

Near-Optimal Joint Object Matching via Convex Relaxation

Yuxin Chen*, Leonidas J. Guibas† and Qi-Xing Huang‡

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Abstract

Joint matching over a collection of objects aims at aggregating information from a large collection of similar instances (e.g. images, graphs, shapes) to improve maps between pairs of them. Given multiple objects and matches computed between a few object pairs in isolation, the goal is to recover an entire collection of maps that are (1) globally consistent, and (2) close to the provided maps — and under certain conditions provably the ground-truth maps. Despite recent advances on this problem, the best-known recovery guarantees are limited to a small constant barrier — none of the existing methods find theoretical support when more than 50% of input correspondences are corrupted. Moreover, prior approaches focus mostly on fully similar objects, while it is practically more demanding to match instances that are only partially similar to each other (e.g., different views of a single physical object).

In this paper, we propose an algorithm to jointly match multiple objects that exhibit only partial similarities, given a few (possibly highly incomplete) pairwise matches that are densely corrupted. By encoding a consistent partial map collection into a 0-1 semidefinite matrix, we propose to recover the ground-truth maps via a parameter-free convex program called MatchLift, following a spectral method that pre-estimates the total number of distinct elements to be matched. Numerically, this program can be efficiently solved via alternating direction methods of multipliers (ADMM) along with a greedy rounding strategy. Theoretically, MatchLift exhibits near-optimal error-correction ability, i.e. in the asymptotic regime it is guaranteed to work even when a dominant fraction $1 - \Theta\left(\frac{\log^2 n}{\sqrt{n}}\right)$ of the input maps behave like random outliers. Furthermore, MatchLift succeeds with minimal input complexity, namely, perfect matching can be achieved as soon as the provided maps form a connected map graph. We evaluate the proposed algorithm on various benchmark data sets including synthetic examples and real-world examples, all of which confirm the practical applicability and usefulness of MatchLift.

Index Terms: Joint graph matching, shape mapping, cycle consistency, dense error correction, partial similarity, convex relaxation, spectral methods, robust PCA, matrix completion, graph clustering, ADMM, MatchLift

1 Introduction

Finding consistent relations across multiple objects is a fundamental scientific problem spanning many fields. A partial list includes jigsaw puzzle solving [1, 2], structure from motion [3, 4], re-assembly of fragmented objects and documents [5, 6], and DNA/RNA shotgun assembly sequencing [7]. Compared with the rich literature in pairwise matching (e.g. of graphs, images or shapes), joint matching of multiple objects has not been well explored. A naive approach for joint object matching is to pick a base object and perform *pairwise matching* with each of the remaining objects. However, as pairwise matching algorithms typically generate noisy results, the performance of such approaches is often far from satisfactory in practice. This gives rise to the question as to how to aggregate and exploit information from all pairwise maps that one computes, in order to improve joint object matching in a consistent and efficient manner.

In this paper, we represent each object as a discrete set of points or elements, and investigate the problem of joint matching over n different sets, for which the input / observation is a collection of pairwise maps

*Y. Chen is with the Department of Electrical Engineering, Stanford University (email: yxchen@stanford.edu).

†L. J. Guibas is with the Department of Computer Science, Stanford University (email: guibas@stanford.edu).

‡Q-X. Huang is with the Department of Computer Science, Stanford University (email: huangqx@stanford.edu).

computed in isolation. A natural and popular criterion to preserve the global relational compatibility is called cycle-consistency, i.e., that composition of maps between two objects should be independent of the connecting path chosen. Such criterion has recently been invoked in many algorithms [3, 8–11] to detect outliers among the pairwise input maps. These works have shown experimentally that one can use inconsistent cycles to prune outliers, provided that the corruption rate is sufficiently small.

Despite the empirical advances of these works, little is known on the theoretical side, namely, under what conditions can the underlying ground-truth maps be reliably recovered. Recent work by [12] provided the first theoretical guarantee for robust and consistent joint matching. However, there are several fundamental issues left unaddressed that must be faced in order to accommodate practical challenges.

1. **Dense Input Errors:** The state-of-the-art results (e.g. [12]) did not provide theoretical support when more than 50% of the input matches are corrupted. This gives rise to the question regarding their applicability in the presence of highly noisy sources, in which case the majority of the input maps can be corrupted. Observe that as the number n of objects to be matched increases, the amount of pairwise maps one can obtain significantly exceeds n . As a result, dense error correction is information theoretically possible as long as the global consistency across pairwise maps can be appropriately exploited. While one would expect an ideal algorithm to work even when most input maps are random outliers, the challenge remains as to whether there exist *computationally feasible* methods that can provably detect and separate dense outliers.
2. **Partial Similarity:** To the best of our knowledge, all prior approaches dealt only with a restricted scenario where the ground-truth maps are given by full isomorphisms (i.e. *one-to-one* correspondences between any two sets). In reality, a collection of objects usually exhibit only partial similarity, as in the case of images of the same scene but from different camera positions. These practical scenarios require consistent matching of multiple objects that are only partially similar to each other.
3. **Incomplete Input Maps:** Computing pairwise maps across all object pairs are often expensive, sometimes inadmissible, and in fact unnecessary. Depending on the characteristics of input sources, one might be able to infer unobserved maps from a small sample of noisy pairwise matches. While [12] considered incomplete inputs, the tradeoff between the undersampling factor and the error-correction ability remains unknown.

All in all, practical applications require matching partially similar objects from a small fraction of densely corrupted pairwise maps — a goal this paper aims to achieve.

1.1 Contributions

This paper is concerned with joint object matching under dense input errors. Our main contributions in this regard are three-fold.

