Geometric MDS Performance for Large Data Dimensionality Reduction and Visualization

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Abstract. Multidimensional scaling (MDS) is a widely used technique for mapping data from a high-dimensional to a lower-dimensional space and for visualizing data. Recently, a new method, known as Geometric MDS, has been developed to minimize the MDS stress function by an iterative procedure, where coordinates of a particular point of the projected space are moved to the new position defined analytically. Such a change in position is easily interpreted geometrically. Moreover, the coordinates of points of the projected space may be recalculated simultaneously, i.e. in parallel, independently of each other. This paper has several objectives. Two implementations of Geometric MDS are suggested and analysed experimentally. The parallel implementation of Geometric MDS is developed for multithreaded multi-core processors. The sequential implementation is optimized for computational speed, enabling it to solve large data problems. It is compared with the SMACOF version of MDS. Python codes for both Geometric MDS and SMACOF are presented to highlight the differences between the two implementations. The comparison was carried out on several aspects: the comparative performance of Geometric MDS and SMACOF depending on the projection dimension, data size and computation time. Geometric MDS usually finds lower stress when the dimensionality of the projected space is smaller.

Key words: dimensionality reduction, multidimensional scaling, Geometric MDS, large-scale data, multi-core implementation, SMACOF, Python codes.

1. Introduction

Every day we receive an enormous amount of data, information, and knowledge. Data becomes information when it is contextualised and related to a specific problem or solution. Data mining is an important part of knowledge discovery processes in various fields and sectors, such as medicine, economics, finance, telecommunications. Data mining helps uncover hidden information from vast amounts of data, which is valuable for recognising important facts, relationships, trends, and patterns. As data sets become increasingly large, more efficient ways of visualizing, analysing and interpreting the information they contain are needed. Comprehension of data is challenging, especially when

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the data relates to a complex object, a phenomenon that is described by many parameters, attributes, or features. Such data are called multidimensional, and the main goal is to make some visual insight into the data set being analysed. Visual information can be perceived much faster than textual information. For human perception, the multidimensional data must be represented in a low-dimensional space, usually two or three dimensions.

The main task in dimensionality reduction is to represent objects with a smaller set of more "compressed" features. Another reason for reducing the dimensionality or visualization of the data is to reduce the computational load for further data processing. Dimensionality reduction techniques enable the extraction of meaningful information hidden in the data. In addition, one of the most important purposes of data visualization is to get an idea of how close or far away certain points of the analysed multidimensional data are from each other.

Consider the multidimensional data set as an array $X = \{X_i = (x_{i1}, \ldots, x_{in}), i = 1, \ldots, m\}$ of *n*-dimensional data points $X_i \in \mathbb{R}^n$, $n \ge 3$. A data point X_i is the result of an observation of some object or phenomenon that depends on *n* features. Dimensionality reduction means finding a set of coordinates (locations) of points $Y_i = (y_{i1}, \ldots, y_{id})$, $i = 1, \ldots, m$, in a lower-dimensional space (d < n), where the particular point $X_i \in \mathbb{R}^n$ is represented by $Y_i \in \mathbb{R}^d$. If $d \le 3$, then dimensionality reduction results can be presented visually for more convenient human decision-making.

Dimensionality reduction or data visualization techniques play an important role in machine learning (Murphy, 2022; Zhou, 2021; Ray et al., 2021; Dzemyda et al., 2013; Dos Santos and Brodlie, 2004; Buja et al., 2008). Such visualization is useful, especially in exploratory analysis: they provide insights into similarity relationships in highdimensional data that would be unlikely to be obtained without visualization (Lee and Verleysen, 2007; Van Der Maaten et al., 2009; Markeviciute et al., 2022; Xu et al., 2019; Bernatavičienė et al., 2007; Kurasova and Molyte, 2011; Borg and Groenen, 2005; Dzemyda and Kurasova, 2006; Dzemyda et al., 2013; Jolliffe, 2002; Karbauskaitė and Dzemyda, 2015, 2016; Groenen et al., 1995). The classical dimensionality reduction methods for data visualization include linear principal component analysis (PCA) (Jolliffe, 2002; Jackson, 1991) and multidimensional scaling (MDS) (Torgerson, 1958; Borg and Groenen, 2005; Borg et al., 2018). PCA seeks to reduce the dimensionality of the data by finding orthogonal linear combinations (principal components) of the original variables with the highest variance (Jackson, 1991; Medvedev et al., 2011). The interpretation of principal components can sometimes be complex. PCA cannot cover non-linear structures consisting of arbitrarily shaped clusters or manifolds because it describes the data in terms of a linear subspace. Since global methods such as PCA and MDS cannot represent the local non-linear structure, there have been developed methods that preserve the raw local distances, i.e. the values of distances between nearest neighbours. Isomap, Local Linear Embedding (LLE), Hessian Local Linear Embedding and Laplacian Eigenmaps are traditional methods that attempt to preserve local Euclidean distances from the original space (Wang et al., 2021; Espadoto et al., 2021). More recent methods that focus on local structure preservation are t-Distributed Stochastic Neighbour Embedding (t-SNE) (Van der Maaten and Hinton, 2008) and UMAP (McInnes et al., 2018). A comprehensive

review of dimensionality reduction methods is presented in Murphy (2022), Espadoto *et al.* (2021), Wang *et al.* (2021), Vachharajani and Pandya (2022), Dzemyda *et al.* (2013), Lee and Verleysen (2007), Van Der Maaten *et al.* (2009), Xu *et al.* (2019).

Artificial neural networks can also be applied to reduce dimensionality and visualize data (Dzemyda *et al.*, 2007). A feedforward neural network is used for a topographic, structure-preserving, dimension-reducing transformation of data, with the additional ability to incorporate various degrees of related subjective information. There is also a neural network architecture developed specifically for topographic mapping, known as a Self-Organizing Map (SOM) (Kohonen, 2001; Stefanovic and Kurasova, 2011), which uses implicit lateral connections in the output layer of neurons. SOM is a technique used for both clustering and data dimensionality reduction (Dzemyda *et al.*, 2007, 2013). In addition, there is a special learning rule, similar to backpropagation, which allows an ordinary feedforward artificial neural network to learn the Sammon mapping, which is a special case of metric MDS, in an unsupervised way. The learning rule for this type of neural network is known as SAMANN (Mao and Jain, 1995; Ivanikovas *et al.*, 2007; Medvedev *et al.*, 2011; Dzemyda *et al.*, 2007).

