

Labeling Isometric and Almost Isometric n -Point Configurations in \mathbb{R}^D

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July 2016

Abstract

Consider two n -point configurations X and Y in \mathbb{R}^D with the same distance distribution and distinct distances. We begin with proposing an algorithm that tries to find a bijection $\psi : X \rightarrow Y$ such that X and Y are congruent under ψ , or determines that no such bijection exists. Next, we discuss extending this algorithm to the case where X and Y have small multiplicities of recurring distances, then to the case where X and Y have “almost the same” distance distribution.

1 Introduction

1.1 Background

We are working towards the extension of the Procrustes Problem, which was solved by Peter Schönemann [5].

Theorem 1.1 (Procrustes Problem). *If we have*

$$X = \{x_1, \dots, x_n\}, Y = \{y_1, \dots, y_n\} \subset \mathbb{R}^D$$

such that

$$|x_i - x_j| = |y_i - y_j| \text{ for } i, j = 1, \dots, n. \quad (\dagger)$$

Then there exists $T \in O(D)$ and $\bar{y} \in \mathbb{R}^D$ such that $y_i = T(x_i) + \bar{y}$ for $i = 1, \dots, n$. $O(D)$ refers to the orthogonal group in dimension D .

1.2 Previous Results

Particularly, we're interested in extending the Procrustes by relaxing the (†) condition. There are two ways of approaching this relaxation:

1. $\{|x_i - x_j| : i \neq j\} = \{|y_i - y_j| : i \neq j\}$;
2. $|x_i - x_j| \approx |y_i - y_j|$ for $i, j = 1, \dots, n$.

[1] and [4] studied the former relaxation with some constraints. In both of their approaches, the first step is to find a suitable labeling of the points such that the conditions of the Procrustes Problem are held under the labeling. In other words, their goal is first to find a labeling, then find the $T \in O(D)$. Overall, this relaxation transforms the Procrustes Problem into:

Problem 1.1 (Unlabeled Procrustes Problem). *If we have*

$$X = \{x_1, \dots, x_n\}, Y = \{y_1, \dots, y_n\} \subset \mathbb{R}^D$$

such that $|x_i - x_j| = |y_{\pi(i)} - y_{\pi(j)}|$ for $i, j = 1, \dots, n$ and $\pi \in \text{Sym}_n$, then does there exist π and $T \in O(D)$ and $\bar{y} \in \mathbb{R}^D$ such that

$$y_{\pi(i)} = T(x_i) + \bar{y}$$

for $i = 1, \dots, n$?

[1] noticed that it is possible that no such π or T exists. [2] explores that exact case.

The latter relaxation was studied in [3] and yielded the following result:

Theorem 1.2. *Let $\{x_1, \dots, x_n\}$ and $\{y_1, \dots, y_n\}$ be two n -point configurations in \mathbb{R}^D with distinct x_i, y_i respectively. Suppose*

$$(1 + \delta)^{-1} \leq \frac{|y_i - y_j|}{|x_i - x_j|} \leq 1 + \delta, \forall i \neq j.$$

For any $\varepsilon > 0$, there exists $\delta > 0$ and a Euclidean motion $\Phi_0 : x \rightarrow Tx + x_0$ such that

$$|y_i - \Phi_0(x_i)| \leq \varepsilon \text{diam} \{x_1, \dots, x_n\}$$

for $i = 1, \dots, n$.

We explore an alternative way of matching the points with the constraint that the pairwise distances are distinct.

2 Matching Two n -Point Configurations with same Distance Distribution

2.1 Preliminaries

Definition 2.1. *If we have an n -point configuration A in a metric space (M, d) , we define*

$$\text{dist}(A) := \{d(p, q) \mid p \neq q \in A\}.$$

Definition 2.2. *Let A be an n -point configuration in a metric space (M, d) and p, q be points in A . The **edge between p and q** is the tuple*

$$e_{p,q} := \left(\{p, q\}, d(p, q) \right).$$

We denote $e_{p,q}[1] = \{p, q\}$ and $e_{p,q}[2] = d(p, q)$.

Note $e_{p,q} = e_{q,p}$.

Definition 2.3. *Let A be an n -point configuration in a metric space (M, d) , the **edge set of A** is defined as*

$$E(A) := \{e_{p,q} \mid p \neq q \in A\}.$$

Note if $|A| = n$, then $|E(A)| = \binom{n}{2}$.

