DISTANCES, ORDERS AND SPACES

Pascal Préa

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Pascal Préa [1,2]

[1] Aix-Marseille Université, CNRS, LIS, Marseille, France
(pascal.prea@lis-lab.fr)

[2] École Centrale de Marseille, Marseille, France

ABSTRACT: A dissimilarity d on a set X is said to be *Robinson* if there exists a total order, said *compatible*, on X such that $x < y < z \implies d(x,z) \geq \max\{d(x,y), d(y,z)\}$. Roughly speaking, d is Robinson if the points of X can be represented on a line *ie.* Robinson dissimilarities generalize line distances.

In this paper, we define k-dimensional Robinson dissimilarities, which generalize the possibility, for a metric set (X,d), to be embedded into a k-dimensional Euclidean space. This generalization is more flexible than the classical embedding and we show that every dissimilarity on an n-set X is $(\log n)$-dimensional Robinson. We give an $O(n^3)$ algorithm which builds such an embedding. This algorithm is based on an incremental algorithm to recognize Robinson dissimilarities.

KEYWORDS: Robinson dissimilarities, embeddings, incremental algorithms.

1 Introduction

Given a finite set set X, a *dissimilarity* on X is a symmetrical function $X \times X \longmapsto \mathbb{R}^+$ such that $\forall x \in X, d(x,x) = 0$ (we say that (X,d) is a dissimilarity space). Dissimilarities generalize distances (a distance is dissimilarity with the triangular inequality).

Given a dissimilarity on a set X, a fundamental problem is to derive "geometrical" properties of X from d, or to characterize dissimilarities from which such properties can be obtained. For instance Robinson dissimilarities (Robinson 1951) correspond to points on a line. These dissimilarities were invented to solve seriation problems in Archeology, but they are now a classical tool for seriation problems in any field. They are also linked with Pyramids (Diday 1986, Durand & Fichet 1988), the standard model with overlapping classes. Moreover, they play an important role ro recognize tractable cases for TSP (Çela & al. 2023).

In this paper, we generalize Robinson dissimilarities to *k(-dimensional)-Robinson* dissimilarities, which represent the fact for X to be embedded into a k-dimensional space. This embedding is less strict than the usual Euclidean

embedding. We show that, if d is a dissimilarity on a set X with $|X| = n$, then d is $(\log n)$-Robinson and we give a $O(n^3)$ algorithm which builds such an embedding. This algorithm is based on an incremental algorithm to recognize Robinson dissimilarities which is presented in the last section.

2 Robinson dissimilarities

A dissimilarity space (X,d) is *Robinson* if there exists a total order, which is said to be *compatible*, on X such that

$$x < y < z \implies d(x,z) \geq \max\{d(x,y), d(y,z)\} \tag{1}$$

Let (X,d) a dissimilarity space and $<$ be an order on X. Notice that $<$ is a compatible order of (X,d) (which is thus a Robinson space) if and only if:

$$x \leq y < z \leq t \implies d(y,z) \leq d(x,t) \tag{2}$$

Given a total order $<$ on X and $x,y,z \in X$, we say that y is *between* x and z for $<$ if $x < y < z$ or $z < y < x$. The set of the elements between x and z is an *interval* for $<$ and we denote it by $[x,z]_<$. Notice that $[x,z]_< = [z,x]_<$.

3 Multidimensional Robinson dissimilarities

Let (X,d) a dissimilarity space and $k \in \mathbb{N}_1$. We say that (X,d) is *k-Robinson* if there exist k orders $<_1, <_2, \ldots, <_k$ such that:

$$\forall x,y,z,t \in X, (\forall 1 \leq i \leq k, y,z \in [x,t]_{<_i}) \implies d(y,z) \leq d(x,t)$$

We say that (X,d) is *k-quasi-Robinson* if there exist k orders $<_1, <_2, \ldots, <_k$ such that:

$$\forall x,y,z \in X, (\forall 1 \leq i \leq k, y \in [x,z]_{<_i}) \implies d(x,z) \geq \max\{d(x,y), d(y,z)\}$$

If $k = 1$, it is equivalent for a dissimilarity space to be Robinson or 1-quasi-Robinson. For $k \geq 2$, then if (X,d) is k-Robinson, then (X,d) is k-quasi-Robinson, but the converse is false (see Figure 1). Notice in addition that, if (X,d) is k-(quasi-)Robinson, then (X,d) is $k+1$-(quasi-)Robinson. The smallest k such that (X,d) is the *Robinson dimension* of (X,d).