An alternative view of dimensionality reduction is offered by multidimensional scaling (Borg and Groenen, 2005). Multidimensional scaling (MDS) is a classical non-linear approach that maps an original high-dimensional data set onto a lower-dimensional data set, but does so in an attempt to preserve the proximities between the corresponding data points. It is one of the most popular methods for multidimensional data visualization (Murphy, 2022; Borg *et al.*, 2018; Dzemyda *et al.*, 2013). Despite the fact that MDS demonstrates great versatility, it is computationally demanding. This can be challenging when the data amount increases. Traditional MDS approaches are limited when analysing very large data sets, as they require long computational time and large amounts of memory. Until now, there have been various studies aimed at creating a new solution or modifying the MDS to analyse large amounts of data and speed up the visualization process (Orts *et al.*, 2019; Qiu and Bae, 2012; Pawliczek *et al.*, 2014; Ingram *et al.*, 2008; Medvedev *et al.*, 2011; Ivanikovas *et al.*, 2007).

The input data for MDS is a symmetric $m \times m$ matrix $\mathbf{D} = \{d_{ij}, i, j = 1, ..., m\}$ of proximities (similarities or dissimilarities) between the pairs of objects. More often, the dissimilarities are used to represent the proximities, however, the formulation of the multidimensional scaling problem and its solving methods remain the same in both cases. A lower dissimilarity value means that the objects are more similar. As dissimilarity, d_{ij} can be a distance between points X_i and X_j , i, j = 1, ..., m (Dzemyda *et al.*, 2013). The smaller the distance, the closer the points are.

One of the most popular algorithms for MDS is SMACOF (De Leeuw, 1977; De Leeuw and Mair, 2009). The algorithm is based on the majorization approach (Groenen *et al.*, 1995), which iteratively replaces the original objective function with an auxiliary majorization function that is much easier to optimize. Majorization is not just an algorithm, it is more of a prescription for constructing optimization algorithms. The principle of majorization consists in constructing a auxiliary function that majorizes a certain objective function (De Leeuw and Mair, 2009). Experimental studies have shown that SMACOF is

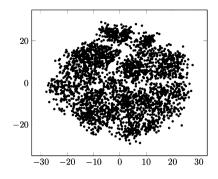
the most accurate algorithm compared to others (Orts *et al.*, 2019; Ingram *et al.*, 2008). There are a number of popular software implementations of SMACOF (De Leeuw and Mair, 2009; Pedregosa *et al.*, 2011; Orts *et al.*, 2019; MATLAB, 2012).

An alternative to SMACOF the Geometric MDS proposed in Dzemyda and Sabaliauskas (2020) and Dzemyda and Sabaliauskas (2021a), and developed in Sabaliauskas and Dzemyda (2021), Dzemyda and Sabaliauskas (2021b, 2021c), Dzemyda et al. (2022), Dzemyda and Sabaliauskas (2022). This will be discussed in more detail further in the paper. Figure 1 illustrates the visualization using Geometric MDS of 10, 50, and 100dimensional datasets generated with Gaussian and Ellipsoidal cluster generators (Handl and Knowles, 2005). The dimensionality of the multidimensional data has been reduced to 2 (d = 2). The Gaussian cluster generator is based on a standard cluster model using multivariate normal distributions, and the ellipsoidal cluster generator creates ellipsoidal clusters with the major axis at an arbitrary orientation. For each cluster, data points are generated at a Gaussian distributed distance from a uniformly random point on the major axis, in a uniformly random direction, and are rejected if they lie outside the boundary (Handl and Knowles, 2005). As the datasets are generated using cluster generators for large high-dimensional data sets with large numbers of clusters, some noisy data may be included. The example in Fig. 1 illustrates the use of visualization techniques to identify outliers, patterns in data in the form of clusters, relationships, and trends. The figure clearly shows clusters of analysed data and the location of individual multidimensional data among the rest.

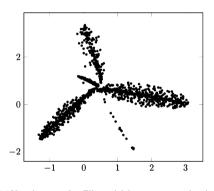
The novelty of this paper consists in the experimental investigation of parallelization of recently developed Geometric Multidimensional Scaling. The process of parallelization can lead to faster dimensionality reduction and data visualization process and to the possibility to process a large-scale multidimensional data. Theoretically proved properties of Geometric MDS are applied in parallelization. In addition, Geometric MDS is also experimentally compared with SMACOF, which is one of the most popular implementations of MDS. It was found that Geometric MDS is superior to the SMACOF in most cases.

2. Geometric MDS: Multidimensional Scaling From a Geometric Point of View

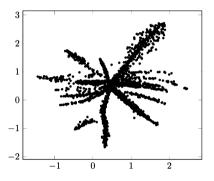
MDS looks for coordinates of points Y_i representing X_i in the lower-dimensional Euclidean space \mathbb{R}^d by minimizing the stress function (Dzemyda *et al.*, 2013). There are several MDS implementations with different stress functions (see review in Dzemyda *et al.*, 2013). However, their minimization is rather complicated. In Dzemyda and Sabaliauskas (2020), Sabaliauskas and Dzemyda (2021), Dzemyda and Sabaliauskas (2021a, 2021c) a new approach, called Geometric MDS, with low computational complexity is proposed, making metric MDS applicable to large-scale data. This method is valuable and advantageous because the stress function and the multidimensional scaling have been considered from a geometrical point of view. The new interpretation of the stress function allows us to find the proper step size and direction of descent forward to the minimum of the stress function analytically, if we consider and move a single point of the projected space. A special property of the new approach is that there is no need for an analytical expression of