Definition 2.4. *An n -point configuration, $A \subset (M, d)$, has **distinct distances** if for every $e, e' \in E(A)$, $e[1] = e'[1]$ if and only if $e[2] = e'[2]$.*

Definition 2.5. *If $A = \{a_1, \dots, a_n\} \subset (M, d)$ is an n -point configuration, the **distance matrix of A** , $\text{distmat}(A)$, is the $n \times n$ matrix*

$$(\text{distmat}(A))_{ij} = d(a_i, a_j).$$

Note $\text{distmat}(A)$ is an $n \times n$ symmetric matrix due to the definition of a metric. Its diagonal entries are all zero.

Our labeling problem can be reformulated as:

Problem 2.1 (Reformulated Unlabeled Procrustes Problem). *Let X and Y n -point configurations such that they have distinct distances and $\text{dist}(X) = \text{dist}(Y)$. We want $\pi \in \text{Sym}_n$ such that $|x_i - x_j| = |y_{\pi(i)} - y_{\pi(j)}|$ for all $i, j = 1, \dots, n$ if such a π exists.*

2.2 The Matching Algorithm

Algorithm 1 This Algorithm assumes the assumptions of **Problem 2** and returns a labeling π , or returns NULL if no labeling exists.

Require: $X, Y \subset \mathbb{R}^D$ have distinct distances, $|X| = |Y|$, $\text{dist}(X) = \text{dist}(Y)$

- 1: **function** DISTINCTDISTANCEMATCHING(X, Y)
- 2: $n \leftarrow |X| = |Y|$
- 3: $\text{distmat}(X) \leftarrow \text{DISTMAT}(X)$
- 4: $\text{distmat}(Y) \leftarrow \text{DISTMAT}(Y)$
- 5: $E(X) \leftarrow \{e_{x_1, x_2} \mid x_1 \neq x_2 \in X\}$
- 6: $E(Y) \leftarrow \{e_{y_1, y_2} \mid y_1 \neq y_2 \in Y\}$
- 7: $\text{SORT}(E(X), \prec), \text{SORT}(E(Y), \prec) \quad \triangleright (e_1 \prec e_2 \Leftrightarrow e_1[2] < e_2[2])$
- 8: $\varphi : E(X) \rightarrow E(Y)$ such that $E(X)[i] \mapsto E(Y)[i], \forall i \in \{1, \dots, \binom{n}{2}\}$
- 9: $\text{first_iteration} = \text{true}$
- 10: **for** $x' \in X$ **do**
- 11: $E_{x'} \leftarrow \{e_{x, x'} \mid x \neq x' \in X\}$
- 12: $y' \leftarrow \bigcap_{e' \in \varphi(E_{x'})} e'[1]$
- 13: **if** $y' = \emptyset$ **then**
- 14: **return** NULL
- 15: $\psi : X \rightarrow Y$ such that $x' \mapsto y'$ and $e[1] \setminus x' \mapsto \varphi(e)[1] \setminus y', \forall e \in E_{x'}$
- 16: $\psi^* \leftarrow \psi$
- 17: **if** first_iteration **then**
- 18: **for** $x \in X$ **do**
- 19: **if** $\psi^* \neq \psi$ **then**
- 20: **return** NULL
- 21: $\text{first_iteration} = \text{false}$
- 22: ψ naturally induces a $\pi \in \text{Sym}_n \quad \triangleright (\text{since } \psi \text{ is bijective})$
- 23: **return** π

3 Matching Two n -Point Configurations with almost small multiplicities of Recurring Distances

In this section, we still assume that X and Y are still n -point configurations with the property $\text{dist}(X) = \text{dist}(Y)$. However, we will slightly relax the constraint from the previous section that X and Y have distinct distances.

Definition 3.1. Let $A \in \mathbb{R}^D$ be an n -point configuration. We say that A has **recurring distance** $d \in \text{dist}(A)$ if there exists $m \in \mathbb{N}$ such that $m > 1$ and $\underbrace{\{d, \dots, d\}}_m \subseteq \text{dist}(A)$. The maximal possible m is the **multiplicity** of d .

Informally, a recurring distance, d , is a value that appears more than once in $\text{dist}(A)$ and its multiplicity is exactly how many d occur in $\text{dist}(A)$.

Because $\text{dist}(X) = \text{dist}(Y)$, X and Y share the same number of recurring distances as well as the same multiplicities for every recurring distance. From now on, we will let $r \in \mathbb{N}$ denote how many recurring distances there are in X and Y .