1. **Algorithms:** Inspired by the recent evidence on the power of convex relaxation, we propose to solve the joint matching problem via a semidefinite program called MatchLift. The algorithm relaxes the binary-value constraints, and attempts to maximize the compatibility between the input and the recovered maps. The program is established upon a semidefinite conic constraint that relies on the total number m of distinct elements to be matched. To this end, we propose to pre-estimate m via a spectral method. Our methodology is essentially *parameter free*, and can be solved by scalable optimization algorithms.
2. **Theory:** We derive performance guarantees for exact matching. Somewhat surprisingly, MatchLift admits perfect map recovery even in the presence of dense input corruptions. Our findings reveal the near-optimal error-correction ability of MatchLift, i.e. as n grows, the algorithm is guaranteed to work even when a dominant fraction – more precisely, a fraction $1 - \Omega\left(\frac{\log^2 n}{\sqrt{n}}\right)$ – of the inputs behave as random outliers. Besides, while the presence of partial similarity unavoidably incurs more severe types of input errors, MatchLift exhibits a strong recovery ability nearly order-wise equivalent to that in the full-similarity scenario, as long as the fraction of each object being disclosed is bounded away from zero. Finally, in many situations, MatchLift succeeds even with minimal input complexity, in the sense that it can reliably fill in all unobserved maps based on very few noisy partial inputs, as soon as the provided maps form a connected graph. This is information theoretically optimal.

3. **Practice:** We have evaluated the performance of MatchLift on several benchmark datasets. These datasets include several synthetic examples as well as real examples from several popular benchmarks. Experimental results on synthetic examples corroborate our theoretical findings. On real datasets, the quality of the maps generated by MatchLift outperforms the state-of-the-art object matching and graph clustering algorithms.

1.2 Prior Art

There has been numerous work studying the problem of object matching, either in terms of shape mapping, graph matching, or image mapping, which is impossible to enumerate. We list below a small sample of development on joint object matching, as well as its relation and distinction to the well-renowned graph clustering problem.

- **Object Matching.** Early work on object matching focused primarily on matching pairs of objects in isolation (e.g. [13–15]). Due to the limited and biased information present in an isolated object pair, pairwise matching techniques can easily, sometimes unavoidably, generate false correspondences. Last few years have witnessed a flurry of activity in joint object matching, e.g. [9–12], which exploited the global cycle-consistency criterion to prune noisy maps. The fundamental understanding has recently been advanced by [12]. Nevertheless, none of the prior work have demonstrated provable recovery ability when the majority of input maps/correspondences are outliers, nor were they able to accommodate practical scenarios where different objects only exhibit partial similarity. Recent work [16] employed spectral methods for denoising in the full-similarity case. However, the errors considered therein are modeled as Gaussian-Wigner additive noise, which is not applicable in our setting. Another line of work [17, 18] proposed to recover global rigid transform between points via convex relaxation, where the point coordinates might only be partially observed. While this line of work is relevant, the problem considered therein is more specialized than the point-based joint matching studied in this paper; also, none of these paradigms are able to enable dense error correction.
- **Matrix Completion and Robust PCA.** In a broader sense, our approach is inspired by the pioneering work in low-rank matrix completion [19, 20] and robust principal component analysis [21–25], which reveal the power of convex relaxation in recovering low-dimensional structures among high-dimensional objects. In fact, the ground truth herein is equivalent to a block-constant low-rank matrix [26], as occurred in various graph-related problems. Nevertheless, their theoretical analyses fail to provide tight bounds in our setting, as the low-rank matrix relevant in our cases is highly sparse as well. That said, additional structural assumptions need to be incorporated in order to achieve optimal performance.
- **Graph Clustering.** The joint matching problem can be treated as a structured graph clustering (GC) problem, where graph nodes represent points on objects and the edge set encodes all correspondences. In this regard, any GC algorithm [27–32] provides a heuristic to estimate graph matching. Nevertheless, there are several intrinsic structural properties herein that are not explored by any generic GC approaches. First, our input takes a block-matrix form, where each block is highly *structured* (i.e. doubly-substochastic), *sparse*, and inter-dependent. Second, the points belonging to the same object are *mutually exclusive* to each other. Third, the corruption rate for different entries can be highly non-symmetric – when translated into GC languages, this means that in-cluster edges might suffer from an order-of-magnitude larger error rate than inter-cluster edges. As a result, the findings for generic GC methods do not deliver encouraging guarantees when applied to our setting. Detailed theoretical and empirical comparisons are provided in Sections 4 and 5, respectively.

1.3 Organization

The rest of the paper is organized as follows. Section 2 formally presents the problem setup, including the input model and the expected output. Our two-step recovery procedure – a spectral method followed by a convex program called MatchLift – is described in Section 3. A scalable alternating direction method of multipliers (ADMM) together with a greedy rounding strategy is also introduced in Section 3. Section 4 presents the main theoretical performance guarantees for our method under a natural randomized model.

All proofs of the main theorems are deferred to the appendices. We introduce numerical experiments demonstrating the practicability of our method in Section 5, as well as empirical comparison with other best-known algorithms. Finally, Section 6 concludes the paper with a summary of our findings.

2 Problem Formulation and Preliminaries

This section presents the problem setup for matching multiple partially similar objects, and introduces an algebraic form for representing a collection of pairwise maps.

2.1 Terminology

Below we formally define several important notions that will be used throughout this paper.

- **Set.** We represent objects to be matched as discrete sets. For example, these sets can represent the vertex sets in the graph matching problem, or encode feature points when matching images.
- **Partial Map.** Given two discrete sets \mathcal{S} and \mathcal{S}' , a subset $\phi \subset \mathcal{S} \times \mathcal{S}'$ is termed a partial map if each element of \mathcal{S} (resp. \mathcal{S}') is paired with *at most one* element of \mathcal{S}' (resp. \mathcal{S}) — in particular, not all elements need to be paired.
- **Map Graph.** A graph $\mathcal{G} = (\mathcal{V}, \mathcal{E})$ is called a map graph w.r.t. n sets $\mathcal{S}_1, \dots, \mathcal{S}_n$ if (i) $\mathcal{V} := \{\mathcal{S}_1, \dots, \mathcal{S}_n\}$, and (ii) $(\mathcal{S}_i, \mathcal{S}_j) \in \mathcal{E}$ implies that pairwise estimates on the partial maps ϕ_{ij} and ϕ_{ji} between \mathcal{S}_i and \mathcal{S}_j are available.

2.2 Input and Output

The input and expected output for the joint object matching problem are described as follows.

