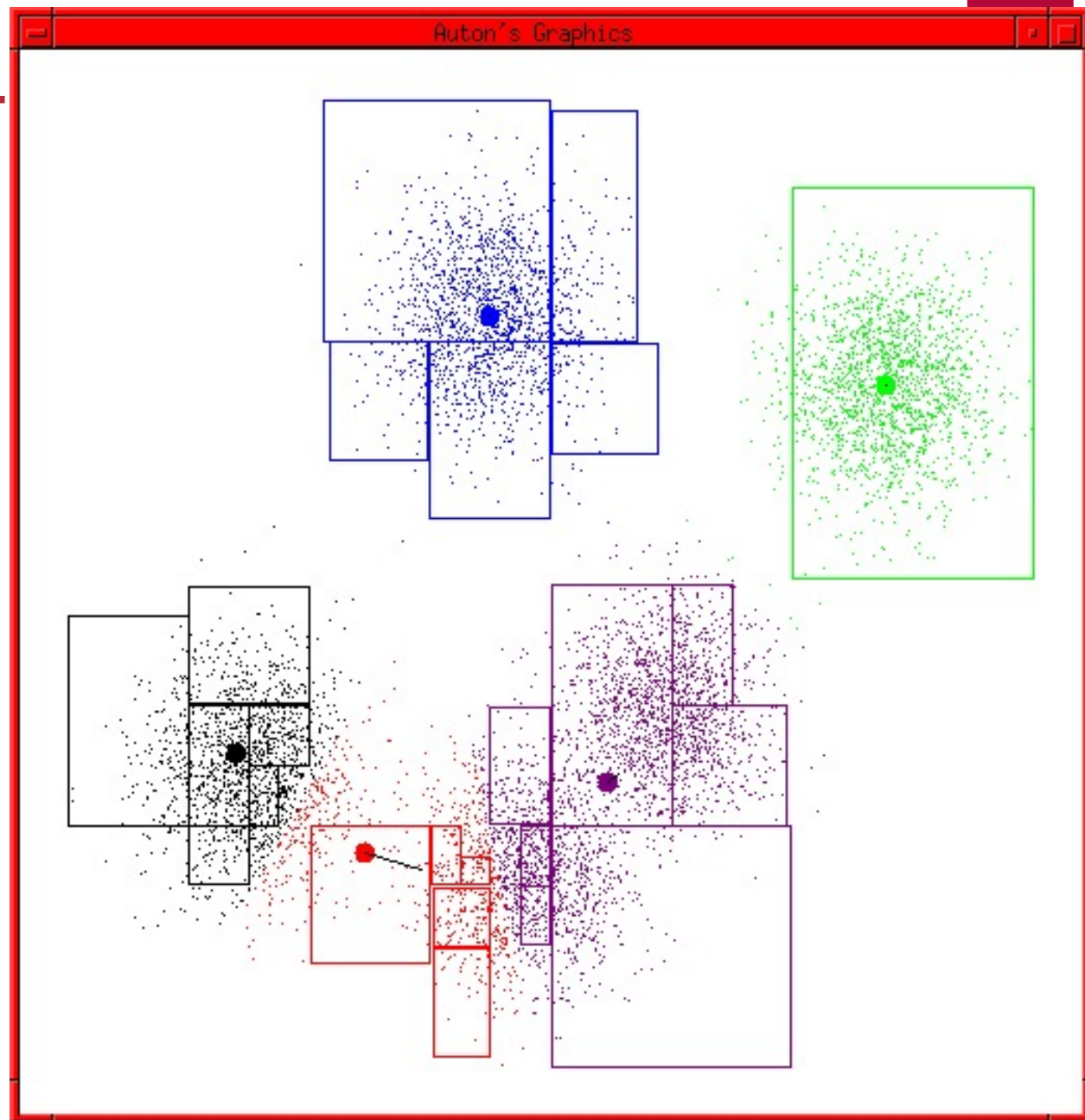
K-means continues.



