Distance-based dimensionality reduction for big data

Master's thesis defense

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1. Introduction, motivation and

objectives

Introduction, motivation and objectives

- Dimensionality reduction (DR) aims to project a dataset into a low-dimensional space.
- Most DR techniques are based on the inter-individual distance matrix ⇒ they have quadratic memory complexity.
- There are algorithms that extend classical MDS to the big data settings.
- In this master's thesis, we adapt one of these algorithms to any generic distance-based DR method.

2. State of the art

A few dimensionality reduction techniques

Non-classical MDS

The SMACOF algorithm minimizes metric stress using a majorization technique (Kruskal, 1964a; Kruskal, 1964b).

Isomap

Preserves geodesic distances between points in a manifold (Tenenbaum, de Silva, and Langford, 2000).

LMDS

A repulsive term between distant points is added to the stress function (Chen and Buja, 2009).

t-SNE

Models similarities between points as conditional probabilities (Maaten and Hinton, 2008).

Multidimensional scaling for big data

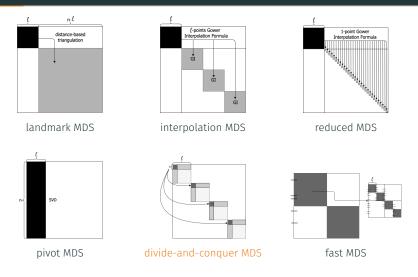


Figure 1: Schematic representation of the six MDS algorithms for big data described in Delicado and Pachón-García, 2024 (Source: original publication).

3. Specification and design of the

solution

Divide-and-conquer dimensionality reduction (1/3)

Algorithm 1 Divide-and-conquer dimensionality reduction

Require: $\mathbf{D} = (\delta_{ij})$, the $n \times n$ matrix of observed distances; \mathcal{M} , the DR method; l, the partition size; c, the amount of connecting points; q, the embedding's dimensionality; and arg, \mathcal{M} 's specific parameters.

Ensure: \widetilde{Y} , a configuration in a q-dimensional space.

- 1: if $n \leq l$ then return $\mathcal{M}(D, q, arg)$
- 2: end if
- 3: Let $k = \lceil \frac{n-l}{l-c} \rceil$
- 4: Randomly partition the data: $\mathcal{P} = \{\mathcal{P}_0, \mathcal{P}_1, \dots, \mathcal{P}_k\}$ where

$$|\mathcal{P}_i| = \begin{cases} l & \text{if } i = 0\\ l - c & \text{if } 0 < i \le (n - l) \bmod k\\ l - c - 1 & \text{if } (n - l) \bmod k < i \le k \end{cases}$$

Divide-and-conquer dimensionality reduction (2/3)

- 5: Sample c connecting points from \mathcal{P}_0 : $\mathcal{C} \subset \mathcal{P}_0$
- 6: Extract distance matrix of \mathcal{P}_0 : $D_{\mathcal{P}_0} = D[\mathcal{P}_0, \mathcal{P}_0]$
- 7: Apply DR method to \mathcal{P}_0 : $\widetilde{Y}_0 = \mathcal{M}(D_{\mathcal{P}_0}, q, arg)$
- 8: Extract embedding of $C: \widetilde{Y}_{C} = \widetilde{Y}_{0}[C,:]$
- 9: Extract distance matrix of \mathcal{C} : $D_{\mathcal{C}} = D[\mathcal{C}, \mathcal{C}]$

Divide-and-conquer dimensionality reduction (3/3)

```
10. for i = 1 to k do
             Extract distance matrix of \mathcal{P}_i: D_{\mathcal{P}_i} = D[\mathcal{P}_i, \mathcal{P}_i]
11.
             Stack connecting points to \mathcal{P}_i: D_{\text{stack}} = [D_{\mathcal{C}}; D_{\mathcal{P}_i}]
12:
            Project the stacked data: \widetilde{Y}_{\text{stack}} = \mathcal{M}(D_{\text{stack}}, q, arg)
13·
            Split embeddings: \widetilde{Y}_{c}^{(i)} = \widetilde{Y}_{\text{stack}}[:c,:] and \widetilde{Y}_{i} = \widetilde{Y}_{\text{stack}}[(c+1):,:]
14:
            Align first and current embeddings: \widetilde{\mathbf{Y}}_i = \text{Procrustes}(\widetilde{\mathbf{Y}}_c, \widetilde{\mathbf{Y}}_c^{(i)}, \widetilde{\mathbf{Y}}_i)
15:
16: end for
17: Combine all embeddings: \widetilde{Y}' = [\widetilde{Y}_0; \widetilde{Y}_1; \dots; \widetilde{Y}_b]
18: Retrieve original row ordering: order = argsort([\mathcal{P}_0; \mathcal{P}_1; \dots; \mathcal{P}_k])
19: Set original ordering: \widetilde{Y}' = \widetilde{Y}' [order, :]
20: Apply PCA to center and rotate data: \widetilde{Y} = PCA(\widetilde{Y}', q)
               return Y
```