- **Input (Noisy Pairwise Maps).** Given n sets $\mathcal{S}_1, \dots, \mathcal{S}_n$ with respective cardinality m_1, \dots, m_n and a (possibly sparse) map graph \mathcal{G} , the input to the recovery algorithm consists of partial maps $\phi_{ij}^{\text{in}} ((i, j) \in \mathcal{G})$ between \mathcal{S}_i and \mathcal{S}_j estimated in isolation, using any off-the-shelf *pairwise* matching method. Note that the input maps ϕ_{ij}^{in} one obtain might not agree, partially or totally, with the ground truth.
- **Output (Consistent Global Matching).** The main objective of this paper is to detect and prune incorrect pairwise input maps in an efficient and reliable manner. Specifically, we aim at proposing a tractable algorithm that returns a full collection of partial maps $\{\phi_{ij} \mid 1 \leq i, j \leq n\}$ that are (i) globally consistent, and (ii) close to the provided pairwise maps – and under some conditions provably the ground-truth maps.

As will be detailed later, the key idea of our approach is to explore global consistency across all pairwise maps. In fact, points across different objects must form several clusters, and the ground-truth maps only exhibit in-cluster edges. We will introduce a novel convex relaxation tailored to the structure of the input maps (Section 3) and investigate its theoretical performance (Section 4).

2.3 Joint Matching in Matrix Form

In the same spirit as most convex relaxation techniques (e.g., [12, 30]), we use matrices to encode maps between objects. Specifically, we encode a partial map $\phi_{ij} : \mathcal{S}_i \mapsto \mathcal{S}_j$ as a binary matrix $\mathbf{X}_{ij} \in \{0, 1\}^{|\mathcal{S}_i| \times |\mathcal{S}_j|}$ such that $\mathbf{X}_{ij}(s, s') = 1$ iff $(s, s') \in \phi_{ij}$. Valid partial map matrices \mathbf{X}_{ij} shall satisfy the following doubly sub-stochastic constraints:

$$\mathbf{0} \leq \mathbf{X}_{ij} \mathbf{1} \leq \mathbf{1}, \quad \mathbf{0} \leq \mathbf{X}_{ij}^\top \mathbf{1} \leq \mathbf{1}. \quad (1)$$

Symbol	Description
$\mathbf{1}$	ones vector: a vector with all entries one
\mathbf{X}_{ij}	(i, j) -th block of a block matrix \mathbf{X} .
$\langle \mathbf{A}, \mathbf{B} \rangle$	matrix inner product, i.e. $\langle \mathbf{A}, \mathbf{B} \rangle = \text{tr}(\mathbf{A}^\top \mathbf{B})$.
$\text{diag}(\mathbf{X})$	a column vector formed from the diagonal of a square matrix \mathbf{X}
$\text{Diag}(\mathbf{x})$	a diagonal matrix that puts \mathbf{x} on the main diagonal
\mathbf{e}_i	i th unit vector, whose i th component is 1 and all others 0
\otimes	tensor product, i.e. $\mathbf{A} \otimes \mathbf{B} = \begin{bmatrix} a_{1,1}\mathbf{B} & a_{1,2}\mathbf{B} & \cdots & a_{1,n_2}\mathbf{B} \\ a_{2,1}\mathbf{B} & a_{2,2}\mathbf{B} & \cdots & a_{2,n_2}\mathbf{B} \\ \vdots & \vdots & \vdots & \vdots \\ a_{n_1,1}\mathbf{B} & a_{n_1,2}\mathbf{B} & \cdots & a_{n_1,n_2}\mathbf{B} \end{bmatrix}$
$\Omega_{\text{gt}}, \Omega_{\text{gt}}^\perp$	support of \mathbf{X}^{gt} , its complement support
$T_{\text{gt}}, T_{\text{gt}}^\perp$	tangent space at \mathbf{X}^{gt} , its orthogonal complement
$\mathcal{P}_{\Omega_{\text{gt}}}, \mathcal{P}_{\Omega_{\text{gt}}^\perp}$	projection onto the space of matrices supported on Ω_{gt} and Ω_{gt}^\perp , respectively
$\mathcal{P}_{T_{\text{gt}}}, \mathcal{P}_{T_{\text{gt}}^\perp}$	projection onto T_{gt} and T_{gt}^\perp , respectively

Table 1: Summary of Notation and Parameters

We then use an $n \times n$ block matrix $\mathbf{X} \in \{0, 1\}^{N \times N}$ to encode the entire collection of partial maps $\{\phi_{ij} \mid 1 \leq i, j \leq n\}$ over $\{\mathcal{S}_1, \dots, \mathcal{S}_n\}$:

$$\mathbf{X} = \begin{pmatrix} \mathbf{I}_{m_1} & \mathbf{X}_{12} & \cdots & \mathbf{X}_{1n} \\ \mathbf{X}_{21} & \mathbf{I}_{m_2} & \cdots & \mathbf{X}_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{X}_{n1} & \cdots & \cdots & \mathbf{I}_{m_n} \end{pmatrix}, \quad (2)$$

where $m_i := |\mathcal{S}_i|$ and $N := \sum_{i=1}^n m_i$. Note that all diagonal blocks are identity matrices, as each object is isomorphic to itself.

For notational simplicity, we will use \mathbf{X}^{in} throughout to denote the collection of pairwise input maps, i.e. each obtained pairwise estimate ϕ_{ij}^{in} is encoded as a binary map matrix $\mathbf{X}_{ij}^{\text{in}} \in \{0, 1\}^{m_i \times m_j}$ obeying the constraint (1). Some other useful notation is summarized in Table 1.

3 Methodology

This section presents a novel methodology, based on a theoretically rigorous and numerically efficient framework.

3.1 MatchLift: A Novel Two-Step Algorithm

We start by discussing the consistency constraint on the underlying ground-truth maps. Assume that there exists a universe $\mathcal{S} = \{1, \dots, m\}$ of m elements such that i) each object \mathcal{S}_i is a (partial) image of \mathcal{S} ; ii) each element in \mathcal{S} is contained in at least one object \mathcal{S}_i . Then the ground-truth correspondences shall connect points across objects that are associated with the same element.