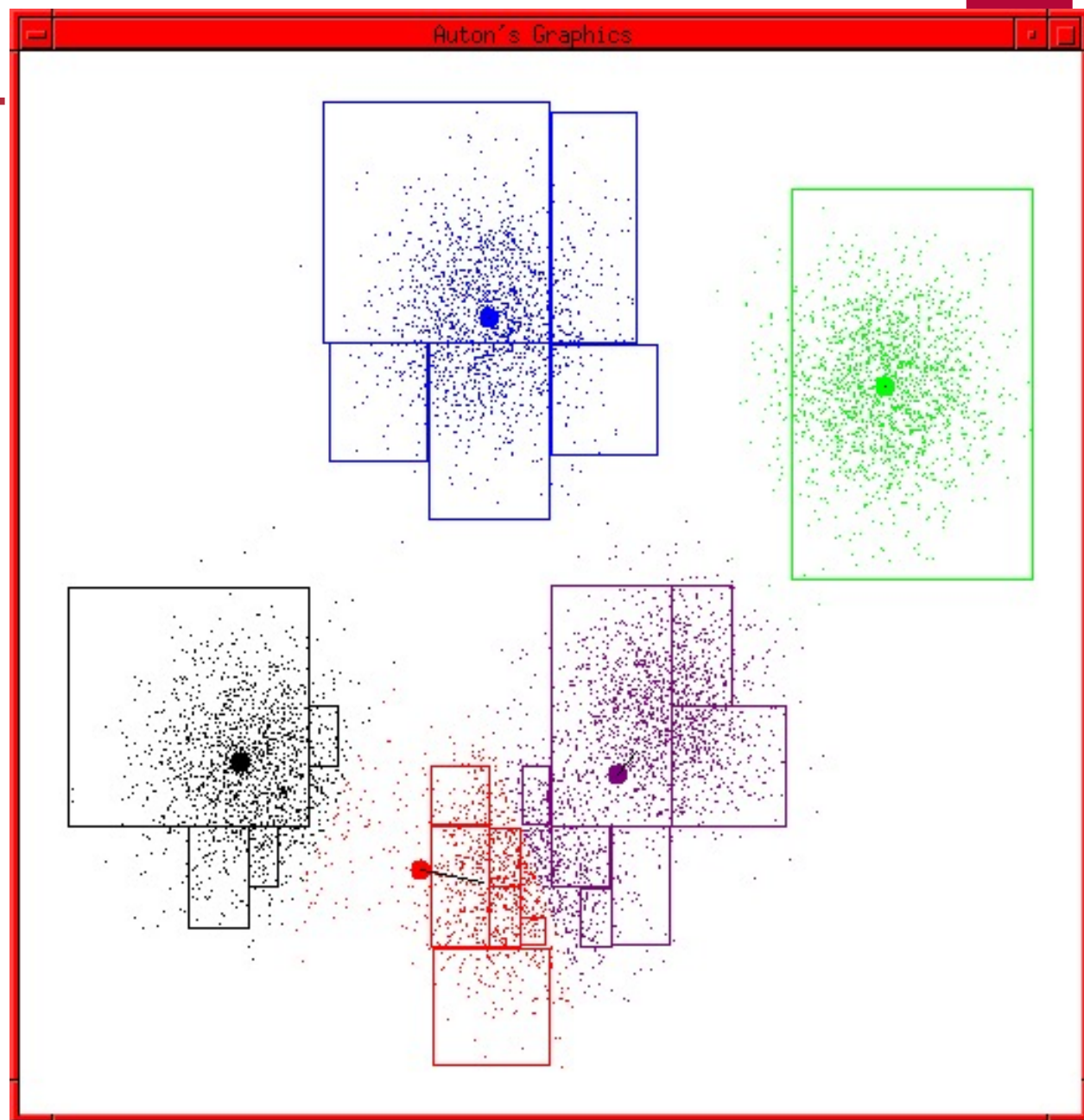
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