

HIGHLIGHTS

- Combines non-convex Riemannian matrix completion method and a dual-basis framework
- Comparable reconstruction to state-of-the-art algorithms
- Provable convergence framework

EUCLIDEAN DISTANCE GEOMETRY

Euclidean Distance Geometry: Given partial pairwise squared distances $\mathbf{D} = [d_{ij}^2]$ in a matrix, where only some entries are known, can we robustly reconstruct the points $\mathbf{P} = [\mathbf{p}_1 \dots \mathbf{p}_n]^T \in \mathbb{R}^{n \times d}$ up to rotation/translation?

Multi-dimensional Scaling (MDS): Recovers \mathbf{P} up to rotation from full information in \mathbf{D} by taking a truncated eigenvalue decomposition of $\mathbf{X} = -\frac{1}{2}(\mathbf{I}_n - \frac{1}{n}\mathbf{1}\mathbf{1}^T)\mathbf{D}(\mathbf{I}_n - \frac{1}{n}\mathbf{1}\mathbf{1}^T)$

Matrix Completion: Algorithms for computing a low-rank matrix $\mathbf{M} \in \mathbb{R}^{n_1 \times n_2}$ given a subset of the entries $\Omega = \{(i, j) \in [n_1] \times [n_2] \mid M_{ij} \text{ is known}\}$. Original methods[1] developed were convex minimizations of the nuclear norm

$$\min_{\mathbf{X} \in \mathbb{R}^{n_1 \times n_2}} \|\mathbf{X}\|_* \text{ subject to } \mathcal{P}_\Omega(\mathbf{X}) = \mathcal{P}_\Omega(\mathbf{M})$$

where \mathcal{P}_Ω is defined as

$$\mathcal{P}_\Omega(\cdot) = \sum_{(i,j) \in \Omega} \langle \cdot, \mathbf{E}_{ij} \rangle \mathbf{E}_{ij}$$

for $\mathbf{E}_{ij} = \mathbf{e}_i \mathbf{e}_j^T$. Many scalable non-convex algorithms for this problem exist.

Problems with existing methods: Distance matrices are a difficult set to optimize over due to the triangle inequality. This leads to poor recovery results with standard matrix completion algorithms on \mathbf{D} .

EXISTING WORK AND GEOMETRIC STRUCTURE

Riemannian Methods for Matrix Completion [2]: A non-convex Riemannian approach to matrix completion.

- **Main idea:** Non-convex gradient descent scheme for matrix completion using entries.
- **Formulation:** Wei et al. define the following optimization program

$$\min_{\mathbf{X} \in \mathbb{R}^{n \times n}} \langle \mathbf{X} - \mathbf{M}, \mathcal{P}_\Omega(\mathbf{X} - \mathbf{M}) \rangle \text{ subject to } \text{rank}(\mathbf{X}) = r$$

The algorithm is a gradient descent scheme on the manifold of rank r matrices with a tangent space at the l -th iterate $\mathbf{X}_l = \mathbf{U}_l \Sigma_l \mathbf{V}_l^T$ defined as $\mathbb{T}_l = \{\mathbf{U}_l \mathbf{Z}_1^T + \mathbf{Z}_2 \mathbf{V}_l \mid \mathbf{Z}_1, \mathbf{Z}_2 \in \mathbb{R}^{n \times r}\}$. To update to \mathbf{X}_{l+1} , the update is taken in the gradient descent direction projected onto the manifold \mathbb{T}_l , then retracted back to the rank r manifold. More specifically

$$\mathbf{X}_{l+1} = \text{SVD}_r(\mathbf{X}_l + \eta_l \mathcal{P}_{\mathbb{T}_l} \mathcal{P}_\Omega(\mathbf{M} - \mathbf{X}_l))$$

with SVD_r defined as the truncated SVD of rank r and η_l computed through an exact line search

- **Pros:** Proven convergence results, efficient implementation
- **Cons:** Poor recovery for the EDG problem

DUAL BASIS APPROACH

Idea: Instead of optimizing over distance matrices, move to Gram matrices for easier computability.