Formally speaking, let the binary matrix $\mathbf{Y}_i \in \{0, 1\}^{m_i \times m}$ encode the underlying correspondences between each point and the universe, i.e. for any $s_i \in \mathcal{S}_i$ and $s \in \mathcal{S}$,

$$\mathbf{Y}_i(s_i, s) = 1, \quad \text{iff } s_i \text{ corresponds to } s.$$

This way one can express

$$\mathbf{X} = \mathbf{Y}\mathbf{Y}^\top$$

with $\mathbf{Y} = (\mathbf{Y}_1^\top, \dots, \mathbf{Y}_n^\top)^\top$, which makes clear that

$$\text{rank}(\mathbf{X}) = m.$$

This is equivalent to the graph partitioning setting with m cliques. Consequently, a natural candidate is to seek a low-rank and positive semidefinite (PSD) matrix to approximate the input. However, this strategy does not effectively explore the sparsity structure underlying the map collection.

To obtain a more powerful formulation, the proposed algorithm is based on the observation that even under dense input corruption, we are often able to obtain reliable estimates on m – the universe size, using spectral techniques. This motivates us to incorporate the information of m into the formulation so as to develop tighter relaxation. Specifically, we lift \mathbf{X} with one more dimension and consider

$$\begin{bmatrix} m & \mathbf{1}^\top \\ \mathbf{1} & \mathbf{X} \end{bmatrix} = \begin{bmatrix} \mathbf{1}^\top \\ \mathbf{Y} \end{bmatrix} \begin{bmatrix} \mathbf{1} & \mathbf{Y}^\top \end{bmatrix} \succeq \mathbf{0}, \quad (3)$$

which is strictly tighter than merely imposing $\mathbf{X} \succeq \mathbf{0}$. Intuitively, the formulation (3) entitles us one extra degree of freedom to assist in outlier pruning, which turns out to be crucial in “debiasing” the errors. Encouragingly, this tightened constraint leads to remarkably improved theoretical guarantees, as will be shown in Section 4. In the following, we formally present our two-step matching procedure.

- **Step I: Estimating m .** We estimate m by tracking the spectrum of the input \mathbf{X}^{in} . According to common wisdom (e.g. [33]), a block-sparse matrix \mathbf{X}^{in} must first be trimmed in order to remove the undesired bias effect caused by over-represented rows / columns. One candidate trimming procedure is provided as follows.

- **Trimming Procedure.** Set d_{\min} to be the smallest vertex degree of \mathcal{G} , and we say a vertex is *over-represented* if its vertex degree in \mathcal{G} exceeds $2d_{\min}$. Then for each overrepresented vertex i , randomly sample $2d_{\min}$ edges incident to it and set to zero all blocks $\mathbf{X}_{ij}^{\text{in}}$ associated with the remaining edges.

With this trimming procedure, we propose to pre-estimate m via Algorithm 1.

Algorithm 1 Estimating the size m of the universe \mathcal{S}

- 1) trim \mathbf{X}^{in} , and let $\tilde{\mathbf{X}}^{\text{in}}$ be the output.
 - 2) perform eigenvalue decomposition on $\tilde{\mathbf{X}}^{\text{in}}$; denote by λ_i the i th largest eigenvalue.
 - 3) **output:** $\hat{m} := \arg \max_{M \leq i < N} |\lambda_i - \lambda_{i+1}|$, where $M = \max\{2, \max_{1 \leq i \leq n} m_i\}$.
-

In short, Algorithm 1 returns an estimate of m via spectral methods, which outputs the number of dominant principal components of \mathbf{X}^{in} .

- **Step II: Map Recovery.** Now that we have obtained an estimate on m , we are in position to present our optimization heuristic that exploits the structural property (3). In order to guarantee that the recovery is close to the provided maps ϕ_{ij}^{in} , one alternative is to maximize correspondence agreement (i.e. the number of compatible non-zero entries) between the input and output. This results in an objective function:

$$\sum_{(i,j) \in \mathcal{G}} \langle \mathbf{X}_{ij}^{\text{in}}, \mathbf{X}_{ij} \rangle.$$

Additionally, since a *non-negative* map matrix \mathbf{X} is inherently sparse, it is natural to add an ℓ_1 regularization term to encourage sparsity, which in our case reduces to

$$\langle \mathbf{1} \cdot \mathbf{1}^\top, \mathbf{X} \rangle.$$

Since searching over all 0-1 map matrices is intractable, we propose to relax the binary constraints. Putting these together leads to the following semidefinite program referred to as *MatchLift*:

$$\begin{aligned}
(\text{MatchLift}) \quad & \underset{\mathbf{X} \in \mathbb{R}^{N \times N}}{\text{maximize}} && \sum_{(i,j) \in \mathcal{G}} \langle \mathbf{X}_{ij}^{\text{in}}, \mathbf{X}_{ij} \rangle - \lambda \langle \mathbf{1} \cdot \mathbf{1}^\top, \mathbf{X} \rangle \\
& \text{subject to} && \mathbf{X}_{ii} = \mathbf{I}_{m_i}, \quad 1 \leq i \leq n, \\
& && \mathbf{X} \succeq \mathbf{0}, \\
& && \begin{bmatrix} m & \mathbf{1}^\top \\ \mathbf{1} & \mathbf{X} \end{bmatrix} \succeq \mathbf{0}.
\end{aligned} \tag{4}$$

Remark 1. Here, λ represents the regularization parameter that balances the compatibility to the input and the sparsity structure. As we will show, the recovery ability of MatchLift is not sensitive to the choice of λ . By default, one can set

$$\lambda = \frac{\sqrt{|\mathcal{E}|}}{2n}, \tag{5}$$

which results in a *parameter-free* formulation.

Remark 2. Careful readers will note that the set of doubly stochastic constraints (1) can be further added into the program. Nevertheless, while enforcement of these constraints (1) results in a strictly tighter relaxation, it only leads to marginal improvement when (4) is present. As a result, we remove them for the sake of computational efficiency. We note, however, that in the scenario where m is difficult to estimate, imposing (1) will “become crucial in allowing a constant fraction (e.g. 50%) of error rate, although dense error correction might not be guaranteed.