(a) Visualization of a data set generated with the Gaussian cluster generator, $m=1248,\,n=10,\,d=2,\,20$ clusters



(b) Visualization of a data set generated with the Gaussian cluster generator, $m=2707,\,n=10,\,d=2,\,40$ clusters



(c) Visualization of an Ellipsoidal data set generated with the Gaussian cluster generator, m = 1506, n = 100, d = 2, 4 clusters

(d) Visualization of an Ellipsoidal data set generated with the Gaussian cluster generator, m=3140, n=50, d=2, 10 clusters

Fig. 1. Examples of dimensionality reduction results obtained using Geometric MDS.

the stress function. Even more, no linear search, which is used for local descent in optimization, is needed. Theoretical investigation showed that the step direction determined by Geometric MDS coincides with the direction of the steepest descent method (Dzemyda and Sabaliauskas, 2021a). The step size found analytically is such that it guarantees stress reduction in that direction.

A Geometric MDS has the advantage that it can use the simplest stress function, and there is no need to normalize it based on the number of data points and the scale of proximities. In Dzemyda and Sabaliauskas (2021a) it is shown that Geometric MDS does not depend on the scale of proximities and can therefore use a simple stress function, such as a raw stress function:

$$S(Y_1, \dots, Y_m) = \sum_{i=1}^m \sum_{j=i+1}^m (d_{ij} - d_{ij}^*)^2,$$
(1)

where d_{ij}^* is the Euclidean distance between points Y_i and Y_j in a lower-dimensional space:

$$d_{ij}^* = \sqrt{\sum_{l=1}^{d} (y_{il} - y_{jl})^2}.$$
 (2)

The optimization problem is to find the minimum of the function $S(\cdot)$ (1), and the optimal coordinates of the points $Y_i = (y_{i1}, \ldots, y_{id}), i = 1, \ldots, m$:

$$\min_{Y_1,\ldots,Y_m \in \mathbb{R}^d} S(Y_1,\ldots,Y_m).$$
(3)

Let there be some initial configuration of points Y_1, \ldots, Y_m . The simplest way to minimize the stress $S(\cdot)$ by Geometric MDS is a consecutive changing of positions of separate points Y_1, \ldots, Y_m many times (iterations). This realization is denoted by GMDS1. One iteration of GMDS1 changes all points $Y_i = (y_{i1}, \ldots, y_{id})$ in consecutive order when *i* runs from 1 to *m* once. Here we compute a new position $Y_i^* = (y_{i1}^*, \ldots, y_{id}^*)$ of $Y_i = (y_{i1}, \ldots, y_{id})$ when the positions of remaining points $Y_1, \ldots, Y_{j-1}, Y_{j+1}, \ldots, Y_m$ are fixed, and update Y_j by Y_j^* then. So, Geometric MDS recalculates the coordinates of a single *d*-dimensional point Y_j . The stopping condition may be the number of iterations or the value of decrease of the stress function (1). Convergence of this algorithm to the local minimum of the stress is proved.

The core formula of Geometric MDS, when defining the transition from Y_j to the new position Y_i^* , is as follows:

$$Y_j^* = \frac{1}{m-1} \sum_{\substack{i=1\\i \neq j}}^m A_{ij},$$
(4)

where the point A_{ij} lies on the line between Y_i and Y_j , $i \neq j$, at a distance d_{ij} from Y_i (see Dzemyda and Sabaliauskas, 2020; Dzemyda and Sabaliauskas, 2021a; Sabaliauskas and Dzemyda, 2021):

$$A_{ij} = Y_i + (Y_j - Y_i) \frac{d_{ij}}{d_{ij}^*}.$$
(5)

Let's consider a generalized way to update the entire set of points $Y = \{Y_1, \ldots, Y_m\}$ to $Y^* = \{Y_1^*, \ldots, Y_m^*\}$. The main equation of Geometric MDS (4) is used for defining the transition from Y_j to the new position Y_j^* . The raw stress function (1) decreases after Y_j goes to Y_j^* , that is, moving any of the projected points by the Geometric MDS method reduces the stress (Dzemyda and Sabaliauskas, 2021a). But in the generalized scenario, all points Y_1, \ldots, Y_m change their coordinates to Y_1^*, \ldots, Y_m^* simultaneously and independently of each other during a single iteration of stress minimization. Therefore, we perform a simultaneous repositioning of all points Y_1, \ldots, Y_m of the projected

space in the directions determined by the Geometric MDS strategy for individual points. We update Y_1, \ldots, Y_m by Y_1^*, \ldots, Y_m^* after all the points Y_1^*, \ldots, Y_m^* are computed, only. This realization is denoted by GMDS*m*. One iteration of GMDS*m* changes all points $Y_i = (y_{i1}, \ldots, y_{id})$ simultaneously once. The convergence is proved theoretically in Dzemyda and Sabaliauskas (2022).

3. Multi-Core Implementation of Geometric MDS

The GMDS*m* version of the Geometric MDS updates all points Y_1, \ldots, Y_m in a lower dimension simultaneously and independently of each other. This enables the use of parallel computing for updating each Y_i independently, i.e. in parallel (see Algorithm 1). Such possibility of parallelization has been investigated further in this paper utilising multicore processing. The computational resources of a single personal computer were used in this work. Experiments using different number of CPU threads were performed to obtain preliminary estimates of the efficiency of multi-core implementation of GMDS*m*.