Constructing Dual Basis and Sampling Operator: Following [3], accessible information is in the form of

$$D_{ij} = \|\mathbf{p}_i - \mathbf{p}_j\|_2^2 = \|\mathbf{p}_i\|_2^2 + \|\mathbf{p}_j\|_2^2 - 2\mathbf{p}_i^T \mathbf{p}_j = X_{ii} + X_{jj} - 2X_{ij}$$

Defining $\mathbf{w}_\alpha = \mathbf{E}_{\alpha_1, \alpha_1} + \mathbf{E}_{\alpha_2, \alpha_2} - \mathbf{E}_{\alpha_1, \alpha_2} - \mathbf{E}_{\alpha_2, \alpha_1}$ for $\alpha = (\alpha_1, \alpha_2)$, we can represent accessible information as $\langle \mathbf{X}, \mathbf{w}_\alpha \rangle$. Given this new basis and its Gram matrix \mathbf{H} defined by $\mathbf{H}_{\alpha, \beta} = \langle \mathbf{w}_\alpha, \mathbf{w}_\beta \rangle$, the dual or bi-orthogonal basis can be constructed as

$$\mathbf{v}_\alpha = \sum_{\beta} \mathbf{H}_{\alpha, \beta}^{-1} \mathbf{w}_\beta$$

This allows us to define an analogous sampling operator for the dual basis problem:

$$\mathcal{R}_\Omega(\cdot) := \frac{L}{m} \sum_{\alpha \in \Omega} \langle \cdot, \mathbf{w}_\alpha \rangle \mathbf{v}_\alpha$$

This problem defined on Gram matrices is mathematically equivalent to standard matrix completion on the squared distance matrix, although as \mathcal{R}_Ω is not self-adjoint we consider a computable surrogate instead.

Defining Computable Surrogate and Optimization Program: We construct a computable surrogate and its corresponding objective function as follows:

$$\mathcal{R}_\Omega^* \mathcal{R}_\Omega(\cdot) := \frac{L^2}{m^2} \sum_{\alpha, \beta} \langle \cdot, \mathbf{w}_\alpha \rangle \langle \mathbf{v}_\alpha, \mathbf{v}_\beta \rangle \mathbf{w}_\beta$$

$$\min_{\mathbf{X} \in \mathbb{R}^{n_1 \times n_2}} \langle \mathbf{X} - \mathbf{M}, \mathcal{R}_\Omega^* \mathcal{R}_\Omega(\mathbf{X} - \mathbf{M}) \rangle \text{ subject to } \text{rank}(\mathbf{X}) = r$$

RIEEDG

Algorithm: Fusing the dual-basis approach with the efficient Riemannian scheme presented in [2].

- **Main idea:** Define a similar algorithm as in [2], but substituting our computable surrogate operator $\mathcal{R}_\Omega^* \mathcal{R}_\Omega$
- **Pros:** Provable convergence framework given good enough initialization
- **Cons:** Slower time complexity with similar reconstruction results as other non-convex algorithms [3]

Algorithm: RieEDG

Input: $\mathcal{P}_\Omega(\mathbf{D})$: The observed distance information; k : the dimension of the datapoints; η : the step size

Initialize $\mathbf{X}_0 = \text{EVD}_k(\mathcal{R}_\Omega^* \mathcal{R}_\Omega(\mathbf{X})) = \mathbf{U}_0 \Lambda_0 \mathbf{U}_0^T$

for $l = 0, 1, 2 \dots$ **do**

$\mathbf{G}_l = \mathcal{R}_\Omega^* \mathcal{R}_\Omega(\mathbf{X} - \mathbf{X}_l)$