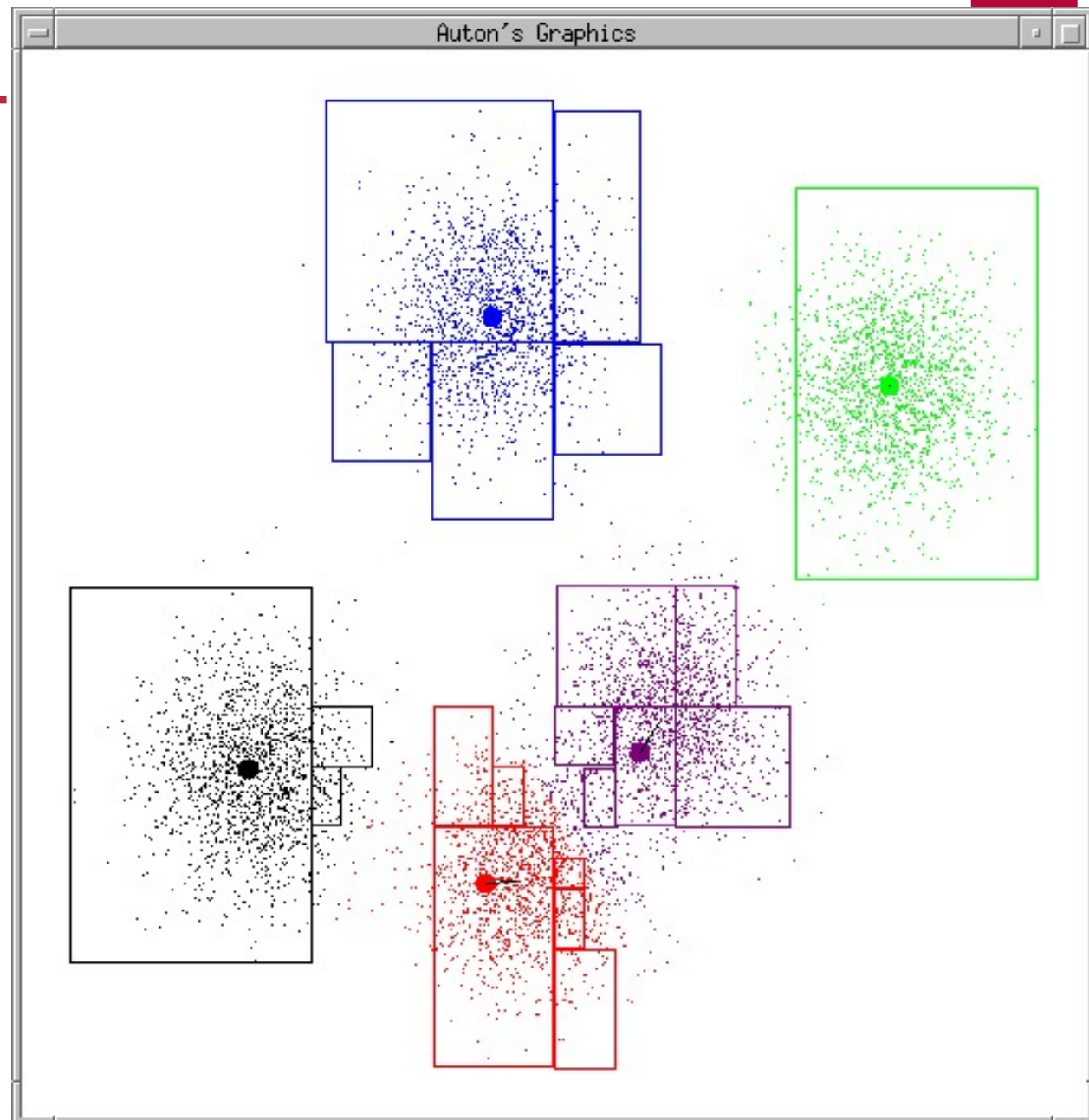
K-means continues.