Orthogonal Procrustes transformation's derivation

Let $A \in \mathbb{R}^{c \times q}$ be the target configuration and $B \in \mathbb{R}^{c \times q}$ the corresponding testee. We wish to fit B to A by rigid motions. That is, we want to find the best orthogonal matrix T such that $A \simeq BT$. We will measure the \simeq relation with the sum-of-squares criterion L and try to minimize it:

$$\min_{\mathsf{T}\in\mathsf{O}(q)}\mathsf{L}(\mathsf{T})=\min_{\mathsf{T}\in\mathsf{O}(q)}\mathsf{tr}(\mathsf{A}-\mathsf{BT})(\mathsf{A}-\mathsf{BT})',$$

Berge, Kiers, and Commandeur (1993) found the following solution. Let $U\Sigma V'$ be the singular value decomposition of A'B, where U'U=I,V'V=I, and Σ is the diagonal matrix with the singular values. Then, L(T) is minimal if T=VU'.

4. Development of the proposal

Python implementation

- divide_conquer implements Algorithm 1 in parallel through the concurrent.futures module.
- · Implementations of DR algorithms used:
 - sklearn.manifold module (Pedregosa et al., 2011) for Isomap and SMACOF.
 - · openTSNE (Poličar, 2023) for t-SNE.
 - A translation of the R library smacofx (Leeuw and Mair, 2009) for LMDS.
- Time complexity is reduced from quadratic (or cubic for Isomap) to linear.
- Space complexity is lowered from $\mathcal{O}(n^2)$ to $\mathcal{O}(l^2)$.

Test datasets

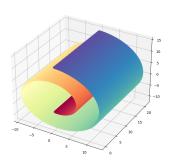


Figure 2: Swiss roll

Unfolded rectangle

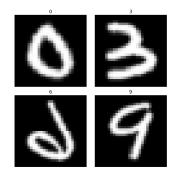


Figure 3: MNIST



Experimental setup

- 1. Tune the bare DR method on a *l* points subset.
- 2. Apply the bare DR method on a larger subset.
- 3. Apply divide-and-conquer DR on a larger subset.
- 4. Apply the bare DR method on the whole dataset (when possible).
- 5. Apply divide-and-conquer DR on the whole dataset.

The testing system was an Asus ROG G513QM-HF026 laptop with

- Windows
- · AMD Ryzen 7 5800H CPU
- · NVIDIA RTX 3060 GPU
- · 16 GB of DDR4-3200MHz RAM
- 1 TB M.2 NVMe PCIe 3.0

5. Experimentation and

evalutation of the proposal

Runtime benchmarks of divide-and-conquer Isomap and divideand-conquer t-SNE

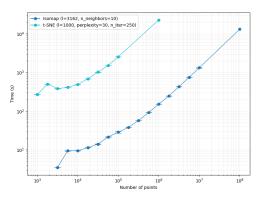


Figure 4: Runtime (s) of divide-and-conquer Isomap and divide-and-conquer t-SNE averaged over 20 experiments. Tests were performed on datasets generated on the Swiss roll manifold with sizes ranging from 10^3 to 10^8 . Data was embedded into \mathbb{R}^2 with different parameter combinations and c=100.

SMACOF's embedding of Swiss roll

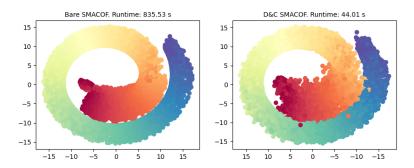


Figure 5: Comparison of the bidimensional embeddings of a 7,500 points Swiss roll dataset by bare (left) and divide-and-conquer (right) SMACOF. The arguments used were $n_iter = 300$, $\varepsilon = 0.001$ and in divide-and-conquer there also were l = 1000 and c = 100. Color represents the angle of rotation along the Swiss roll spiral.

