

Technical notes on the **antyclust** package

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2019-03-14

This document explains the technical and algorithmical background of the R package **antyclust**. **It is a work in progress.** The following topics are covered:

1. A formalization of the anticlustering problem
2. A description of the objective functions used to measure anticluster similarity
3. A documentation of the algorithms used to optimize the objective functions

1 Problem formalization

A set of n d-dimensional data points $X = \{x_i\}$ ($i \in \{1, \dots, n\}$) has to be partitioned into K clusters $C = \{c_k, k = 1, \dots, K\}$, satisfying the following restrictions:

$$\bigcup_{k=1}^K c_k = X \tag{1}$$

$$S_k \cap S_j = \emptyset, \forall k, j \in \{1, \dots, K\}, k \neq j \tag{2}$$

$$|c_k| = |c_j|, \forall k, j \in \{1, \dots, K\} \tag{3}$$

Restriction (1) ensures that each element from the underlying set X is assigned to an anticluster; restriction (2) ensures that each element is assigned to only one anticluster; restriction (3) ensures that each anticluster contains the same number of elements. It follows that $|c_k| = \frac{n}{K} \forall k \in \{1, \dots, K\}$. Restriction (3) is currently implemented for all methods in the **antyclust** package, but it is not an obligatory restriction for anticlustering in general. The objective is to select a partitioning C that maximizes the similarity of the K anticlusters.

2 Objective functions

This section presents definitions of optimal anticluster similarity as assumed in the **antyclust** package.

2.1 The variance objective

Späth (1986) and Valev (1998) independently proposed to maximize the variance criterion used in k-means clustering as the objective in anticlustering. The variance criterion is given by sum of the squared errors between cluster centers (μ_k) and individual data points (Jain, 2010):

$$\sum_{k=1}^K \sum_{x_i \in c_k} \|x_i - \mu_k\|^2 \tag{4}$$

The following plot graphically illustrates efforts to maximize and to minimize the variance criterion in a 2-dimensional feature space for three (anti)clusters, respectively. The partitions employ restrictions (1) - (3), including the restriction of equal (anti)cluster sizes. Optimizing the variance objective is a computationally difficult problem that is usually tackled using heuristic methods (Jain, 2010; Späth, 1986; Valev, 1998).

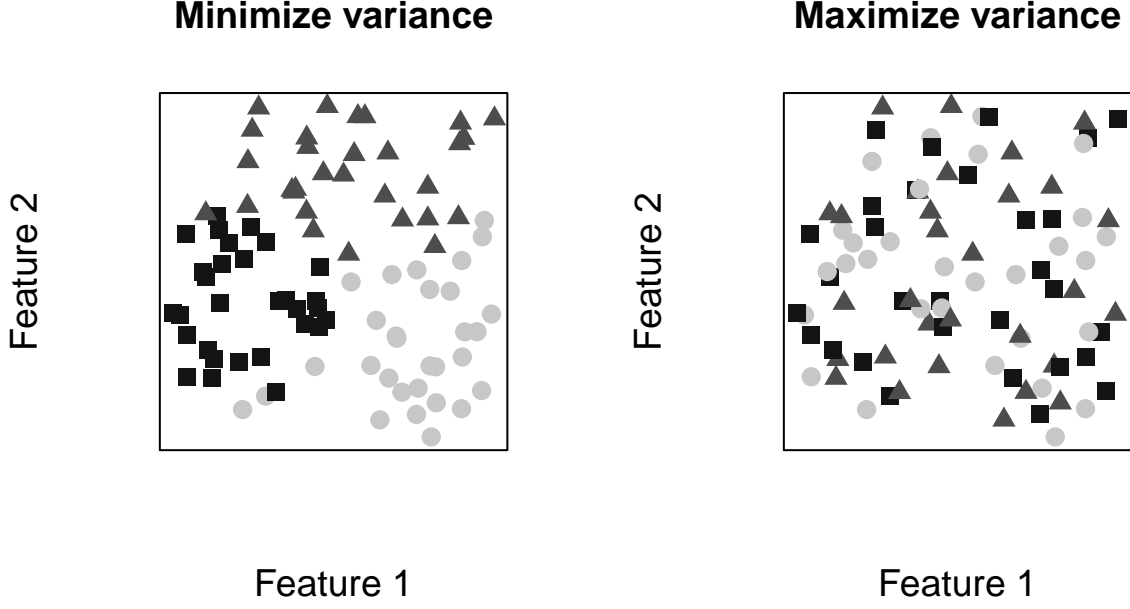


Figure 1: Attempts to minimize and maximize the variance objective, leading to clustering and anticlustering partitions, respectively.

