

Clustering : k -means and k -medoids



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Introduction

A common task in data science is to search in a given data set for indications of internal structure that can be further explored, for example, an intrinsic organization of data into clusters.

Clustering Problem

We start by formulating the basic clustering problem: Given a set of vectors,

$$\mathcal{D} = \{x^{(1)}, x^{(2)}, \dots, x^{(p)}\}$$

arrange them into k distinct subsets, or clusters.

$$\mathcal{D}_\ell = \{x^{(j)} : j \in I_\ell\}, \ell = 1, \dots, k$$

where

$$I_i \cap I_j = \emptyset, \forall i \neq j$$

and

$$\bigcup_{\ell=1}^k I_\ell = \{1, 2, \dots, p\}.$$

The k -means algorithm

Assume that the data set consists of real vectors.

$$\mathcal{D} = \{x^{(1)}, x^{(2)}, \dots, x^{(p)}\}, x^{(j)} \in \mathbb{R}^n$$

that have been partitioned according to some unspecified criterion into k subsets,

$$\mathcal{D}_\ell = \{x^{(j)} : j \in I_\ell\}$$

where I_ℓ is the index set corresponding to partitioning.
We introduce the notation

$$II = \{I_1, I_2, \dots, I_k\}$$

and called II a *partitioning* of the data.

The k -means algorithm

For each cluster \mathcal{D}_ℓ , we define an associated characteristic vector $c^{(\ell)} \in \mathbb{R}^n$ that represents the cluster.

The characteristic vector of the cluster makes it possible to define the *within-cluster tightness* or *within-cluster coherence* of \mathcal{D}_ℓ according to the formula

$$q_\ell = q_\ell(c^{(\ell)}) = \sum_{j \in I_\ell} \|x^{(j)} - c^{(\ell)}\|^2$$

where the distance between the data point and characteristic vector is measured in Euclidean norm.

The quantity of q_ℓ is sometimes referred to as the within-cluster sum of squares(WCSS). Initially, a small WCSS should be taken as an indication of a tight cluster for which the characteristic vector is a good representative of each cluster member.

The k -means algorithm

The *overall tightness*, or *overall coherence*, of the clustering \mathbf{II} is measured by the quantity

$$Q(\mathbf{II}, c^{(1)}, c^{(2)}, \dots, c^{(k)}) = \sum_{\ell=1}^k q_{\ell}$$

The k -means algorithm, also known as *Lloyd's algorithm*, is an iterative procedure searching for an optimal clustering \mathbf{II}_{opt} and corresponding characteristic vectors $c_{opt}^{(1)}, c_{opt}^{(2)}, \dots, c_{opt}^{(k)}$ such that

$$Q(\mathbf{II}_{opt}, c_{opt}^{(1)}, c_{opt}^{(2)}, \dots, c_{opt}^{(k)}) = \min Q(\mathbf{II}, c^{(1)}, c^{(2)}, \dots, c^{(k)})$$

where the minimum is taken over all *partitionings* of the data and all characteristic vectors representing the clusters.

An alternating minimization scheme

The k -means algorithm is based on the following alternating optimization scheme:

- 1 **Updating step** : Given the current partitioning \mathbf{II}_t , update the characteristic vectors

$$Q(\mathbf{II}_t, c_{t+1}^{(1)}, c_{t+1}^{(2)}, \dots, c_{t+1}^{(k)}) = \min Q(\mathbf{II}, c^{(1)}, c^{(2)}, \dots, c^{(k)})$$

where the minimization is over the vectors $c^{(\ell)}$.

- 2 **Assignment step** : Given the updated characteristic vectors $c_{t+1}^{(1)}, c_{t+1}^{(2)}, \dots, c_{t+1}^{(k)}$, update the partitioning,

$$Q(\mathbf{II}_{t+1}, c_{t+1}^{(1)}, c_{t+1}^{(2)}, \dots, c_{t+1}^{(k)}) = \min Q(\mathbf{II}_t, c_{t+1}^{(1)}, c_{t+1}^{(2)}, \dots, c_{t+1}^{(k)}),$$

minimizing over different partitionings into k clusters.