LMDS's embedding of Swiss roll

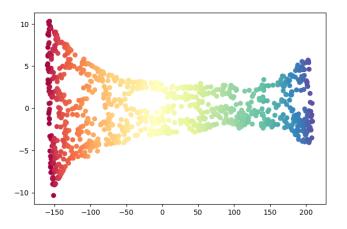


Figure 6: Bidimensional embedding of a 1,000 points Swiss roll dataset computed by LMDS with k=10 and $\tau=0.1$. Color represents the angle of rotation along the Swiss roll spiral.

Divide-and-conquer Isomap's embedding of Swiss roll

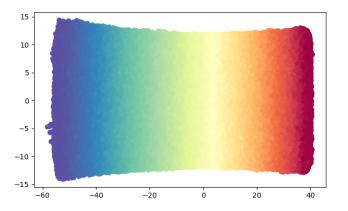


Figure 7: Bidimensional embedding of a 10^8 points Swiss roll dataset computed by divide-and-conquer Isomap with k = 10, l = 3,162 and c = 100. Color represents the angle of rotation along the Swiss roll spiral.

Divide-and-conquer t-SNE's embedding of Swiss roll

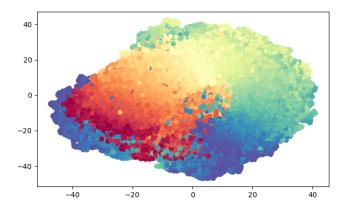


Figure 8: Bidimensional embedding of a 10^6 points Swiss roll dataset computed by divide-and-conquer t-SNE with $l=1,000,\,c=100,\,Perp=30$ and $n_iter=250$. Color represents the angle of rotation along the Swiss roll spiral.

SMACOF's embedding of a 5,000 points subset of MNIST (1/2)

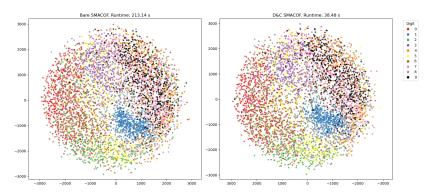


Figure 9: Bidimensional embeddings of a 5,000 points subset of MNIST by bare (left) and divide-and-conquer (right) SMACOF. The arguments we used were $n_iter = 300$, $\varepsilon = 0.001$ and in divide-and-conquer there also were l = 1000 and c = 100.

SMACOF's embedding of a 5,000 points subset of MNIST (2/2)

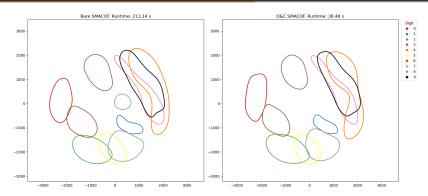


Figure 10: Kernel density estimation of the bidimensional embeddings of a 5,000 points subset of MNIST by bare (left) and divide-and-conquer (right) SMACOF. The arguments we used were $n_iter = 300$, $\varepsilon = 0.001$ and in divide-and-conquer there also were l = 1000 and c = 100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

Divide-and-conquer SMACOF's embedding of the whole MNIST

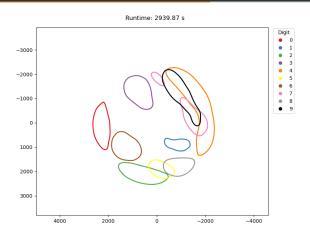


Figure 11: Kernel density estimation of the bidimensional embeddings of the whole MNIST dataset by divide-and-conquer SMACOF. The arguments used were $n_iter = 300$, $\varepsilon = 0.001$, l = 1000 and c = 100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

LMDS's embedding of a 5,000 points subset of MNIST

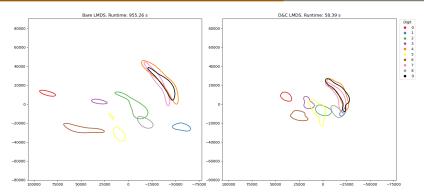


Figure 12: Kernel density estimation of the bidimensional embeddings of a 5,000 points subset of MNIST by bare (left) and divide-and-conquer (right) LMDS. The arguments used were k=10, $\tau=1$ and in divide-and-conquer there also were l=1000 and c=100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