The purpose of the experiment is to compare the time required to compute new points Y_1^*, \ldots, Y_m^* in the projected space from Y_1, \ldots, Y_m , when using a sequential programming implementation of GMDSm with the computing power of one CPU core (single thread), and respectively when using a parallel algorithm using 2, 4, 6, 8, 10, 12 CPU threads. Here each point Y_i^* is computed using Eq. (4), and all points Y_1, \ldots, Y_m change their coordinates to Y_1^*, \ldots, Y_m^* at once simultaneously and independently of each other during a single iteration. One iteration is a recalculation (optimal updating) of all points $Y_i = (y_{i1}, \ldots, y_{id}), i = 1, \ldots, m$ once. The pseudo-code of the algorithm used for experiments is provided in Algorithm 1 (Dzemyda et al., 2022). Note that only the time taken to compute the new coordinates was measured during the experiments (see the part of Algorithm 1 that is marked as running in parallel). The California Housing dataset (Pace and Barry, 1997) was used for the experiments. The dataset consists of data collected in the 1990 California Census, it has 20640 examples of block groups, each containing an average of 1425.5 people living in a geographically compact area. There are 10 different features per example: 8 numeric attributes, one predictive attribute and the target. For preliminary experiments and due to limited computing resources, only part of the data was used (see Table 1), i.e. data of 500, 1000, 1500, 2000, 2500, 3000, 3500, 4000, 4500 and 5000 observations were analysed, and the dimensionality of the multidimensional data was reduced to 2. This increase in data size reveals a trend which demonstrates that with further increases in data size, the use of parallel computing becomes more efficient. The experiments resulted in an average dependence of the time to calculate new point coordinates (without taking into account the full visualization process) on the size of the data when using different numbers of CPU threads for parallel processing. Each experiment was repeated 10 times to obtain more reliable data, so Table 1 presents the average value of the generalized data. For the experiments we used an AMD Ryzen 5 2600 processor, 4000 MHz (6 cores, 12 threads) with high multiprocessor performance, with 16GB of DDR4-2666 1333 MHz RAM, and the Python-Anaconda (Anaconda, 2022) environment was used as the software.

Algorithm 1: A parallel Geometric MDS algorithm for simultaneous calculation of new points Y_i^* in the projected space

Data: *n* is the dimensionality of data, *m* is the number of data points, *d* is the dimensionality of the projected space;

 $D = (d_{ij})$ is a symmetric $m \times m$ distance matrix containing the distances, taken pairwise, between *n*-dimensional data points;

iterations is the number of transitions from Y_1, \ldots, Y_m to the new position Y_1^*, \ldots, Y_m^* . Y_1, \ldots, Y_m is a set of coordinates of points in a lower-dimensional space (d < n), where the particular point $X_i \in \mathbb{R}^n$, $i = 1, \ldots, m$ is represented by $Y_i \in \mathbb{R}^d$. One iteration is a change of all points $Y_i = (y_{i1}, \ldots, y_{id})$ once; j^0 is an index of the point among Y_1, \ldots, Y_m , whose coordinates will be recalculated during one iteration.

Result: Y^* is $m \times d$ matrix of the projected points; S_1 – value of stress function $Y \leftarrow$ initial values of $m \times d$ matrix;

for *it* \leftarrow 1 *to iterations* **do**

for each $j^0 \in \{1, ..., m\}$ in parallel do for each $i \in \{1, ..., m\} \setminus \{j^0\}$ do $w \leftarrow d_{ij^0} / \sqrt{\sum_{k=1}^d (y_{ik} - y_{j^0k})^2};$ for $k \leftarrow 1$ to d do $A_{ik} \leftarrow y_{ik} + (y_{j^0k} - y_{ik})w;$ end for $k \leftarrow 1$ to d do $y_{j^0k}^* \leftarrow \frac{1}{m-1} \sum_{j=1, j \neq j^0}^m A_{jk};$ end $Y_{j^0} \leftarrow Y_{j^0}^*;$ /* $Y_{j^0}^*$ are final coordinates of the optimized point */ end

end

for $i \leftarrow 1$ to m do $d_{ii} \leftarrow 0;$ for $j \leftarrow i + 1$ to m do $\begin{pmatrix} d_{ij}^* \leftarrow \sqrt{\sum_{k=1}^d (y_{ik}^* - y_{jk}^*)^2}; \\ d_{ji}^* \leftarrow d_{ij}^*; \\ /^* d_{ij}^* \text{ is a symmetric } m \ge m \text{ distance matrix containing the distances,} \\ \text{ taken pairwise, between } d\text{-dimensional data points*/} \\ end$ end

 $S_1 \leftarrow S(Y_1^*, \dots, Y_m^*)$. /* MDS stress $S(\cdot)$ defined by (1) */

Average dependence of the computation time of new coordinates (not considering the complete visualization process) on the data size when using different numbers of CPU threads for parallel processing, n = 8; the dimensionality of multidimensional data was reduced to 2 (d = 2).

Number of points, <i>m</i>	Sequential	Number of CPU threads (parallel processing)								
	algorithm	2	4	6	8	10	12			
100	0.0445	0.6492	0.7018	0.8365	1.0246	1.2450	1.3315			
500	1.1770	1.1783	0.9902	1.0372	1.2384	1.4430	1.5087			
1000	4.8455	2.9246	1.9148	1.7087	1.9091	2.0435	2.0459			
1500	10.8324	5.8416	3.4209	2.8879	2.9647	3.0228	3.0353			
2000	19.1812	9.8804	5.5490	4.3496	4.3921	4.4029	4.3415			
2500	30.3274	15.3864	8.2581	6.2743	6.3550	6.2527	6.2111			
3000	43.6203	21.5616	11.6450	8.7684	8.7479	8.6064	8.2878			
3500	59.4401	28.9800	15.8067	11.5609	11.2593	11.0542	10.6939			
4000	79.0386	37.7887	20.7712	15.1054	14.3880	14.3113	13.4704			
4500	99.2351	47.8711	25.8535	18.8063	18.0433	17.4048	16.8381			
5000	124.6468	59.0778	31.7166	23.1730	22.3510	21.3549	20.6990			

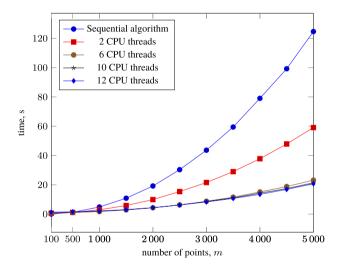


Fig. 2. Average dependence of the computation time of new coordinates (not taking into account the complete visualization process) on the size of the data, when using different numbers of CPU threads for parallel processing, n = 8.

