# An Introduction to Multidimensional Scaling and ISOMAP

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- In this case the distance *d* above is the usual Euclidean distance.
- There are cases where the matrix D is valid distance matrix, but still there exists no set of vectors  $x_1, ..., x_N$  in any  $R^d$  with perfect  $||x_i x_j|| = d_{ij}$ . Such a distance is called non-Euclidean distance.

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4. A d-dimensional MDS coordinates of n objects is derived from the coordinate matrix  $X = E_d \Lambda_d^{\frac{1}{2}}$ , where  $E_d$  is the matrix of d eigenvectors and  $\Lambda_d$  is the diagonal matrix of d eigenvalues of B, respectively



#### Take the square the elements of D



#### Construct the J matrix



#### Construct double centering matrix



Solve the largest 2 eigenvalues and eigenvector of B

$$\lambda_1 = 9724.168, \ \lambda_2 = 3160.986, \quad \mathbf{e_1} = \begin{pmatrix} -0.637\\ 0.187\\ -0.253\\ 0.704 \end{pmatrix}, \ \mathbf{e_2} = \begin{pmatrix} -0.586\\ 0.214\\ 0.706\\ -0.334 \end{pmatrix}$$

Use that to construct the final MDS coordinates.

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Use that to construct the final MDS coordinates.



## **Stress Majorization**

In the classical MDS algorithm the cost function that we are trying to optimize is called the stress function and it is given by :

$$Stress_D(x_1,x_2,\ldots,x_N) = \left(\sum_{i 
eq j=1,...,N} \left(d_{ij} - \|x_i - x_j\|
ight)^2
ight)^{1/2}$$

In this function we try to find  $x_1, ..., x_N$  in a certain dimension d such that Stress $(x_1, ..., x_N)$  is as small as possible

## **Stress Majorization**

The stress function has a more general form as :

$$\sigma(X) = \sum_{i < j \leq n} w_{ij} (d_{ij}(X) - \delta_{ij})^2$$

Here wij weight between a pair of points (i,j) that represents the confidence in in the similarity between points (i,j).

 $\delta_{ij}$  the given distance between the points i,j

## **Stress Majorization**

"Pressing" the data into 2 dimensions enables us to visualize the data. However, that comes with a price : high stress function value (which correlates wit distorted representation )



MDS is implemented in Sklearn



Example

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Example

MDS is implemented in Sklearn

A selection from the 64-dimensional digits dataset







Example

Application



### Measuring distance between two persistence diagrams



### Bottleneck distance between two persistent diagrams

• Given two persistence diagrams X and Y, let η be a bijection between points in the diagram. The following two distances are commonly used in the context of PH to measure the distance between two persistence diagrams:

$$W_{\infty}(X,Y) = \inf_{\eta: X \to Y} \sup_{x \in X} \|x - \eta(x)\|_{\infty}$$

$$W_q(X,Y) = \left[\inf_{\eta:X\to Y} \sum_{x\in X} \|x-\eta(x)\|_{\infty}^q\right]^{1/q}$$

### Bottleneck distance between two persistent diagrams



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- The orientation of the result MDS "picture" is arbitrary.
- When we obtain MDS coordinates that have non-zero stress, we should remember that the distances among the resulting items are distorted representations of the relationships given by the input data. This distortion is greater when the stress is greater.
- That being said, we, in general, can rely on the larger distances as being more accurate than smaller distances.

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The main idea in non-metric MDS :

- The actual values given to us are not that meaningful
- *Ranking among different points* is important
- Non-metric MDS finds a low-dimensional representation, which respects the ranking of distances as much as possible

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- Relaxing  $dij \approx d' ij$  from MDS by allowing  $d' ij \approx f(d_{ij})$ , for some monotone function f

Monotonic means :  $d_{ij} < d_{kl} \Leftrightarrow f(dij) \leq f(d_{kl})$ 

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Given a dimension d, non-metric MDS seeks to find an optimal configuration  $X \subset \mathbb{R}^d$  that gives  $f(d_{ij}) \approx d^{\hat{}}ij = ||x_i - x_j||$  as close as possible. •  $f(d_{ij}) = d^*ij$  is only required to preserve the order of  $d_{ij}$ ,

i.e.,  $d_{ij} < d_{kl} \Leftrightarrow f(d_{ij}) \leq f(d_{kl}) \Leftrightarrow d_{ij}^* \leq d_{kl}^*$ 

The stress function for non-metric MDS is given by :

$$Stress = \left(\sum_{i < j} (\hat{d}_{ij} - f(d_{ij}))^2 / \sum d_{ij}^2 
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So the steps for ISOMAP on a given data :

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So the steps for ISOMAP on a given data :

1- Construct the neighborhood graph of the data X using one of the neighborhood graphs we studied earlier in the course

2- Use the Dijekstra algorithm or the Floyd–Warshall algorithm to find the distance between nodes on the graph

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3- Apply MDS on the distance matrix above and extract the coordinates with the desired dimension



A- Euclidian distance might not represent the actual distance between the points in the data.

B- We can construct the neighborhood graph of the data and then compute the geodesic distance between the points of the graph

C- Embedding the space we obtained in B into the plane.

let dist be a  $|V| \times |V|$  array of minimum distances initialized to infinity

```
for each edge (u, v)
dist[u][v] \leftarrow w(u, v) // the weight of the edge (u,v)
for each vertex v
dist[v][v] \leftarrow 0
for k from 1 to |V|
for i from 1 to |V|
if dist[i][j] > dist[i][k] + dist[k][j]
dist[i][j] \leftarrow dist[i][k] + dist[k][j]
end if
```

Floyd algorithm is good to use when we want to compute the distance matrix on a dense graph. When the graph G is sparse, Dijekstra algorithm is a better choice.