This algorithm, all at once, attempts to disentangle the ground truth and outliers as well as predict unobserved maps via convex relaxation, inspired by recent success in sparse and low-rank matrix decomposition [21, 22]. Since the ground truth matrix is *simultaneously* low-rank and sparse; existing methodologies, which focus on dense low-rank matrices, typically yield loose, uninformative bounds in our setting.

Finally, we note that our matching algorithm and main results are well suited for a broad class of scenarios where each pairwise input can be modeled as a (partial) permutation matrix. For instance, our setting subsumes phase correlation [34], angular synchronization [35], and multi-signal alignment [36] as special cases.

3.2 Alternating Direction Methods of Multipliers (ADMM)

Most advanced off-the-shelf SDP solvers like SeDuMi or MOSEK are typically based on interior point methods, and such second-order methods are unable to handle problems with large dimensionality. For practical applicability, we propose a first-order optimization algorithm for approximately solving MatchLift, which is a variant of the ADMM method for semidefinite programs presented in [37]. Theoretically it is guaranteed to converge. Empirically, it is often the case that ADMM converges to modest accuracy within a reasonable amount of time, and produces desired results with the assistance of appropriate rounding procedures. This feature makes ADMM practically appealing in our case since the ground-truth matrix is known to be a 0-1 matrix, for which moderate entry-wise precision is sufficient to ensure good rounding accuracy. The details of the ADMM algorithm are deferred to Appendix A.

3.3 Rounding Strategy

As MatchLift solves a relaxed program of the original convex problem, it may return fractional solutions. In this case, we propose a greedy rounding method to generate valid partial maps. Given the solution $\tilde{\mathbf{X}}$ to MatchLift, the proposed strategy proceeds as in Algorithm 2. One can verify that this simple deterministic rounding strategy returns a matrix that encodes a consistent collection of partial maps. Note that \mathbf{v}_i^T denotes the i th row of a matrix \mathbf{V} .

Algorithm 2 Rounding Strategy

initialize compute the top r eigenvalues $\Sigma = \text{diag}(\sigma_1, \dots, \sigma_r)$ and eigenvectors $U = (\mathbf{u}_1, \dots, \mathbf{u}_r)$ of \hat{X} , where r is an estimate of the total number distinctive points to be recovered. Form $V = U\Sigma^{\frac{1}{2}}$.

repeat

1) Let O be a unitary matrix that obeys $O\mathbf{v}_1 = \mathbf{e}_1$, and set $V \leftarrow VO^\top$.

2) For each of the remaining rows \mathbf{v}_i belonging to each set \mathcal{S}_j ($i \in \mathcal{S}_j$), perform

$$\mathbf{v}_i \leftarrow \mathbf{e}_1, \quad \text{if } \langle \mathbf{v}_i, \mathbf{v}_1 \rangle > 0.5 \text{ and } i = \arg \max_{l \in \mathcal{S}_j} \langle \mathbf{v}_l, \mathbf{v}_1 \rangle.$$

3) All indices i obeying $\mathbf{v}_i = \mathbf{e}_1$ are declared to be matched with each other, and are then removed.

Repeat 1) for the next row that has not been fixed.

until all the rows of V have been fixed.

4 Theoretical Guarantees: Exact Recovery

Our heuristic algorithm MatchLift recovers, under a natural randomized setting, the ground-truth maps even when only a vanishing portion of the input correspondences are correct. Furthermore, MatchLift succeeds with minimal input complexity, namely, the algorithm is guaranteed to work as soon as those input maps that coincide with the ground truth maps form a connected map graph.

4.1 Randomized Model

In the following, we present a natural randomized model, under which the feature of MatchLift is easiest to interpret. Specifically, consider a universe $[m] := \{1, 2, \dots, m\}$. The randomized setting consider herein is generated through the following procedure.

- For each set \mathcal{S}_i ($1 \leq i \leq n$), each point $s \in [m]$ is included in \mathcal{S}_i independently with probability p_{set} .
- Each $\mathbf{X}_{ij}^{\text{in}}$ is observed / computed independently with probability p_{obs} .
- Each observed $\mathbf{X}_{ij}^{\text{in}}$ coincides with the ground truth independently with probability $p_{\text{true}} = 1 - p_{\text{false}}$.
- Each *observed but incorrect* $\mathbf{X}_{ij}^{\text{in}}$ is independently drawn from a set of partial map matrices satisfying

$$\mathbb{E}\mathbf{X}_{ij}^{\text{in}} = \frac{1}{m} \mathbf{1} \cdot \mathbf{1}^\top, \quad \text{if } \mathbf{X}_{ij}^{\text{in}} \text{ is observed and corrupted.} \quad (6)$$

Remark 3. The above mean condition (6) holds, for example, when the augmented block (i.e. that obtained by enhancing \mathcal{S}_i and \mathcal{S}_j to have all m elements) is drawn from the entire set of permutation matrices or other symmetric groups uniformly at random. While we impose (6) primarily to simplify our presentation of the analysis, we remark that this assumption can be significantly relaxed without degrading the matching performance.

Remark 4. We also note that the outliers do not need to be generated in an i.i.d. fashion. Our main results hold as long as they are jointly independent and satisfy the mean condition (6).

4.2 Main Theorem: Near-Optimal Matching

We are now in position to state our main results, which provide theoretical performance guarantees for our algorithms.

Theorem 1 (Accurate Estimation of m). *Consider the above randomized model. There exists an absolute constant $c_1 > 0$ such that with probability exceeding $1 - \frac{1}{m^5 n^5}$, the estimate on m returned by Algorithm 1 is exact as long as*

$$p_{\text{true}} \geq \frac{c_1 \log^2(mn)}{\sqrt{np_{\text{obs}}p_{\text{set}}}}. \quad (7)$$

Proof. See Appendix B. □

Theorem 1 ensures that one can obtain perfect estimate on the universe size or, equivalently, the rank of the ground truth map matrix via spectral methods. With accurate information on m , MatchLift allows perfect matching from densely corrupted inputs, as revealed below.