The results obtained (see Table 1 and Fig. 2) demonstrate that even using two CPU threads for parallel processing, it is possible to achieve significant process speedup (on average up to 2 times) compared to sequential computing. The results obtained conclude that when using more than two processor threads for parallel computing, it is possible to increase the speed of the process of calculating new coordinates in the projected space on average up to 6 times. It is also worth noting that the difference in time increases with increasing the size of the analysed data, that is, the larger the size of the data used for visualization, the more efficient the process of parallel computing (see Fig. 2). It should be

noted that these data were obtained as a result of initial experiments, using the computational power of a personal computer. Consider also that the CPU used for the experiments had 6 cores and 12 threads, so, as can be seen from the results, the efficiency of calculations using more than 6 threads is not high. The results are promising, which indicates the feasibility of using Geometric MDS to visualize large-scale multidimensional data using the computing power of a supercomputer or cluster for parallel computing.

4. Sequential Implementation of Geometric MDS

Although MDS demonstrates great versatility, it is computationally expensive and not scalable, and requires handling the entire data distance or other proximity matrix. This can be challenging when we deal with large-scale data. In this section, we propose the optimized Python implementation of GMDSm for sequential computations.

MDS-type algorithms require to update a matrix of $m \times m$ of all pairwise Euclidean distances d_{ij}^* between points Y_i and Y_j , i.e. for m points a $m \times m$ matrix is required, each entry of which defines a non-negative distance between the pair of points. Calculations of Euclidean distance occur frequently in machine learning and computer vision (Albanie, 2019; Song *et al.*, 2016). The element of the distance matrix is defined by (2).

However, as noted in Albanie (2019), the Euclidean distance matrix can be computed in another way, which is equivalent to (2):

$$\left(d_{ij}^* \right)^2 = (Y_i - Y_j)(Y_i - Y_j)^T = \|Y_i\|^2 - 2Y_i Y_j^T + \|Y_j\|^2,$$
(6)

where the first and last terms are the norms of vectors and can be calculated separately.

This method defined by (6) of calculating the Euclidean distance matrix has been implemented and used in Geometric MDS Python programming implementation. Applying (6) allows to speed up the visualization significantly because of much simpler calculations, and also allows to extend the dimensionality reduction for relatively large-scale data. Python function code for calculating distance matrix from a given $m \times d$ matrix $Y = (Y_i = (y_{i1}, \ldots, y_{id}), i = 1, \ldots, m)$ is presented in Listing 1. This code can be used to compute a $m \times m$ distance matrix from $m \times n$ matrix $X = (X_i = (x_{i1}, \ldots, x_{in})), i = 1, \ldots, m$, too.

The algorithm in Listing 1 returns 10^{-5} values on the diagonal of the proximity matrix and in other situations when points among X_i and X_j coincide (the coincidence is only possible before Geometric MDS starts). This avoids dividing by zero for further calculations.

The Geometric MDS and SMACOF programming implementation code for Python has been optimized for computational speed. The Geometric MDS stress minimization method as well as the SMACOF implementation are featured by the fact that all points in low-dimensional space change their coordinates simultaneously and independently of each other during one iteration of stress minimization. We provide a function code implemented to calculate new coordinates of points in *d*-dimensional space in one iteration using Geometric MDS and SMACOF (see Listings 2 and 3), as well as the code of a fully

```
i import numpy as np
def euclidean_distances(Y):
Q = np.einsum("ij,ij->i", Y, Y)[:, np.newaxis]
distances = -2 * Y @ Y.T
distances += Q
distances += Q.T
np.maximum(distances, 1e-10, out = distances)
return np.sqrt(distances)
```

Listing 1. Python function code for calculating distance matrix.

```
1 #dm is a symmetric m x m matrix of proximities between n-dimensional
data points;
2 #Y is m x d matrix of the projected points (initial configuration of
points);
3 #Y_new is matrix of size m x d of new positions of projected points.
4 def correction(dm, dist, Y):
5 B = 1 - np.divide(dm, dist)
6 B[np.arange(len(B)), np.arange(len(B))] -= B.sum(axis = 1)
7 Y_new = Y + 1 / (len(Y) - 1) * np.dot(B, Y)
8 return Y_new
```

Listing 2. Python function code for calculating new points $Y^* = \{Y_1^*, \dots, Y_m^*\}$ in a low-dimensional space using Geometric MDS.

```
1 #dm is a symmetric m x m matrix of proximities between n-dimensional
data points;
2 #Y is m x d matrix of the projected points (initial configuration of
points);
3 #Y_new is matrix of size m x d of new positions of projected points.
4 def correction2(dm, dist, Y):
5 B = -np.divide(dm, dist)
6 B[np.arange(len(B)), np.arange(len(B))] -= B.sum(axis = 1)
7 Y_new = 1 / len(Y) * np.dot(B, Y)
8 return Y_new
```

Listing 3. Python function code for calculating new points $\hat{Y}^* = {\{\hat{Y}_1^*, \dots, \hat{Y}_m^*\}}$ in a low-dimensional space using SMACOF (optimized version of SMACOF, referred to as SMACOF-Fast).

working Python program that implements the sequential Geometric MDS algorithm (see Listing 4), where the stopping condition for stress (1) minimization is either the iteration number or the accuracy. In one iteration (see Listings 2 and 3), the coordinates of all points in the projected space are recalculated once. Listing 3 presents the function code for calculating new coordinates using SMACOF, adapted from the scikit-learn library (Pedregosa *et al.*, 2011) and updated to improve computation speed (this implementation is denoted as SMACOF-Fast). In this implementation, the stress majorization, which is also known

```
1 #dm is a symmetric m x m matrix of proximities between n-dimensional
      data points;
2 #Y is m x d matrix of the projecting points (initial configuration of
      points);
3 #eps refers to the accuracy of the final stress value;
4 #max it is a maximum number of iterations;
5 #dist is a distance m x m matrix of points Y;
6 def GMDSm(dm, Y, eps, max it):
7
    S old, it = np.inf, 0
  np.fill diagonal(dm, 1e-5)
8
   while it < max it:
9
      dist = euclidean distances(Y)
10
      S = np.sum((dm - dist) ** 2) / 2
      if abs(S old - S) < eps:
13
       break
14
      Y = correction(dm, dist, Y)
      #use Y = correction2(dm, dist, Y) for SMACOF-Fast
15
      S \text{ old} = S
16
      it += 1
18
  return Y
```

Listing 4. Main program of Geometric MDS.

as the Guttman transform, guarantees monotonic stress convergence and is more powerful than traditional methods such as gradient descent.