An alternating minimization scheme : Updating step

The mapping

$$\mathbb{R}^n \rightarrow \mathbb{R}, c^{(\ell)} \mapsto Q(\mathbf{II}, c^{(1)}, c^{(2)}, \dots, c^{(\ell)}, \dots, c^{(k)}),$$

where $c^{(1)}, c^{(2)}, \dots, c^{(\ell-1)}, c^{(\ell+1)}, \dots, c^{(k)}$, is differential, so a minimizer must be a critical point for the function. In light of the fact that only $q_{(\ell)}$ depends on $c^{(\ell)}$, the critical points with respect to $c^{(\ell)}$ are found by solving the equation

$$\nabla_{c^{(\ell)}} Q(\mathbf{II}, c^{(1)}, c^{(2)}, \dots, c^{(k)}) = \nabla_{c^{(\ell)}} q_{\ell}(c^{(\ell)}) = 0,$$

which can be written componentwise as

$$\begin{aligned} \frac{\partial}{\partial c_i^{(\ell)}} \sum_{j \in \mathbf{I}_{\ell}} \sum_{q=1}^n (x_q^{(j)} - c_q^{(\ell)})^2 &= -2 \sum_{j \in \mathbf{I}_{\ell}} (x_i^{(j)} - c_i^{(\ell)}) \\ &= -2 \left(\sum_{j \in \mathbf{I}_{\ell}} x_i^{(j)} - |\mathcal{D}_{\ell}| c_i^{(\ell)} \right), 1 \leq i \leq n. \end{aligned}$$

An alternating minimization scheme : Updating step

where $|\mathcal{D}_\ell|$ denotes the cardinality of the cluster \mathcal{D}_ℓ , that is, the number of index in the set I_ℓ .

Therefore, from condition gives

$$c^{(\ell)} = \frac{1}{|\mathcal{D}_\ell|} \sum_{j \in I_\ell} x^{(j)}.$$

since the coherence, which is quadratic function of $c^{(\ell)}$, increases without bounded as $c^{(\ell)}$ goes to infinity, its unique critical point must correspond to the global minimum.

An alternating minimization scheme : Assignment step

Consider now the minimization with respect to the partitioning, keeping the characteristic vectors $c^{(\ell)}$ fixed.

Given $x^{(j)} \in \mathcal{D}_\ell$, where \mathcal{D}_ℓ refers to the current partitioning, if

$$\|x^{(j)} - c^{(m)}\| < \|x^{(j)} - c^{(\ell)}\|, \text{ for some } m \neq \ell,$$

then reassigning $x^{(j)}$ to \mathcal{D}_m will decrease the overall tightness.

k -means algorithm(Lloyd's algorithm)

Given the number of clusters $k = 2, 3, \dots$, and a tolerance $\tau > 0$.

Initialize: Assign a partitioning \mathbf{I}_0 , $\Delta Q = \infty$.

while $\Delta Q > \tau$ **do**

Updating step: For each I_ℓ in \mathbf{I}_t , compute the cluster centroid

$$c_{t+1}^{(\ell)} = \frac{1}{|\mathcal{D}_\ell|} \sum_{j \in I_\ell} x^{(j)}.$$

Assignment step: For each $x^{(j)}$, find the closest cluster centroid, and assign $x^{(j)}$ to the corresponding cluster. This defines the new partitioning \mathbf{I}_{t+1} .

 Update

$$\Delta Q = |Q(\mathbf{I}_t, c_t^{(1)}, \dots, c_t^{(j)}) - Q(\mathbf{I}_{t+1}, c_{t+1}^{(1)}, \dots, c_{t+1}^{(j)})|,$$

and advance the counter $t \rightarrow t + 1$.

end while

An alternating minimization scheme

The assignment step in the k -means algorithm can be interpreted in geometric terms as follows. The k -means algorithm induces a *Voronoi tessellation* of the data space \mathbb{R}^n . Given the cluster centroids $c^{(1)}, c^{(2)}, \dots, c^{(k)}$, the ℓ -th Voronoi set V_ℓ is defined by

$$\begin{aligned} V_\ell &= \{x \in \mathbb{R}^n \mid \|x - c^{(\ell)}\| < \|x - c^{(j)}\| \text{ for all } j, \quad j \neq \ell\} \\ &= \bigcap_{j \neq \ell} \{x \in \mathbb{R}^n \mid \|x - c^{(\ell)}\| < \|x - c^{(j)}\|\}. \end{aligned}$$

Hence, the k -means algorithm can be seen as a method to implicitly subdivide the data space into k Voronoi sets, and assign each data point to a cluster according to which Voronoi set the point belongs to, without explicitly computing the tessellation.

Initial partitioning

- 1 *Random initialization:* Divide the index set $I = \{1, 2, \dots, p\}$ into k random, nonempty subsets. One possibility is to first randomly permute the elements of I and then divide the permuted index vector into k subvectors of roughly equal size.
- 2 *Partitioning by hyperplanes:* Subdivide the data space using hyperplanes. The simplest example is for $k = 2$, when a hyperplane can be used to split the data in two initial clusters.
- 3 *Initial seed partitioning:* Select k points from the data set, $x^{(j_1)}, x^{(j_2)}, \dots, x^{(j_k)}$. These *seed vectors* can be chosen either randomly or, if data indicate the presence of clusters, e.g., in a PCA plot, the choice can be one representative for each presumed cluster. Subsequently, partition the data space \mathbb{R}^n into k Voronoi sets V_j defined by seed vectors. The initial clusters are chosen as $I_\ell = \{j \in I \mid x^{(j)} \in V_\ell\}, 1 \leq \ell \leq k$.