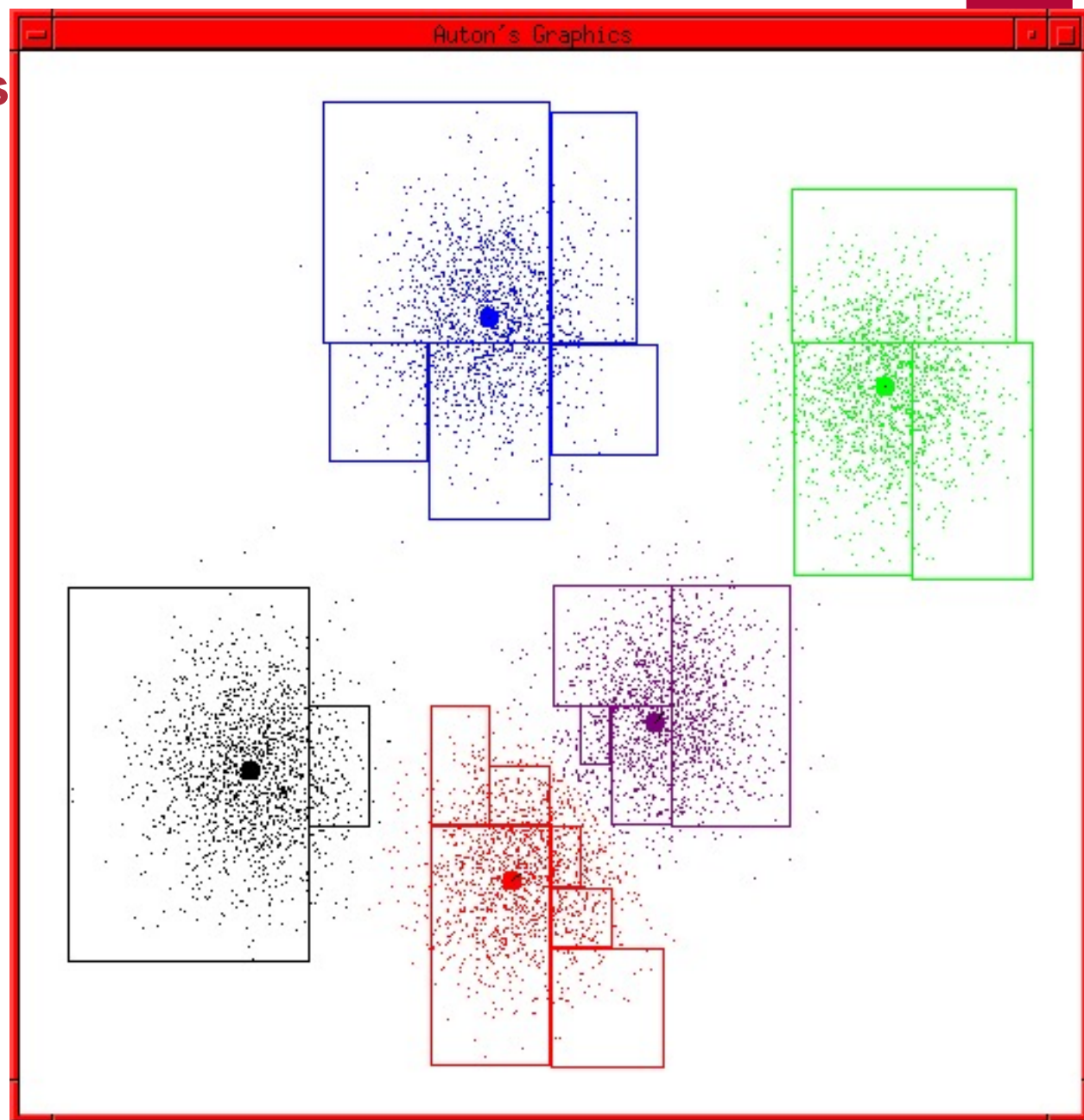
K-means continues.