Geometric MDS becomes the main competitor of SMACOF. We would like to highlight the main differences between the two algorithms. More detailed theoretical comparison of them is provided in Dzemyda and Sabaliauskas (2022).

At first, we define a Geometric MDS step from point Y_j to Y_j^* , j = 1, ..., m that calculates new coordinates of point Y_i^* :

$$Y_j^* = Y_j + \frac{1}{m-1} \sum_{i=1}^m (Y_i - Y_j) f_{ij}, \quad f_{ij} = 1 - \frac{d_{ij}}{d_{ij}^*}, \quad f_{ii} = 0.$$
(7)

SMACOF is an iterative procedure that calculates the Guttman transform (Guttman, 1968) at each iteration and updates simultaneously the coordinates of *m* points $Y_j = (y_{j1}, \ldots, y_{jd}), j = 1, \ldots, m$:

$$\hat{Y}^* = \frac{1}{m} B(Y)Y,\tag{8}$$

where $\hat{Y}_{j}^{*} = (\hat{y}_{j1}^{*}, \dots, \hat{y}_{jd}^{*}), \ j = 1, \dots, m$ are new points, *Y* and \hat{Y}^{*} are $m \times d$ matrices of coordinates of points Y_{j} and \hat{Y}_{j}^{*} , respectively, B = B(Y) is $m \times m$ matrix with elements

$$B_{ij} = -\frac{d_{ij}}{d_{ij}^*}, \qquad B_{ii} = -\sum_{\substack{j=1\\j\neq i}}^m B_{ij}, \quad i, j = 1, \dots, m.$$
(9)

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It is easy to check that SMACOF works very similarly to Geometric MDS. By calculating coordinates of point \hat{Y}_i^* , j = 1, ..., m, (8) and (9) can be expressed as

$$\hat{Y}_{j}^{*} = \frac{1}{m} \sum_{i=1}^{m} \frac{(Y_{j} - Y_{i})d_{ij}}{d_{ij}^{*}}.$$
(10)

It is proved in Dzemyda and Sabaliauskas (2022) that both Geometric MDS and SMACOF make a step from the point Y_j to its new position Y_j^* or \hat{Y}_j^* , respectively in the same direction with different step size. The step size by Geometric MDS is longer by $\frac{m}{m-1}$ times.

Geometric MDS can be expressed in matrix form as well:

$$Y^* = Y + \frac{1}{m-1}\hat{B}(Y)Y,$$
(11)

where $Y_j^* = (y_{j1}^*, \dots, y_{jd}^*)$, $j = 1, \dots, m$ are new points, Y and Y^* are $m \times d$ matrices of coordinates of points Y_j and Y_i^* , respectively, $\hat{B} = \hat{B}(Y)$ is $m \times m$ matrix with elements

$$\hat{B}_{ij} = 1 - \frac{d_{ij}}{d_{ij}^*}, \qquad \hat{B}_{ii} = -\sum_{\substack{j=1\\j\neq i}}^m \hat{B}_{ij}, \quad i, j = 1, \dots, m.$$
(12)

4.1. Computation Time Cost by Geometric MDS and SMACOF

The dimensionality reduction performance was compared using SMACOF (from the scikit-learn library, Pedregosa *et al.*, 2011), SMACOF-Fast (Listing 3) and Geometric MDS algorithms (Listing 2). Experiments were carried out using:

- randomly generated sets of 10-dimensional data points $X_i = (x_{i1}, \ldots, x_{id}), i = 1, \ldots, m$, with coordinates uniformly distributed in the interval (0, 1);
- random proximity matrices with elements uniformly distributed in the interval (0, 1).

The dimensionality was reduced to d = 2. After one iteration, the time taken to calculate the new coordinates of the point positions in the projected space was measured and the stress value $S(\cdot)$ defined by (1) was obtained. Data with different number of points *m* were used: 10, 20, 50, 100, 200, 500, 1000, 2000, 5000, 10000, 15000, and 20000. In order to obtain more accurate and reliable results, each experiment was repeated 100 times. The results of these experiments are shown in Tables 2 and 3. Since the number of computational operations using Geometric MDS and SMACOF-Fast is approximately the same (it can be concluded from Listing 2 and Listing 3), the time required to compute new coordinates using Geometric MDS and SMACOF-Fast is about the same.

The dependency of the calculation time on a number of points using Geometric MDS (see Listing 2) and SMACOF (Pedregosa *et al.*, 2011) is shown in Fig. 3. The number *m* of points (m = 10, 20, 50, 100, 200, 500, 1000, 2000, 5000, 10000, 15000, and 20000)

Table 2 Average dependency of time required to compute new coordinates and Stress function values per iteration on data size when n = 10, using SMACOF-Fast, SMACOF and Geometric MDS.