2.2 The distance objective

In addition to the variance criterion, the **anticlust** package introduces another clustering objective to the anticlustering application. The objective has been developed in the problem domain of cluster editing¹ and is based on a measure of the pairwise dissimilarity between data points (Böcker & Baumbach, 2013; Rahmann et al., 2007). In weighted cluster editing, the optimal objective is found when the sum of all pairwise dissimilarities within-clusters is minimized; for the anticlustering application, the objective is maximized instead.

To formalize the cluster editing objective, we use variables x_{ij} to encode whether two data points x_i and x_j belong to the same anticluster c_k :

$$x_{ij} = \begin{cases} 1 & \text{if } x_i \in c_k \wedge x_j \in c_k \\ 0 & \text{otherwise} \end{cases} \quad (5)$$

Assume that d_{ij} represents a measure of the dissimilarity between two data points x_i and x_j , for example given as the euclidean distance. The cluster editing distance objective is then given as follows (Grötschel & Wakabayashi, 1989; Miyauchi & Sukegawa, 2015):

$$\sum_{1 \leq i < j \leq n} d_{ij} x_{ij} \quad (6)$$

Maximizing the distance objective corresponds to minimizing the average linkage distance between anticlusters. In hierarchical cluster algorithms, the average linkage distance is a quantification of the similarity of two clusters (Bacon, 2001). To appreciate the correspondence of the distance objective and the average linkage

¹Cluster editing has also been studied under different names such as correlation clustering (Bansal, Blum, & Chawla, 2004), clique partition problem (Grötschel & Wakabayashi, 1989), and transitivity clustering (Wittkop et al., 2010).

distance, consider the total sum of all pairwise dissimilarities. The total sum can be partitioned into the sum of distances within-clusters and the sum of distances between-clusters:

$$d_{\text{within}} = \sum_{1 \leq i < j \leq n} d_{ij} x_{ij} \quad (7)$$

$$d_{\text{between}} = \sum_{1 \leq i < j \leq n} d_{ij} (1 - x_{ij}) \quad (8)$$

$$d_{\text{total}} = d_{\text{within}} + d_{\text{between}} \quad (9)$$

Because d_{total} is not influenced by the values of x_{ij} , maximizing d_{within} leads to the same values of x_{ij} as minimizing d_{between} . In the special case of $K = 2$ where we have two partitions A and B , d_{between} can be formulated as follows:

$$d_{\text{between}} = \sum_{i \in A} \sum_{j \in B} d_{ij} \quad (10)$$

This formulation is very close to the average linkage distance that additionally incorporates the cardinalities of the sets A and B (Guha, Rastogi, & Shim, 1998):

$$d_{\text{avg}} = \frac{1}{|A| |B|} \sum_{i \in A} \sum_{j \in B} d_{ij} \quad (11)$$

Given restriction (3) for the anticlustering problem, the partitions A and B are of equal size; therefore, d_{between} is minimized whenever d_{avg} is minimized, which also happens when d_{within} is maximized. Thus, the cluster editing objective is a generalization of the average linkage method on more than two clusters.

3 Algorithmic approaches

Finding optimal data partitions usually corresponds to NP-complete problems (Arabie & Hubert, 1996; Bansal et al., 2004; Jain, 2010). For NP-complete problems, it is often infeasible to find the optimal objective, especially when n is large. To find the optimal solution for moderately large instances, **anticlust** employs integer linear programming. To process larger problem instances, **anticlust** uses heuristic methods based on repeated random sampling.

3.1 NP-completeness

In the following, I show that distance anticlustering is NP-complete. First, distance anticlustering is in NP because the distance objective can be computed in polynomial time for a given partitioning; the summation of all distance values d_{ij} is in $O(n^2)$.

Second, I show that if an efficient algorithm exists to solve distance anticlustering in polynomial time, it is also possible to solve the NP-complete balanced number partitioning problem in polynomial time (Mertens, 2001). In the number partitioning problem, we have a list of positive integers a_1, a_2, \dots, a_n and try to find a subset $A \subset \{1, \dots, n\}$ that minimizes the partition difference

$$E(A) = \left| \sum_{i \in A} a_i - \sum_{j \notin A} a_j \right| \quad (12)$$

In the balanced version of number partitioning, we may impose the restriction of $|A| = \frac{n}{2}$ – assuming that n is even – corresponding to restriction (3) of equal cluster sizes in anticlustering (Mertens, 2001).