K-means continues.



K-means terminates





K-means Questions

- What is it trying to optimize?
- Are we sure it will terminate?
- Are we sure it will find an optimal clustering?
- How should we start it?
- How could we automatically choose the number of centers?

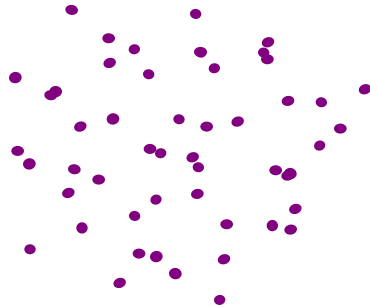


K-means objective

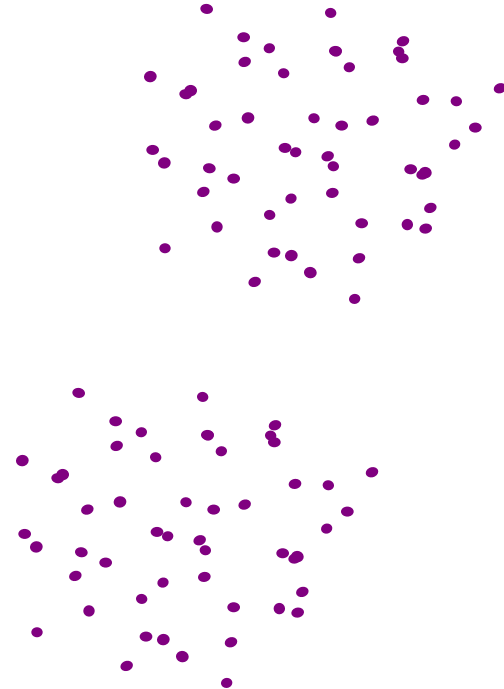
Given a set of observations $(\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_n)$, where each observation is a d -dimensional real vector, k -means clustering aims to partition the n observations into k ($\leq n$) sets $\mathbf{S} = \{S_1, S_2, \dots, S_k\}$ so as to minimize the within-cluster sum of squares (WCSS) (i.e. [variance](#)). Formally, the objective is to find:

$$\arg \min_{\mathbf{S}} \sum_{i=1}^k \sum_{\mathbf{x} \in S_i} \|\mathbf{x} - \boldsymbol{\mu}_i\|^2 = \arg \min_{\mathbf{S}} \sum_{i=1}^k |S_i| \text{Var } S_i$$

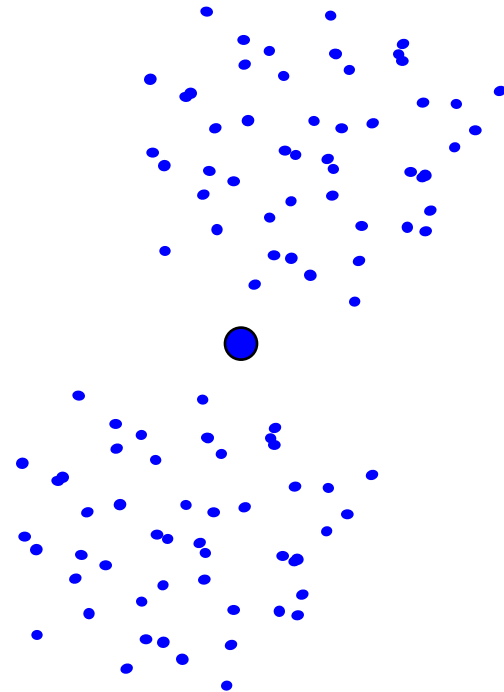
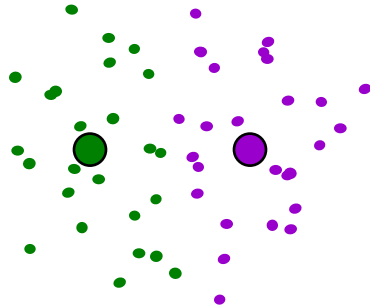
Will we find the optimal configuration?



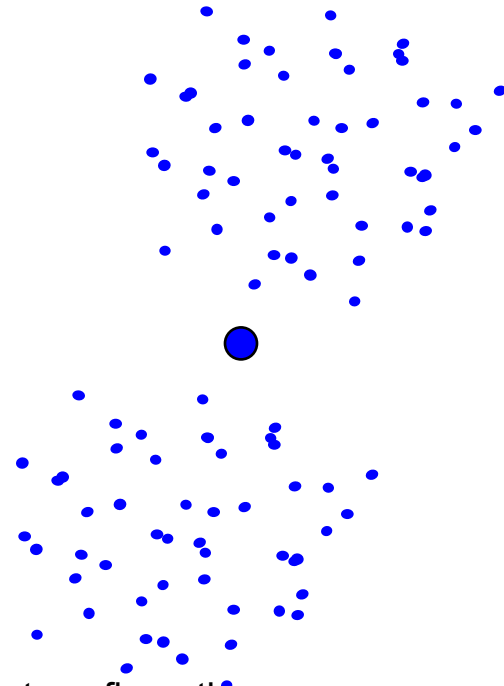
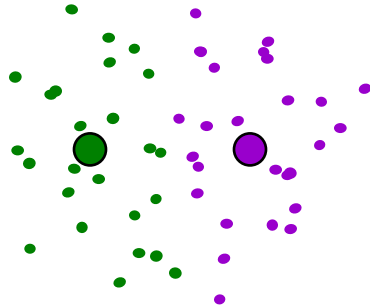
Not necessarily.