We also give the following order on the recurring distances:

$$d_k \text{ denotes the } k^{\text{th}} \text{ smallest recurring distance.} \quad (\ddagger)$$

Notice that (\ddagger) gives the following ordering:

$$d_1 < d_2 < \dots < d_{r-1} < d_r.$$

Likewise, we let m_k denote the multiplicity of d_k for each $k \in \{1, \dots, r\}$. For this section, we assume that each multiplicity is *small*, i.e. $m_k \ll n$ for all $k \in \{1, \dots, r\}$.

Example 3.1. Let's take $A \subset \mathbb{R}^D$ with the following with the following hypothetical distance distribution:

$$\text{dist}(A) = \{1, 2, 2, 2, 2, 3, 4, 4, 4, 5, 5, 6\}.$$

Notice the numbers 2, 4, and 5 are recurring.

So the number of recurring distance in A is 3.

We also see $d_1 = 2$, $d_2 = 4$, and $d_3 = 5$.

Finally, $m_1 = 4$, $m_2 = 3$, $m_3 = 2$.

Next, we consider the edges with these recurring distances.

Definition 3.2. Let $A \in \mathbb{R}^D$ be an n -point configuration with r_A recurring distances. For each $k \in \{1, \dots, r_A\}$, we define the set $E_{d_k}(A) \subseteq E(A)$ such that

$$E_{d_k}(A) = \{e \in E(A) \mid e[2] = d_k\}.$$

We also define the ordered set $R(A)$ such that

$$R(A)[k] = E_{d_k}(A), \quad \forall k \in \{1, \dots, r_A\}.$$

In our two n -point configuration case concerning X and Y , it follows from definition that:

- $|R(X)| = |R(Y)| = r$;
- $|E_{d_x}(X)| = |E_{d_k}(Y)| = m_k$ for each $k \in \{1, \dots, r\}$.

Notice that for each $k \in \{1, \dots, r\}$, $E_{d_x}(X)$ is bijective to $E_{d_k}(Y)$ under a total of $m_k!$ labelings. In other words, any $\sigma_k \in \text{Sym}_{m_k}$ induces a bijection, $\varphi_k : E_{d_x}(X) \rightarrow E_{d_k}(Y)$. Given $R(X)$ and $R(Y)$, we can define a bijection $\varphi : E(X) \rightarrow E(Y)$ such that

$$\varphi(e_x) = \begin{cases} e_y \text{ where } e_x[2] = e_y[2] & \text{if } e[2] \text{ is a nonrecurring distance} \\ \varphi_k(e_x), \text{ for } k \text{ where } e_x \in E_{d_x}(X) & \text{if } e[2] \text{ is a recurring distance} \end{cases}.$$

Notice that there are total of $\prod_{k=1}^r (m_k!)$ different φ .

Algorithm 2 This Algorithm assumes X and Y are n -point configurations such that X and Y possess small recurring distance multiplicities and $\text{dist}(X) = \text{dist}(Y)$. It returns a labeling π , or returns NULL if no such labeling exists.

Require: $X, Y \subset \mathbb{R}^D$ with small multiplicities of recurring distances, $|X| = |Y|$, $\text{dist}(X) = \text{dist}(Y)$

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1: function DISTINCTDISTANCEMATCHING2( $X, Y$ )
2:    $n \leftarrow |X| = |Y|$ 
3:    $\text{distmat}(X) \leftarrow \text{DISTMAT}(X)$ 
4:    $\text{distmat}(Y) \leftarrow \text{DISTMAT}(Y)$ 
5:    $E(X) \leftarrow \{e_{x_1, x_2} \mid x_1 \neq x_2 \in X\}$ 
6:    $E(Y) \leftarrow \{e_{y_1, y_2} \mid y_1 \neq y_2 \in Y\}$ 
7:    $\text{SORT}(E(X), \prec), \text{SORT}(E(Y), \prec) \quad \triangleright (e_1 \prec e_2 \Leftrightarrow e_1[2] < e_2[2])$ 
8:   for each  $\prod_{k=1}^r (m_k!)$  permutations of the recurring distances do
9:     Take the corresponding bijection  $\varphi : E(X) \rightarrow E(Y)$  where

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$$E(X)[i] \mapsto \begin{cases} E(Y)[i] & \text{for nonrecurring distances} \\ \varphi_k(E(X)[i]), \text{ for } k \text{ where } e_x \in E_{d_x}(X) & \text{for recurring distances} \end{cases}$$