To convert the number partitioning formulation into a formulation of distance anticlustering, we define d_{ij} as the absolute difference representing the dissimilarity of two numbers:

$$d_{ij} := |a_i - a_j| \quad (13)$$

We thus obtain

$$E(A) = \sum_{i \in A} \sum_{j \notin A} d_{ij} \quad (14)$$

Using variables $x_{ij} \in \{0, 1\}$ to represent whether two numbers are both either in A or not, i.e.,

$$x_{ij} = \begin{cases} 1 & \text{if } (x_i \in A \wedge x_j \in A) \vee (x_i \notin A \wedge x_j \notin A) \\ 0 & \text{otherwise} \end{cases}$$

we obtain $E(A)$ as the distance anticlustering objective:

$$E(A) = \sum_{1 \leq i < j \leq n} d_{ij} x_{ij} \quad (15)$$

Hence, balanced number partitioning is a special case of distance anticlustering where

- a) $K = 2$
- b) each element is described by exactly one integer
- c) d_{ij} is the absolute difference

If a polynomial-time algorithm exists that solves distance anticlustering, we can therefore solve the NP-complete balanced number partitioning in polynomial time. Hence, distance anticlustering is NP-complete.

3.2 Integer linear programming

Despite the NP-complete nature of cluster editing, integer linear programming has been successfully used to find optimal solutions even for relatively large problem instances (Böcker, Briesemeister, & Klau, 2011; L. H. N. Lorena, Quiles, Carvalho, & Lorena, 2018). Integer linear programming identifies the values of *decision variables* that optimize a linear objective function, while employing constraints on the decision variables that are implemented as mathematical inequalities. In the case of anticlustering, integer linear programming can be used to maximize the distance objective in (6) while constraints ensure that the anticlustering conditions in (1) - (3) are met.

For the integer linear programming formulation of distance anticlustering, a problem instance is represented as an undirected complete graph $G = (V, E)$ (cf. Schaeffer, 2007). Each vertex $v \in V = (1, \dots, n)$ represents an element from the input data that has to be assigned to an anticluster. Edges are unordered pairs $\{i, j\} \in E$; the short form ij will be used to refer to edges. A cost function $w : E \rightarrow \mathbb{R}$ assigns a weight to the edges where each weight is given by the distance between the two elements that are connected by the edge, i.e., $w(ij) = d_{ij}$.

The integer linear program returns a subgraph $G' = (V, E')$. For G' , we use decision variables x_{ij} to encode whether two vertices i and j are connected by an edge:

$$x_{ij} = \begin{cases} 1 & \text{if } ij \in E' \\ 0 & \text{otherwise} \end{cases}$$

If two vertices are connected by an edge in G' , the integer linear program ensures that they are also part of a *clique*, that is, a subgraph whose vertices are all connected. Cliques in G' represent anticlusters. Hence, the following two conditions are equivalent:

1. $x_{ij} = 1$
2. Vertices i and j belong to the same anticluster.

The **anticlust** package employs an integer linear program on the basis of formulations that Grötschel and Wakabayashi (1989) introduced to solve cluster editing:

$$\text{Maximize } \sum_{1 \leq i < j \leq n} w(ij) x_{ij} \quad (16)$$

$$-x_{ij} + x_{ik} + x_{jk} \leq 1, \quad \forall 1 \leq i < j < k \leq n, \quad (17)$$

$$x_{ij} - x_{ik} + x_{jk} \leq 1, \quad \forall 1 \leq i < j < k \leq n, \quad (18)$$

$$x_{ij} + x_{ik} - x_{jk} \leq 1, \quad \forall 1 \leq i < j < k \leq n, \quad (19)$$

$$\sum_{1 \leq i < j \leq n} x_{ij} + \sum_{1 \leq k < i \leq n} x_{ki} = \frac{n}{K} - 1, \quad \forall i \in \{1, \dots, n\} \quad (20)$$

$$x_{ij} \in \{0, 1\}, \quad \forall 1 \leq i < j \leq n \quad (21)$$

The inequalities (17) - (19) are called triangular constraints; they ensure that the output graph is a union of disjoint cliques (Grötschel & Wakabayashi, 1989). Constraint (20) ensures that K cliques are returned, each of cardinality $\frac{n}{K}$. Constraint (21) ensures that the decision variables x_{ij} are binary. Cliques are created under the objective to maximize the sum of the edge weights within cliques, i.e., the distance objective is maximized.

The **anticlust** package can use one of the commercial solvers **gurobi** or **CPLEX** or the open source GNU linear programming kit to solve the integer linear program defined in (16) - (21) optimally.

3.2.1 Preprocessing

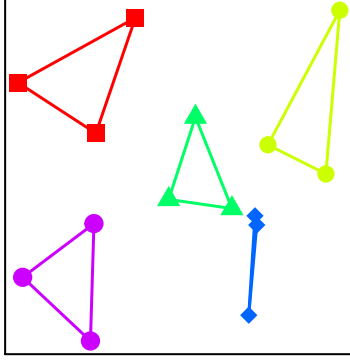
To expand its applicability to larger problem sizes, additional constraints can be added to the integer linear program through a redefinition of the cost function w . This approach results from two considerations:

1. The run time of exact weighted cluster editing is improved if there is an uneven distribution of edge weights (Böcker & Baumbach, 2013; Böcker et al., 2011)
2. It is possible to prevent very similar elements from joining the same anticluster without strongly impairing the quality of the solution

In the cluster editing integer linear programming framework, setting $w(ij) = -\infty$ prevents two vertices i and j from joining the same clique in the output graph (Böcker et al., 2011). This recalculation induces a strong unevenness among edge weights, increasing the problem sizes the integer linear program can be applied to. If only very similar elements are prevented from joining the same anticluster, the quality of the solution is not strongly impaired because the anticlustering objective is based on the idea that similar elements should be in different anticlusters.