Will we find the optimal configuration?



Trying to find good optima



Idea 1: Be careful about where you start

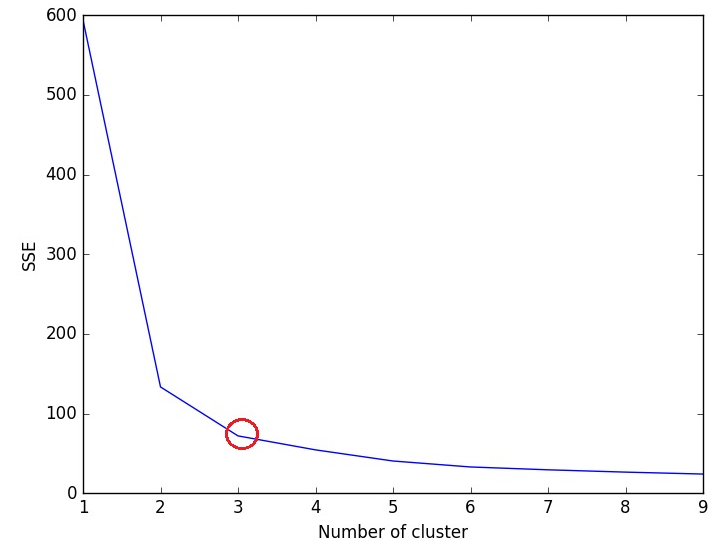
Idea 2: Do many runs of k-means, each from a different random start configuration

Many other ideas floating around. For example:

- Place first center on top of randomly chosen datapoint.
- Place second center on datapoint that's as far away as possible from first center
- Place j 'th center on datapoint that's as far away as possible from the closest of Centers 1 through $j-1$

Choosing the number of clusters (k)

- No simple answer
- Subjective evaluation: Are clusters interpretable?
- Elbow method:
 - Y-axis: **sum of squared errors (SSE)** inside each cluster (the squared difference between points and their cluster center). Alternatively, use **Variance Explained%**.
 - X-axis: number of clusters
 - Determine the elbow in the graph
- Other quality measures:
 - e.g., Silhouette coefficient, Dunn index...