If a metric space (X,d) can be embedded into a $\mathbb{R}^k$, then (X,d) is k-Robinson. But the Robinson dimension of (X,d) is generally smaller. For instance, if $|X| = n$ and d is the constant dissimilarity, then (X,d) is Robinson (its Robinson dimension is 1) although it needs an $n-1$-dimensional Euclidean space to be embedded. Moreover, we have:

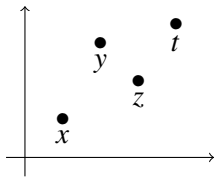

Figure 1. *A set X with four points x,y,z,t. If (X,d) is 2-quasi-Robinson with the two orders represented by the two axis, then no condition is imposed on $d(y,z)$ and we can set $d(y,z) > d(x,t)$. If (X,d) is 2-Robinson (with the same orders), then $d(y,z) \leq d(x,t)$.*

Proposition 1 *The Robinson dimension of a dissimilarity space (X,d) with $|X| = n$ is $\leq \lceil \log_2 \lceil \frac{n}{3} \rceil \rceil + 1$.*

Algorithm 1 returns an approximate value for the Robinson dimension of a dissimilarity space.

Algorithm 1: APPROXIMATE-ROBINSON-DIMENSION

Input: (X,d), a dissimilarity space.
Output: An upper bound on the Robinson dimension of (X,d).
begin
  $X' \leftarrow X$; $k \leftarrow 0$;
  SORT-LINES(X,d) ;
  while $X' \neq \emptyset$ **do**
    $S \leftarrow$ MAXIMAL-ROBINSON-SUBSPACE(X',d) ;
    $X' \leftarrow X' \setminus S$;
    $k \leftarrow k+1$;
  return $\lceil \log_2 k \rceil + 1$;

The function SORT-LINES(X,d), for every $x \in X$, sorts the points of X by increasing values of their distance from x. This function runs in $O(n^2 \log n)$ where $n = |X|$. The function MAXIMAL-ROBINSON-SUBSPACE returns a subset S of X', maximal for inclusion and such that (S,d) is Robinson. This can be easily implemented by a greedy algorithm. We will see in Section 4 that, after SORT-LINES, such a greedy version of MAXIMAL-ROBINSON-SUBSPACE runs in $O(|X'|^2)$. So, as there is at most $n/3$ iterations of the **while** loop, Algorithm 1 runs in $O(n^3)$.

4 An incremental algorithm to recognize Robinson dissimilarities

In order to implement MAXIMAL-ROBINSON-SUBSPACE, we need a function ADD-AND-TEST which takes as entry a dissimilarity space (X,d), a set $S \subset X$ such that (S,d) is Robinson, the PQ-tree $\mathcal{T}_P(S,d)$ and a point $x \in X \setminus S$. A *PQ-tree* (Booth & Lueker 1976) is a data structure which can encode all the compatible orders of a Robinson dissimilarity. ADD-AND-TEST returns the PQ-tree $\mathcal{T}_P(S \cup \{x\},d)$ (If $(S \cup \{x\},d)$ is not Robinson, then $\mathcal{T}_P(S \cup \{x\},d)$ = **none**). The algorithm of ADD-AND-TEST can be sketched as follows:

1. Compute the sets $B_\delta^S := B_\delta(x) \cap S$.
2. Insert the sets B_δ^S into $\mathcal{T}_\mathcal{P}(S,d)$. We get a PQ-tree $\mathcal{T}_\mathcal{P}{}^x(S,d)$.
3. Add the point x to $\mathcal{T}_\mathcal{P}{}^x(S,d)$. We get the PQ-tree $\mathcal{T}_\mathcal{P}(S \cup \{x\},d)$. This will be done in two steps:
   (a) Consider only the points of S the closest from x.
   (b) Consider the other points of S.

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