The **anticlust** package may realize a preprocessing step that groups very similar elements into *preclusters*; elements of the same precluster are prevented from joining the same anticluster thereafter. The preclustering step is formalized by the following integer linear program:

Preclustering



Anticlustering

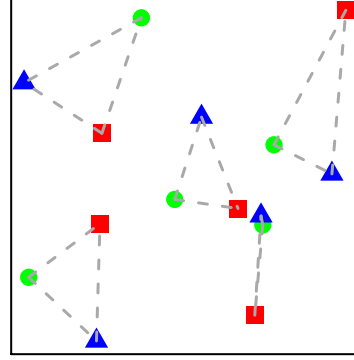


Figure 2: The left-hand plot shows the preclusters each consisting of three elements. The right-hand plot shows the optimal anticlustering under the preclustering restrictions. Elements that are part of the same precluster cannot be part of the same anticluster

$$\text{Minimize } \sum_{1 \leq i < j \leq n} w(ij) x_{ij} \quad (22)$$

$$-x_{ij} + x_{ik} + x_{jk} \leq 1, \quad \forall 1 \leq i < j < k \leq n, \quad (23)$$

$$x_{ij} - x_{ik} + x_{jk} \leq 1, \quad \forall 1 \leq i < j < k \leq n, \quad (24)$$

$$x_{ij} + x_{ik} - x_{jk} \leq 1, \quad \forall 1 \leq i < j < k \leq n, \quad (25)$$

$$\sum_{1 \leq i < j \leq n} x_{ij} + \sum_{1 \leq k < i \leq n} x_{ki} = K - 1, \quad \forall i \in \{1, \dots, n\} \quad (26)$$

$$x_{ij} \in \{0, 1\}, \quad \forall 1 \leq i < j \leq n \quad (27)$$

By minimizing the distance objective (22), the preclustering step solves weighted cluster editing under the restriction of a cluster size K , resulting in $\frac{n}{K}$ clusters.² The number of elements per precluster corresponds to the number of anticlusters. For $K = 2$, the preclustering corresponds to the minimum weight perfect matching problem (Gerards, 1995). The left-hand plot in Figure 2 illustrates the preprocessing step for $K = 3$ and $n = 15$.

In a second step, the cost function is redefined to prevent elements from the same precluster from joining the same anticluster:

$$w(ij) = \begin{cases} -\infty & \text{if } x_{ij} = 1 \\ d_{ij} & \text{if } x_{ij} = 0 \end{cases}$$

Using the redefined edge weights, `antyclust` solves distance anticlustering using the integer linear program defined in (16) - (21). An example of anticlustering employing preclustering restrictions is shown in the right-hand plot of Figure 2. Because the optimal solution sometimes requires to join elements that are part of the same precluster, the preprocessing sometimes precludes an optimal solution.

²This is a feasible preprocessing step because minimizing the distance objective works faster than maximizing the distance objective when group sizes are equal. That is, cluster editing is solved more efficiently using integer linear programming than anticlustering.

3.3 Heuristic anticlustering

To solve larger problem instances that cannot be processed using integer linear programming, a heuristic method based on random sampling is employed in the `anticlust` package. The heuristic may incorporate a preclustering step that often improves the quality of the output. Across a user-specified number of runs, each element is first randomly assigned to an anticluster, then the objective value is computed. In the end, the best assignment is returned as the output. When preclustering is employed, the random assignment is conducted under the restriction that preclustered elements cannot be part of the same anticluster.

3.3.1 Heuristic preclustering

A heuristic cluster algorithm is employed as a preclustering step for the random search procedures. The algorithm is based on k-means clustering but enforces equal cluster sizes. First, $\frac{n}{K}$ cluster centers are initialed using k-means. Then, elements are sequentially assigned to the nearest cluster centers while ensuring that each of the clusters is filled with K elements. The following pseudo code formalizes the heuristic:

```
E = list of n elements
C = list of K/n cluster centers, initialized using k-means
Repeat K times {
  S <- random sequence of C
  for (j in S) {
    1. e = element that is closest to cluster center j
    2. Add e to cluster corresponding to cluster center j
    3. Remove e from E
  }
}
```

4 References

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