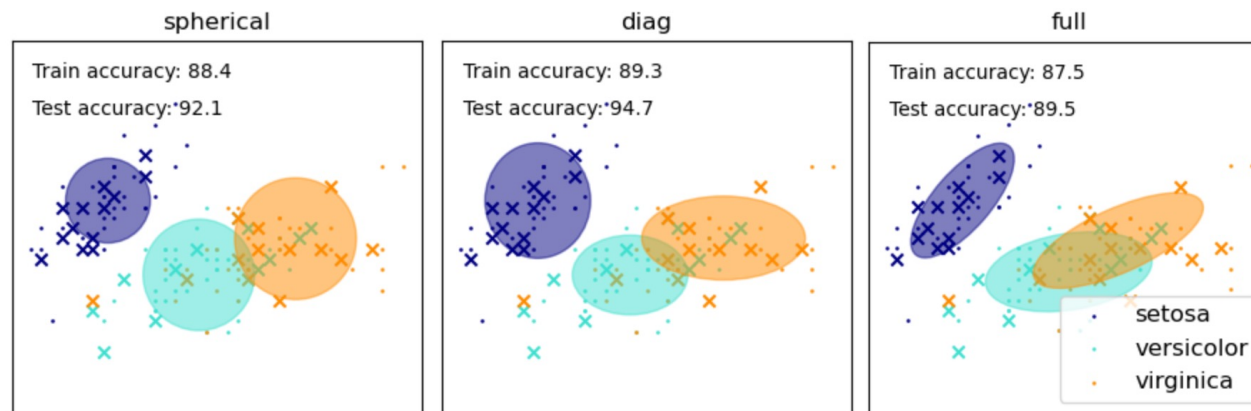




Other Clustering Methods

Model-Based Clustering

- Grounded in Multivariate Normal (Gaussian) Distribution
 - The data is generated from Mixtures of Normal
 - Constraining the covariance matrix to restrict the shape of clusters
- Can choose the number of clusters using criterion such as BIC
- Provides “soft” or probabilistic cluster assignment
- Uses the Expectation-Maximization (EM) algorithm to maximize the Likelihood function (see readings)
- K-means can be considered as a special case

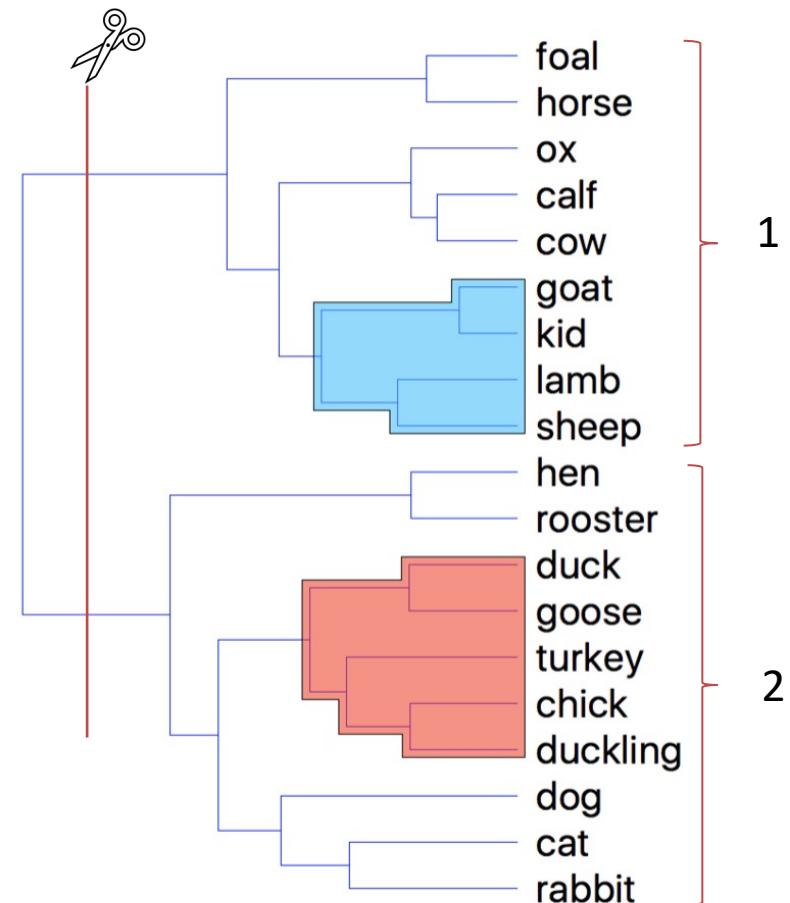


Source: https://scikit-learn.org/stable/auto_examples/mixture/plot_gmm_covariances.html

Hierarchical Clustering

- Merges or splits records in a greedy manner
 - Agglomerative: each observation starts in its own cluster, and pairs of clusters that are most similar to each other are merged
 - Divisive: all observations start in one cluster and clusters are spitted based on similarity
- Allows flexible distance (similarity) measure; not limited to Euclidean Distance
- Produces a **dendrogram** (tree) for the records
- We can choose the number of clusters by cutting the dendrogram at any level