Theorem 2 (Exact and Robust Matching). *Consider the randomized model described above. There exist universal constants $c_0, c_1, c_2 > 0$ such that for any*

$$c_1 \left(\frac{p_{\text{obs}}}{m} + \sqrt{\frac{p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}} \right) \leq \lambda \leq \frac{\sqrt{p_{\text{obs}} \log(mn)}}{p_{\text{set}}}, \quad (8)$$

if the non-corruption rate obeys

$$p_{\text{true}} > \frac{c_0 \log^2(mn)}{\sqrt{np_{\text{obs}} p_{\text{set}}^2}}, \quad (9)$$

then the solution to MatchLift is exact and unique with probability exceeding $1 - (mn)^{-3}$.

Proof. See Appendix C. □

Note that the performance is not sensitive to λ as it can be arbitrarily chosen between $\Theta(\sqrt{\frac{p_{\text{obs}}}{n}})$ and $\Theta(\sqrt{p_{\text{obs}}})$. The implications of Theorem 2 are summarized as follows.

1. **Near-Optimal Recovery under Dense Errors.** Under the randomized model, MatchLift succeeds in pruning all outliers and recovering the ground truth with high probability. Somewhat surprisingly, this is guaranteed to work even when the non-corrupted pairwise maps account for only a *vanishing* fraction of the inputs. As a result, MatchLift achieves near-optimal recovery performance in the sense that as the number n of objects grows, its outlier-tolerance rate can be arbitrarily close to 1. Equivalently speaking, in the asymptotic regime, almost all input maps – more precisely, a fraction

$$1 - \Omega\left(\frac{\log^2 n}{\sqrt{n}}\right) \quad (10)$$

of inputs – can be badly corrupted by random errors without degrading the matching accuracy. This in turn highlights the significance of joint object matching: no matter how noisy the input sources are, perfect matching can be obtained as long as sufficiently many instances are available.

To the best of our knowledge, none of the prior results can support perfect recovery with more than 50% corruptions, regardless of how large n can be. The only comparative performance is reported for the robust PCA setting, where semidefinite relaxation enables dense error correction [24, 25]. However, their condition cannot be satisfied in our case. Experimentally, applying RPCA on joint matching is unable to tolerate dense errors (see Section 5).

2. **Exact Matching of Partially Similar Objects.** The challenge for matching partially similar objects arises in that the overlapping ratio between each pair of objects is in the order of p_{set}^2 while the size of each object is in the order of p_{set} . As correct correspondences only come from overlapping regions, it is expected that with a fixed p_{false} , the matching ability degrades when p_{set} decreases, which coincides with the bound in (9). However, the order of fault-tolerance rate with n is independent of p_{set} as long as p_{set} is bounded away from 0.
3. **Minimal Input Complexity.** Suppose that p_{set} and p_{false} are both constants bounded away from 0 and 1, and that $m = n^{O(\text{poly } \log(n))}$. Condition (9) asserts that: the algorithm is able to separate outliers and fill in all missing maps reliably with no errors, as soon as the input complexity (i.e. the number of pairwise maps provided) is about the order of $n \text{poly } \log(n)$. Recall that the connectivity threshold for an Erdős–Renyi graph $\mathcal{G}(n, p_{\text{obs}})$ is $p_{\text{obs}} > \frac{\log n}{n}$ (see [38]). This implies that MatchLift allows exact recovery nearly as soon as the input complexity exceeds the information theoretic limits.

4.3 Comparison with Prior Approaches

Our exact recovery condition significantly outperforms the best-known performance guarantees, including various SDP heuristics for matching problems, as well as general graph clustering approaches when applied to object matching, detailed below.

- Semidefinite Programming:** The SDP formulation proposed by Wang and Singer [17] admits exact recovery in the *full-similarity* setting when $p_{\text{true}} > c_1$ for some absolute constant $c_1 \approx 50\%$ in the asymptotic regime. One might also attempt recovery by minimizing a weighted sum of nuclear norm and ℓ_1 norm as suggested in matrix completion [19] and robust PCA [21, 22]. In order to enable dense error correction, robust PCA requires the sparse components (which is $\mathbf{X}^{\text{in}} - \mathbf{X}^{\text{gt}}$ here with \mathbf{X}^{gt} denoting the ground truth) to exhibit random signs [24, 25]. This cannot be satisfied in our setting since the sign pattern of $\mathbf{X}^{\text{in}} - \mathbf{X}^{\text{gt}}$ is highly biased (i.e. all non-negative entries of $\mathbf{X}^{\text{in}} - \mathbf{X}^{\text{gt}}$ lying in the support of \mathbf{X}^{gt} have negative signs, while all non-negative entries of $\mathbf{X}^{\text{in}} - \mathbf{X}^{\text{gt}}$ outside the support of \mathbf{X}^{gt} have positive signs).
- Graph Clustering:** Various approaches for general graph clustering have been proposed with theoretical guarantees under different randomized settings [28, 29, 31]. These results typically operate under the assumption that in-cluster and inter-cluster correspondences are independently corrupted, which does not apply in our model. Due to the block structure input model, these two types of corruptions are highly correlated and usually experience order-of-magnitude difference in corruption rate (i.e. $(1 - p_{\text{true}}) \frac{m-1}{m}$ for in-cluster edges and $(1 - p_{\text{true}}) \frac{1}{m}$ for inter-cluster edges). To facilitate comparison, we evaluate the most recent deterministic guarantees obtained by [31]. The key metric D_{max} therein can be easily bounded by $D_{\text{max}} \geq 1 - p_{\text{true}}$ due to a significant degree of in-cluster edge errors. The recovery condition therein requires

$$D_{\text{max}} < \frac{1}{m+1}, \quad \Rightarrow \quad p_{\text{true}} > \frac{m}{m+1},$$

which does not deliver encouraging guarantees compared with $p_{\text{true}} > \Theta\left(\frac{\log^2 n}{\sqrt{n}}\right)$ achieved by MatchLift.

5 Experimental Evaluation

In this section, we evaluate the performance of MatchLift and compare it against [29] and other graph matching methods. We consider both synthetic examples, which are used to verify the exact recovery conditions described above, as well as popular benchmark datasets for evaluating the practicability on real-world images.