Number of points, <i>m</i>							Ratio					
	SMACOF		SMACOF-Fast		Geometric MDS		SMACOF/ SMACOF-Fast		Geometric MDS/SMACOF-Fast			
	Stress	Time, s	Stress	Time, s	Stress	Time, s	Stress ratio	Time ratio	Stress ratio	Time ratio		
10	11.6	0.00019	11.6	0.00003	11.7	0.00003	1.0000000	7.3152985	1.0095220	1.2097938		
20	66.7	0.00036	66.7	0.00005	66.7	0.00006	1.0000000	6.9684499	1.0011044	1.1193416		
50	479.1	0.00056	479.1	0.00009	478.8	0.00009	1.0000000	6.3983110	0.9992541	1.0824801		
100	2128.1	0.00093	2128.1	0.00016	2127.0	0.00017	1.0000000	5.9893883	0.9995029	1.0680442		
200	8946.6	0.00194	8946.6	0.00035	8944.1	0.00037	1.0000000	5.5188379	0.9997245	1.0489429		
500	51188.6	0.00885	51188.6	0.00246	51183.4	0.00267	1.0000000	3.5984986	0.9998969	1.0861585		
1000	218604.8	0.03845	218604.8	0.01419	218594.0	0.01452	1.0000000	2.7084907	0.9999505	1.0227548		
2000	894694.9	0.18151	894694.9	0.06169	894672.8	0.06175	1.0000000	2.9422145	0.9999753	1.0009971		
5000	5106657.3	1.15621	5106657.3	0.32358	5106605.3	0.32322	1.0000000	3.5731957	0.9999898	0.9988961		
10000	21892993.0	5.06069	21892993.0	1.27452	21892881.6	1.26265	1.0000000	3.9706513	0.9999949	0.9906829		
15000	59857709.1	14.66615	59857709.1	3.51422	59857507.2	3.46431	1.0000000	4.1733769	0.9999966	0.9857980		
20000	127201601.7	32.07792	127201601.7	7.49513	127201281.2	7.45024	1.0000000	4.2798321	0.9999975	0.9940104		

 Table 3

 Average dependence of the time required to compute new coordinates and Stress function values per iteration on the data size on random proximity matrices whose element takes uniformly distributed values in the interval (0, 1), using SMACOF-Fast, SMACOF and Geometric MDS.

							Ratio					
Number of points, <i>m</i>	SMACOF		SMACOF-Fast		Geometric MDS		SMACOF/ SMACOF-Fast		Geometric MDS/SMACOF-Fast			
	Stress	Time, s	Stress	Time, s	Stress	Time, s	Stress ratio	Time ratio	Stress ratio	Time ratio		
10	3.812807	0.00018	3.812807	0.00003	3.738581	0.00003	1.00000000	6.20922293	0.98053234	1.01140901		
20	22.02273	0.00036	22.02273	0.00006	21.81476	0.00006	1.00000000	6.34562798	0.99055667	1.02805624		
50	164.0261	0.00057	164.0261	0.00010	163.6571	0.00010	1.00000000	6.00706317	0.99774989	1.02165017		
100	788.3272	0.00112	788.3272	0.00017	787.7029	0.00017	1.00000000	6.75047562	0.99920800	1.02180998		
200	3295.794	0.00230	3295.794	0.00035	3294.927	0.00035	1.00000000	6.49822251	0.99973703	0.99185505		
500	19294.97	0.00888	19294.97	0.00272	19293.7	0.00286	1.00000000	3.26174940	0.99993376	1.04896365		
1000	83109.29	0.03834	83109.29	0.01447	83107.34	0.01415	1.00000000	2.65000273	0.99997652	0.97795314		
2000	339234.9	0.17323	339234.9	0.05924	339231.9	0.05665	1.00000000	2.92438158	0.99999104	0.95639967		
5000	1940633	1.15612	1940633	0.30542	1940628	0.29682	1.00000000	3.78540252	0.99999716	0.97186003		
10000	8348942	5.42154	8348942	1.22316	8348932	1.23237	1.00000000	4.43240329	0.99999878	1.00752752		
15000	22765379	15.06911	22765379	3.37158	22765362	3.44193	1.00000000	4.46944395	0.99999926	1.02086564		
20000	48401588	32.37652	48401588	7.35390	48401562	7.41715	1.00000000	4.40263131	0.99999946	1.00860033		

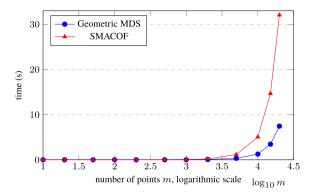


Fig. 3. Dependence of computation time on the number of points when reducing the dimensionality of 10dimensional data (n = 10) to d = 2. The number of points (m = 10, 20, 50, 100, 200, 500, 1000, 2000, 5000, 10000, 15000, and 20000) is given in a logarithmic scale. All results are averaged, as a result of each experiment being repeated 100 times to calculate one iteration.

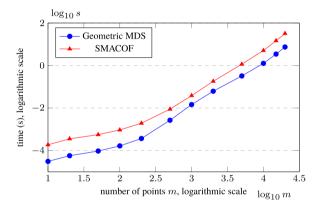


Fig. 4. Dependence of computation time on the number of points when reducing the dimensionality of 10dimensional data (n = 10) to d = 2. The time of calculation and the number of points (m = 10, 20, 50, 100, 200, 500, 10000, 15000, and 20000) are given in a logarithmic scale. All results are averaged, as a result of each experiment being repeated 100 times to calculate one iteration.

is given in a logarithmic scale. Each experiment was repeated 100 times, so the figures summarise the results with the average time and the value of the Stress function for each case of m. In order to provide the results in a more informative form, and to emphasize the differences in the time required to calculate new coordinates using Geometric MDS (see Listing 2) and SMACOF (Pedregosa *et al.*, 2011), the time of calculation and the number of points are additionally given in a logarithmic scale in Fig. 4. The results demonstrate that Geometric MDS (Listing 2) outperforms the SMACOF (from the scikit-learn library, Pedregosa *et al.*, 2011) in all cases in terms of the computation time required to recalculate the coordinates of new points Y_1, \ldots, Y_m in the projected space. Comparing the results obtained using SMACOF, SMACOF-Fast and Geometric MDS, it can be observed that

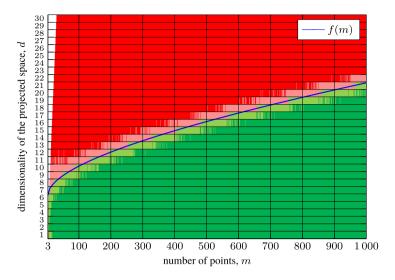


Fig. 5. Stress comparison between Geometric MDS and SMACOF depending on the number of points m and and the dimensionality of the projected space d.

with increasing data size Geometric MDS requires less computing time and the resulting stress is smaller than that by SMACOF under the same conditions (see Fig. 4, Tables 2 and 3). Meanwhile, the time required to recalculate new coordinates using Geometric MDS and SMACOF-Fast is approximately the same.