2-cluster solution

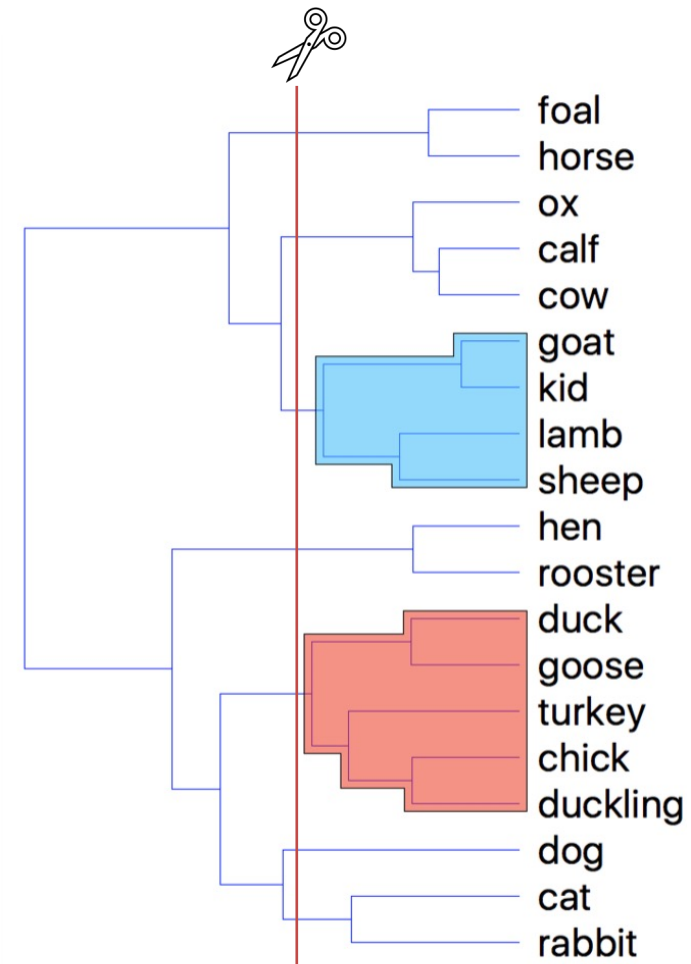


Source: <https://en.wikipedia.org/wiki/File:Orange-data-mining-hierarchical-clustering.png>

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7-cluster solution



Source: <https://en.wikipedia.org/wiki/File:Orange-data-mining-hierarchical-clustering.png>

DBSCAN (Density-based spatial clustering of applications with noise)

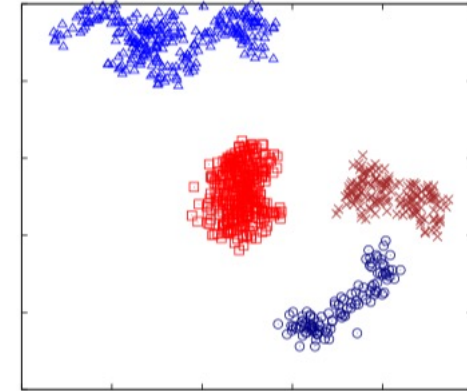
- Groups together points with many nearby neighbors
 - Neighbor is defined using a distance threshold ϵ

Advantages

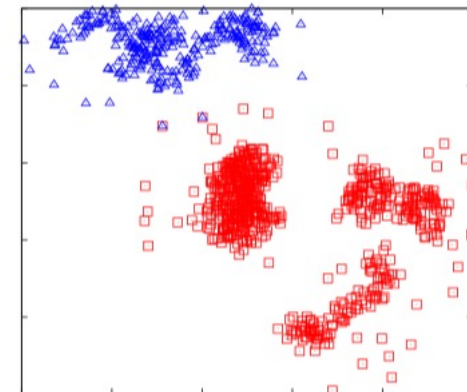
- Finds arbitrarily-shaped clusters
- No need to choose the number of clusters
- Handles outliers (finds noise points that do not belong a cluster)
- Fast for certain databases

Disadvantage

- Can be sensitive to the distance threshold ϵ



(d) $\rho = 0.1, \epsilon = 5000$



(h) $\rho = 0.1, \epsilon = 11300$

Source: Gan, J., & Tao, Y. (2015). DBSCAN revisited: Mis-claim, un-fixability, and approximation. *ACM SIGMOD*



Thank you!

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