5.1 Synthetic Examples

We follow the randomized model described in Section 4 to generate synthetic examples. For simplicity, we only consider the full observation mode, which establishes input maps between all pairs of objects. In all examples, we fix the universe size such that it consists of $m = 16$ points. We then vary the remaining parameters, i.e., n , p_{set} and p_{false} , to assess the performance of an algorithm. We evaluate 31×36 sets of parameters for each scenario, where each parameter configuration is simulated by 10 Monte Carlo trials. The empirical success probability is reflected by the color of each cell. Blue denotes perfect recovery in all experiments, and red denotes failure for all trials.

Figure 1(a) illustrates the phase transition for $p_{\text{set}} = 0.6$, when the number of objects n and p_{false} vary. We can see that MatchLift is exact even when the majority of the input correspondences are incorrect (e.g., 75% when $n = 150$). This is consistent with the theoretical result that the lower bound on p_{true} for exact recovery is $O(\log^2 n / \sqrt{n})$.

Figure 1(c) shows the phase transition for $n = 100$, when p_{set} and p_{false} vary. We can see that MatchLift tolerates more noise when p_{set} is large. This is also consistent with the result that the error-correction ability improves with p_{set} .

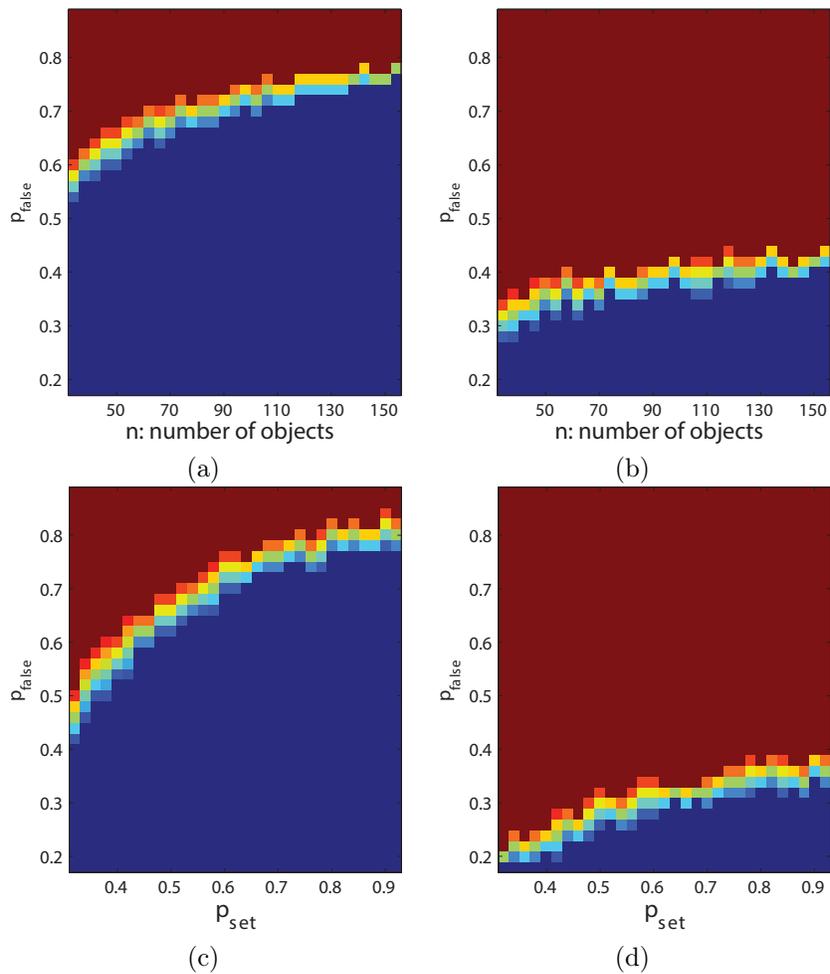


Figure 1: Phase Transition Diagrams of the proposed approach (MatchLift) and [29]. We can see that MatchLift can recover the ground-truth maps even the majority of the input correspondences are wrong, while the exact recovery of [29] requires that the percentage of incorrect correspondences is less than 50%. (a-b) $p_{\text{set}} = 0.6$. (c-d) $n = 100$.



Figure 2: A small benchmark called Building created for matching multiple images with partial similarity. Manually labeled feature points are highlighted.



Figure 3: A small benchmark called Chair created for matching multiple images with partial similarity. Manually labeled feature points are highlighted.

In comparison, Figure 1(b) and Figure 1(d) illustrate the phase transition diagrams achieved by the algorithm proposed in [29]. One can see that MatchLift is empirically superior, as [29] is unable to allow dense error correction in our case.

5.2 Real-World Examples

We have applied our algorithm on six benchmark datasets, i.e., CMU-House, CMU-Hotel, two datasets (Graf and Bikes) from [39]¹ and two new datasets (referred as Chair and Building, respectively) designed for evaluating joint partial object matching. As shown in Figures 2 and 3, the Building data set contains 16 images taken around a building [4], while the Chair data set contains 16 images of a chair model from different viewpoints. In the following, we first discuss the procedure for generating the input to our algorithm, i.e., the input sets and the initial maps. We then present the evaluation setup and analyze the results.

- **Feature points and initial maps.** To make fair comparisons with previous techniques on CMU-House and CMU-Hotel, we use the features points provided in [15] and apply the spectral matching

¹available online: robots.ox.ac.uk/vgg/research/affine

	Input	MatchLift	RPCA	LearnI	LearnII
House	68.2%	100%	92.2%	99.8%	96%
Hotel	64.1%	100%	90.1%	99.8%	90%

Table 2: Matching performance on the hotel and house datasets. We compare the proposed MatchLift algorithm with Robust PCA (RPCA) and two learning based graph matching methods: LearnI [45] and LearnII [15].

algorithm described in [40] to establish initial maps between features points. To assess the performance of the proposed algorithm with sparse input maps, we only match each image with 10 random neighboring images.