The k -medoids algorithm

Denote the distance d between the i th and j th data points as

$$d_{ij} = d(x^{(i)}, x^{(j)}), 1 \leq i, j \leq p,$$

where d is a metric, that is, a binary from $d : \mathbb{R}^n \times \mathbb{R}^n \rightarrow \mathbb{R}_0^+$ such that, for any $x, y, z \in \mathbb{R}^n$,

- ① $d(x, x) = 0$,
- ② $d(x, y) \geq 0$,
- ③ $d(x, y) = d(y, x)$,
- ④ $d(x, y) \leq d(x, z) + d(z, y)$.

Clearly, every norm $\|\cdot\|$ on \mathbb{R}^n defines the metric $d(x, y) = \|x - y\|$.

The k -medoids algorithm

Assume that a partitioning \mathbf{II} of the data into k clusters is given, and associated with each \mathcal{D}_ℓ a characteristic vector *belonging to the cluster*, that is,

$$c^{(\ell)} \in \mathcal{D}_\ell, 1 \leq \ell \leq k.$$

Following what we did for k -means algorithm, we define the within-cluster tightness $q_\ell(c^{(\ell)})$ and the overall tightness $Q(\mathbf{II}, c^{(1)}, \dots, c^{(k)})$ according to the formulas

$$Q(\mathbf{II}, c^{(1)}, \dots, c^{(k)}) = \sum_{\ell=1}^k q_\ell(c^{(\ell)}), \text{ where } q_\ell(c^{(\ell)}) = \sum_{j \in \mathbf{I}_\ell} d(x^{(j)}, c^{(\ell)}).$$

The setting is very similar to the derivation of the k -means algorithm, with the additional requirement that **characteristic vectors must belong themselves to the clusters that represent.**

k -medoids algorithm(partitioning around medoids)

1. **Given** the number of clusters k , a metric d , and a tolerance $\tau > 0$.
2. **Initialize** : Specify the initial set of characteristic vectors, or medoids,

$$\{c_0^{(1)}, c_0^{(2)}, \dots, c_0^{(k)}\}, \quad c_0^{(\ell)} \in \mathcal{D}.$$

Set $t = 0, \Delta Q = \infty$.

3. **Iteration : While** $\Delta Q > \tau$:

Assignment step : For each $x^{(j)}, 1 \leq j \leq p$, find the nearest medoid,

$$d(x^{(j)}, c^{(\ell_*)}) = \min_{1 \leq \ell \leq k} d(x^{(j)}, c^{(\ell)}),$$

and assign $x^{(j)}$ to the cluster \mathcal{D}_{ℓ_*} .

This process determines a partitioning \mathbf{II}_{t+1} into k clusters $\mathcal{D}_1, \dots, \mathcal{D}_k$.

k -medoids algorithm(partitioning around medoids)

Updating step : For each cluster \mathcal{D}_ℓ , $1 \leq \ell \leq k$, calculate the within-cluster tightness

$$q_\ell(x^{(j)}) \text{ for all } x^{(j)} \in \mathcal{D}_\ell,$$

and select the medoid for which the tightness is smallest,

$$c_{t+1}^{(\ell)} = \arg \min \{q_\ell(x^{(j)}) \mid x^{(j)} \in \mathcal{D}_\ell\}.$$

Calculate the overall tightness,

$$Q_{t+1} = Q(\mathbf{I}_{t+1}, c_{t+1}^{(1)}, \dots, c_{t+1}^{(k)}).$$

If $t > 1$, update

$$\Delta Q = |Q_t - Q_{t+1}|,$$

and advance the counter by one, $t \rightarrow t + 1$.

k -means or k -medoids ?

While the idea behind k -means may be very natural and well in line with a mathematical frame of mind, among the arguments in support of k -medoids are the flexibility with respect to how distance between data points is measured and lower sensitivity to presence of cluster outliers that may have a strong adverse effect on the performance of the k -means algorithm.

How to choose k in the algorithm

Consider a data set of p vectors, on which either k -means or k -medoids clustering is applied with a give k , and let the cluster centroids or medoids be $c^{(1)}, c^{(2)}, \dots, c^{(k)}$. The within-cluster mean distance (WCMD) is the average distance of each point from its respective cluster centroid, that is,

$$WCMD_k = \frac{1}{p} \sum_{\ell=1}^k \sum_{j \in I_\ell} d(x^{(j)}, c^{(\ell)}).$$

References

- 1 Daniela Calvetti, Erkki Somersalo, Clustering: K-means and K-medoids. *Mathematics of Data Science: A Computational Approach to Clustering and Classification*.