Divide-and-conquer LMDS's embedding of the whole MNIST

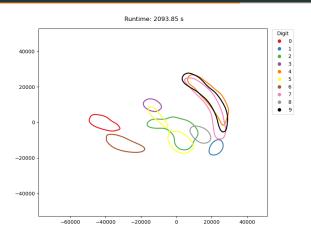


Figure 13: Kernel density estimation of the bidimensional embeddings of the whole MNIST dataset by divide-and-conquer LMDS. The arguments used were $k=10, \ \tau=1, \ l=1000$ and c=100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

Isomap's embedding of a 5,000 points subset of MNIST

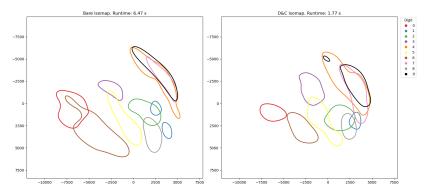


Figure 14: Kernel density estimation of the bidimensional embeddings of a 5,000 points subset of MNIST by bare (left) and divide-and-conquer (right) Isomap. The arguments used were k=5 and in divide-and-conquer there also were l=1000 and c=100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

Divide-and-conquer Isomap's embedding of the whole MNIST

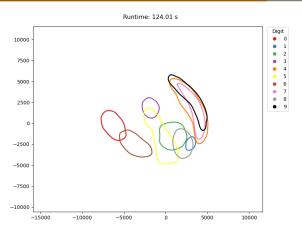


Figure 15: Kernel density estimation of the bidimensional embeddings of the whole MNIST dataset by divide-and-conquer Isomap. The arguments used were $k=5,\ l=1000$ and c=100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

t-SNE's embedding of the whole MNIST

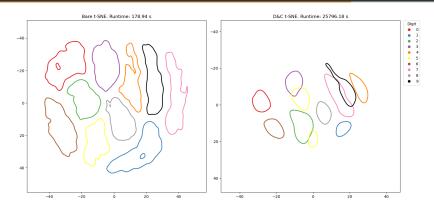


Figure 16: Kernel density estimation of the bidimensional embeddings of the whole MNIST dataset by bare (left) and divide-and-conquer (right) t-SNE. The arguments used were Perp = 20, $n_iter = 100$ and in divide-and-conquer there also were l = 1000 and c = 100. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

t-SNE's insconsistency

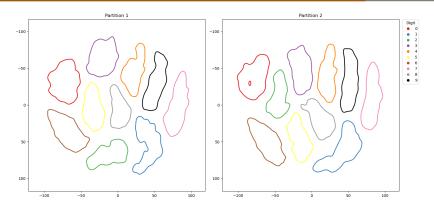


Figure 17: Kernel density estimation of the bidimensional embeddings of two halves of the MNIST dataset. Data was randomly ordered before being splitted. The DR method used was divide-and-conquer t-SNE with Perp=30. Contour lines are at 70% of the maximum estimated density for each digit and embedding.

6. Analysis of sustainability and

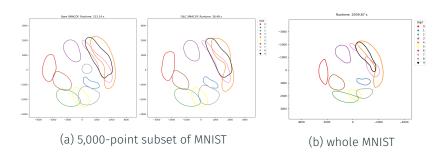
ethical implications

GHG emissions

- Data centers generate significant GHG emissions due to high electricity usage.
- Our divide-and-conquer DR framework reduces runtime and hardware demands, lowering emissions.
- It enables sustainable DR by decreasing dependence on supercomputers and improving efficiency.

Visibility of small communities

- DR methods can emphasize biases.
- However, more data ⇒ more likely to represent small communities.



7. Conclusions

Conclusions

- Developed a general divide-and-conquer framework for distancebased DR methods, reducing time and memory complexities.
- Achieved strong embedding quality on large datasets, notably projecting a 10⁸ points Swiss roll in 3 h on a standard computer.
- Contributed to making advanced DR techniques more accessible and sustainable for big datasets.

Future work:

- Formalize the framework into a Python package.
- Analyze the effect of c on performance and embedding quality.
- Investigate why LMDS cannot unroll the Swiss roll manifold.

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Thank you!

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