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10:   first_iteration = true
11:   for  $x' \in X$  do
12:      $E_{x'} \leftarrow \{e_{x, x'} \mid x \neq x' \in X\}$ 
13:      $y' \leftarrow \bigcap_{e' \in \varphi(E_{x'})} e'[1]$ 
14:     if  $y' = \emptyset$  then
15:       break to next iteration of outer loop
16:      $\psi : X \rightarrow Y$  such that  $\begin{cases} x' \mapsto y' \\ e[1] \setminus x' \mapsto \varphi(e)[1] \setminus y', \quad \forall e \in E_{x'} \end{cases}$ 
17:      $\psi^* \leftarrow \psi$ 
18:     if first_iteration then
19:       for  $x \in X$  do
20:         if  $\psi^* \neq \psi$  then
21:           break to next iteration of outer loop
22:       first_iteration = false
23:      $\psi$  naturally induces a  $\pi \in \text{Sym}_n \quad \triangleright (\text{since } \psi \text{ is bijective})$ 
24:     return  $\pi$ 
25: return NULL

```

4 Matching Two n -Point Configurations with “Almost-Same Distance Distribution”

From [3] we have the following definition for “Almost-Same Distance Distribution”:

Definition 4.1. *We say that X and Y have **almost-same distance distribution**, denoted $\text{dist}(X) \approx \text{dist}(Y)$, if for each $dx_{i,j} \in \text{dist}(X)$, there exists exactly one $dy_{i',j'} \in \text{dist}(Y)$ such that $(1 - \epsilon) < \frac{dx_{i,j}}{dy_{i',j'}} < (1 + \epsilon)$ for some ϵ .*

This condition, if met, would induce a very natural edge bijection. Then we can apply **Algorithm 1**. The following is an algorithm that checks this condition:

Algorithm 3 This Algorithm checks the condition of **Definition 9**.

Require: $X, Y \subset \mathbb{R}^D$

```

1: function ALMOSTSAMEDISTANCE( $X, Y$ )
2:    $E(X) \leftarrow \{e_{x_1, x_2} \mid x_1 \neq x_2 \in X\}$ 
3:    $E(Y) \leftarrow \{e_{y_1, y_2} \mid y_1 \neq y_2 \in Y\}$ 
4:   for  $e_X \in E(X)$  do
5:      $\Phi_X(e_X) := \underset{e_Y \in Y}{\text{argmin}} |e_X[2] - e_Y[2]|$ 
6:   for  $e_Y \in E(Y)$  do
7:      $\Phi_Y(e_Y) := \underset{e_X \in X}{\text{argmin}} |e_Y[2] - e_X[2]|$ 
8:   for  $e_X \in E(X)$  do
9:     if  $e_X \neq \Phi_Y(\Phi_X(e_X))$  then
10:      return FALSE
11:   for  $e_Y \in E(Y)$  do
12:     if  $e_Y \neq \Phi_X(\Phi_Y(e_Y))$  then
13:      return FALSE
14:   return  $\Phi_X$ 

```

5 Performance

We assume D is fixed and constant. We also assume that the sorting algorithm used sorts k items in $\mathcal{O}(k \log k)$ time.

5.1 Algorithm 1

LINE 2 through LINE 6 each take $\mathcal{O}n^2$ time. The sorting on LINE 7 takes $\mathcal{O}(n^2 \log n^2) = \mathcal{O}(n^2 \log n)$ time. The loop on LINE 10 takes $\mathcal{O}(n^2)$ time.

Algorithm 1 takes $\mathcal{O}(n^2 \log n)$ time.

5.2 Algorithm 2

Algorithm 2 is similar to **Algorithm 1** but iterates the loop on LINE 10 to a maximum of $\prod_{k=1}^r (m_k!)$ iterations.

Algorithm 2 takes $\mathcal{O}\left(n^2 \log n + n^2 \prod_{k=1}^r (m_k!)\right)$ time.

5.3 Algorithm 3

We know that $|E(X)| = |E(Y)| = \binom{n}{2} = \mathcal{O}(n^2)$. So the bulk of the work of this algorithm is done in the loops on LINE 6 and LINE 8. They both take $\mathcal{O}(n^2)$ edges and compares each edge with the other $\mathcal{O}(n^2)$ edges, both of which take $\mathcal{O}(n^4)$ time.

Algorithm 3 takes $\mathcal{O}(n^4)$ time.

Acknowledgements

I would like to thank Professor David Speyer for his crucial help and guiding remarks. I would like to thank Dr. Steven Damelin for catalyzing this REU project. I would like to thank Neophytos Charalambides and Cyrus Anderson for various insightful discussions, as well as for several poignant and cherished conversations. I would like to extend my thanks Sean Kelly for his help and invaluable efforts. Lastly, I would also like to thank the University of Michigan Math Department and the NSF for making this REU possible.

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