$\mathbf{W}_l = \mathbf{X}_l + \eta \mathcal{P}_{\mathbb{T}_l} \mathbf{G}_l$

$\mathbf{X}_{l+1} = \text{EVD}_k(\mathbf{W}_l)$

end for

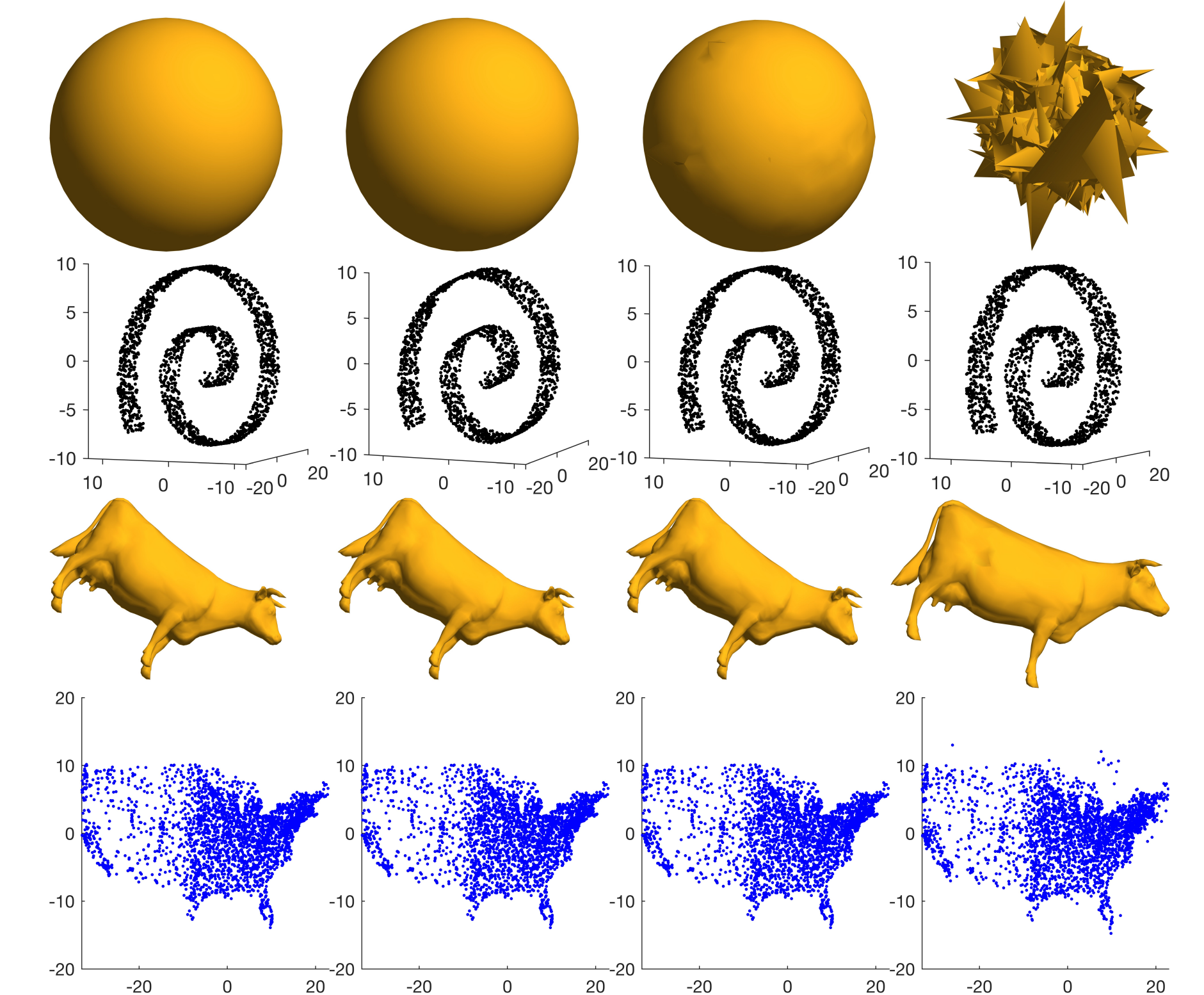
Output: \mathbf{X}_{rev}

NUMERICAL EXPERIMENTS

Synthetic data and Tabulated Results: Various 2- and 3-dimensional datasets were used for testing and are referred to below in increasing size order. The objective of RieEDG is to recover the full set of points \mathbf{P} up to orthogonal transformation from a subset of entries of \mathbf{D} chosen using a Bernoulli sampling model, where each entry has a probability γ of being selected for $\gamma \in [0, 1]$, with an expected γL entries chosen. RieEDG outputs the Gram matrix $\mathbf{X} = \mathbf{P}\mathbf{P}^T$, from which \mathbf{P} can be recovered. The comparison referenced in Table is the relative error between the recovered matrix \mathbf{X}_{rev} and the ground truth matrix \mathbf{X} in Frobenius norm averaged over 10 trials. Each run was terminated after 500 iterations or a relative difference of 10^{-7} in Frobenius norm.

Dataset	$\gamma = 5\%$	3%	2%	1%	5% Timing (sec)
Sphere (3D)	6.2e-07	1.2e-06	9.52e-03	1.08	4.62
Swiss Roll (3D)	5.04e-07	8.84e-07	1.14e-06	0.0604	30.9
Cow (3D)	5.58e-07	8.62e-06	1.50e-06	0.0095	67.4
U.S. Cities (2D)	5.90e-07	1.613-03	0.0168	0.0796	135

Reconstructed Images: Below are images of the reconstructed datasets. From left to right, the sampling rate goes from 5% to 1% as in the table above.



REFERENCES

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