4.2. Comparative Efficiency of Geometric MDS and SMACOF

To compare which method, Geometric MDS or SMACOF, better reduces the MDS stress $S(\cdot)$ defined by (1), we tested these methods on random proximity matrices whose elements take uniformly distributed values in the interval (0, 1). 101 tests were performed with different m = 3, ..., 1000 and d = 1, ..., 30. The experiments were carried out using the programming implementations provided earlier in this section: SMACOF-Fast based on Listing 3 and Geometric MDS based on Listing 2. The results of the experiments are given in Fig. 5, which visually shows which algorithm, SMACOF or Geometric MDS, get a smaller value of the stress function $S(\cdot)$ after updating the points $Y_1, ..., Y_m$ only once, i.e. after one iteration, depending on the number m of points and the dimensionality d of the projected space. The red colour indicates cases in which SMACOF stress was superior to Geometric MDS in all cases, the light red indicates cases in which SMACOF stress was superior to Geometric MDS in more cases, the light green indicates cases in which SMACOF in all cases.

Figure 5 discloses a very interesting and essential difference between the performance of Geometric MDS and SMACOF. Geometric MDS finds smaller stress when the dimensionality of projected space is lower. This is particularly important for the multidimensional data visualization task.

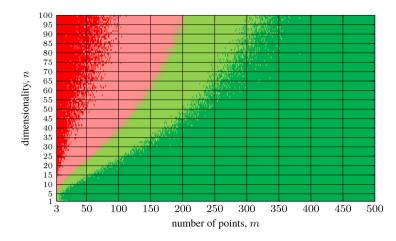


Fig. 6. Stress comparison between Geometric MDS and SMACOF depending on the number of points *m* and the dimensionality *n*, when reducing the dimensionality to d = 2.

In Fig. 5, we present a curve that would allow us to choose either Geometric MDS or SMACOF in order to get a better stress after their first step and assuming that the values of the elements of the symmetric distance matrix are uniformly distributed in the interval (0, 1). The upper part of the parabola was used to determine the distinguishing curve. The optimization package of the MAPLE software (Bernardin *et al.*, 2021) was used to define the coefficients in the equation that is as follows in this particular case:

$$f(m) \approx 0.003646m + 0.362826\sqrt{m-3} + 6.782796.$$

Particular case is d = 2, because it is the most used in multidimensional data visualization. Figure 6 illustrates the values of the stress function $S(\cdot)$ depending on the number of points m and the dimensionality n of the analysed multidimensional data. By random generating a set of *m n*-dimensional points X_1, \ldots, X_m , the distance matrix $m \times m$ was calculated. Then a random set of 2-dimensional (d = 2) points Y_1, \ldots, Y_m was selected as a starting position for optimization of the stress. The coordinates of these points were recalculated in one iteration using Geometric MDS and SMACOF (see Listings 2 and 3). The colours in Fig. 6 were labelled using the same rules as in Fig. 5: red means that in 101 such independent trials, SMACOF gave better stress in all cases, light red means that SMACOF gave better stress in more cases than Geometric MDS, light green means that Geometric MDS gave better stress in more cases than Geometric MDS but not in all cases, and light green means that Geometric MDS gave better stress in all cases. Figure 6 shows visually whether Geometric MDS or SMACOF gives lower stress depending on the number of points *m* and on the dimensionality *n* after one iteration of 101 tries. The results in Fig. 6 can be useful for choosing an appropriate method when the dimensionality n of the data to be analysed and the number of points n are known.

5. Conclusions

A new Geometric MDS method with the low computational complexity has been proposed in Dzemyda and Sabaliauskas (2021a, 2022). Geometric MDS provides an iterative procedure to minimize MDS stress where coordinates of a particular point of the projected space are moved to a new position defined analytically. Such a change in position is easily interpreted geometrically. Moreover, the coordinates of points of the projected space may be recalculated simultaneously, i.e. in parallel, independently of each other during one iteration of stress minimization. Two implementations of Geometric MDS are suggested and analysed experimentally: parallel and sequential.

A parallel implementation of Geometric MDS is developed for dimensionality reduction using multithreaded multi-core processors. Based on the results obtained, we can claim that Geometric MDS allows to implement parallel computing using multithreaded multi-core processors, as a result, the time to calculate new coordinates of points in the low-dimensional space may be reduced by 6 times on average, depending on the data size.

The sequential implementation given in this paper is optimized for computational speed, enabling it to solve large data problems. It is compared with the SMACOF version of MDS. Python codes for both Geometric MDS and SMACOF are presented to highlight the differences between the two implementations. These codes were optimized for computational speed. The comparison was carried out on several aspects: the comparative performance of Geometric MDS and SMACOF depending on the projection dimension, data size and computation time. Geometric MDS usually finds lower stress when the dimensionality of the projected space is smaller.

SMACOF is an iterative procedure that calculates the Guttman transform. We discovered that the procedure of updating lower-dimensional points by Geometric MDS is a generalization of the Guttman transform. If we apply $\frac{1}{m}$ instead of $\frac{1}{m-1}$ in Eq. (7) to all the lower-dimensional points Y_1, \ldots, Y_m simultaneously and independently of each other during a single iteration of stress minimization, we get the same result as with the Guttman transform. However, it is proved in Dzemyda and Sabaliauskas (2022) that both Geometric MDS (GMDSm) and SMACOF make a step from the point Y_j to its new position in the same direction with different step size. The step size by Geometric MDS is $\frac{m}{m-1}$ times longer. In both cases, the stress (1) decreases. Moreover, Geometric MDS allows to update coordinates of a single point Y_j , and it guarantees the decrease of the stress, too. These are the reasons for further exploring the potential of Geometric MDS and applying the advantages discovered.

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