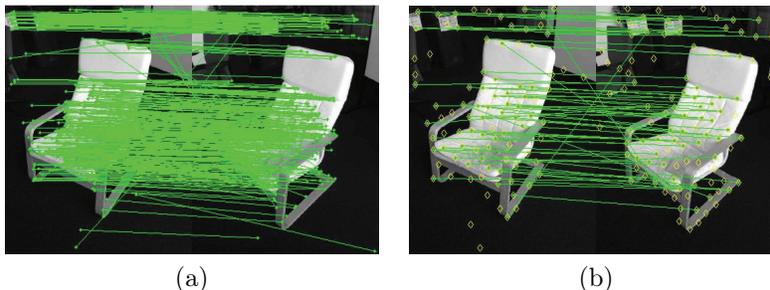


Figure 4: A map between dense SIFT feature points (a) is converted into a map between sampled feature points (b).

- To handle raw images in Chair, Building, Graf and Bikes, we apply a different strategy to build feature points and initial maps. We first detect dense SIFT feature points [41] on each image. We then apply RANSAC [42] to obtain correspondences between each pair of images. As SIFT feature points are over-complete, many of them do not appear in the resulting feature correspondences between pairs of views. Thus, we remove all feature points that have less than 2 appearances in all pair-wise maps. We further apply farthest point sampling on the feature points until the sampling density is above $0.05w$, where w is the width of the input images. The remaining feature points turn out to be much more distinct and thus are suitable for joint matching (See Figure 4). For the experiments we have tested, we obtain about 60 – 100 features points per image.
- **Evaluation protocol.** On CMU-House and CMU-Hotel, we count the percentage of correct feature correspondences produced by each algorithm. On Chair, Building, Graf and Bikes, we apply the metric described in [43], which evaluates the deviations of manual feature correspondences. As the feature points computed on each image do not necessarily align with the manual features, we apply [44] to interpolate feature level correspondences into pixel-wise correspondences for evaluation.
- **Results.** Table 2 shows the results of various algorithms on CMU-House and CMU-Hotel. We can see that even with moderate initial maps, MatchLift recovers all ground-truth correspondences. In contrast, the method of [29] can only recover 92.2% and 90.1% ground-truth correspondences on CMU-House and CMU-Hotel, respectively. Note that, MatchLift also outperforms state-of-the-art learning based graph matching algorithms [15, 45]. This shows the the advantage of joint object matching.

Figure 5 and Figure 6 illustrate the results of MatchLift on Chair, Building, Graf and Bikes. As these images contain noisy background information, the quality of the input maps is lower than those on House and Hotel. Encouragingly, MatchLift still recovers almost all manual correspondences. Moreover, MatchLift significantly outperforms [29], as the fault-tolerance rate of [29] is limited by a small constant barrier.

Another interesting observation is that the improvements on Graf and Bikes (each has 6 images) are lower than those on Chair and Building (each has 16 images). This is consistent with the common

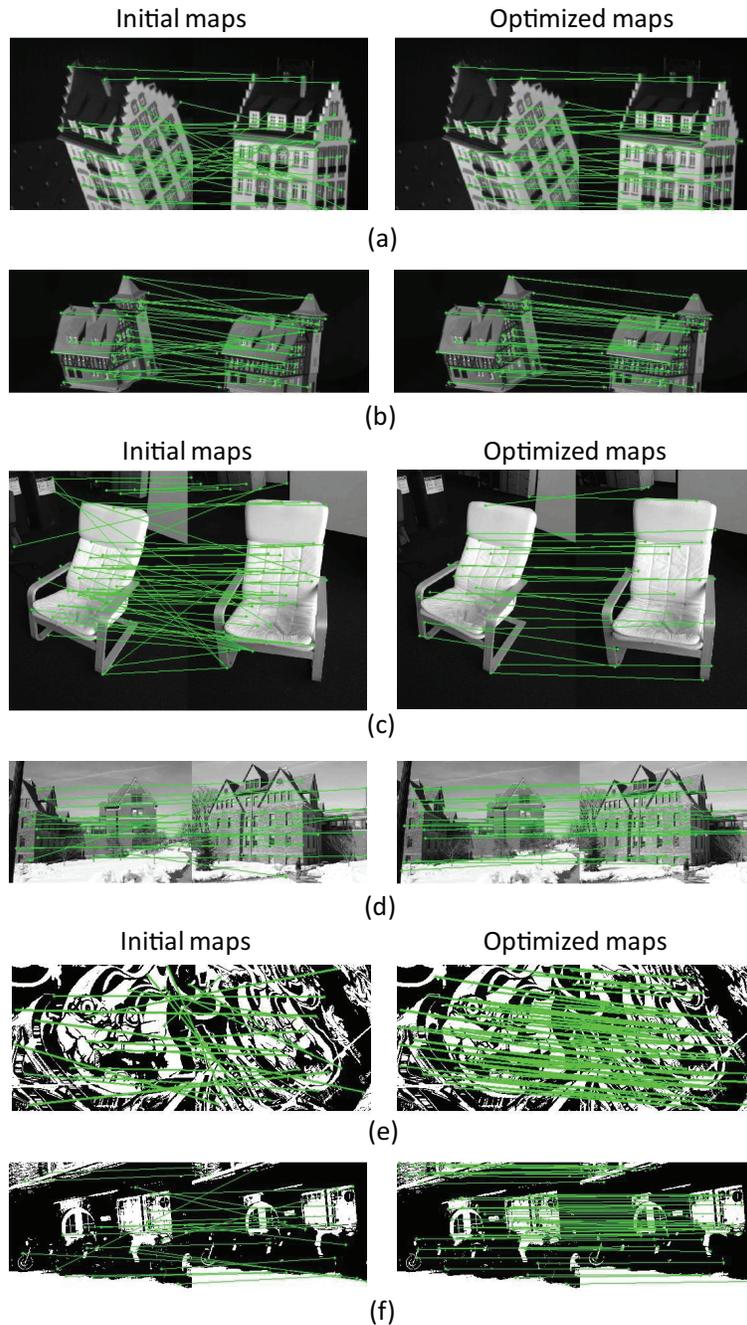


Figure 5: Comparisons between the input maps and the output of MatchLift on six benchmark datasets: (a) CMU Hotel, (b) CMU House, (c) Chair, (d) Building, (e) Graf, and (f) Bikes. The optimized maps not correct incorrect correspondences as well as fill in missing correspondences (generated by paths through intermediate shapes). One representative pair from each dataset is shown here.

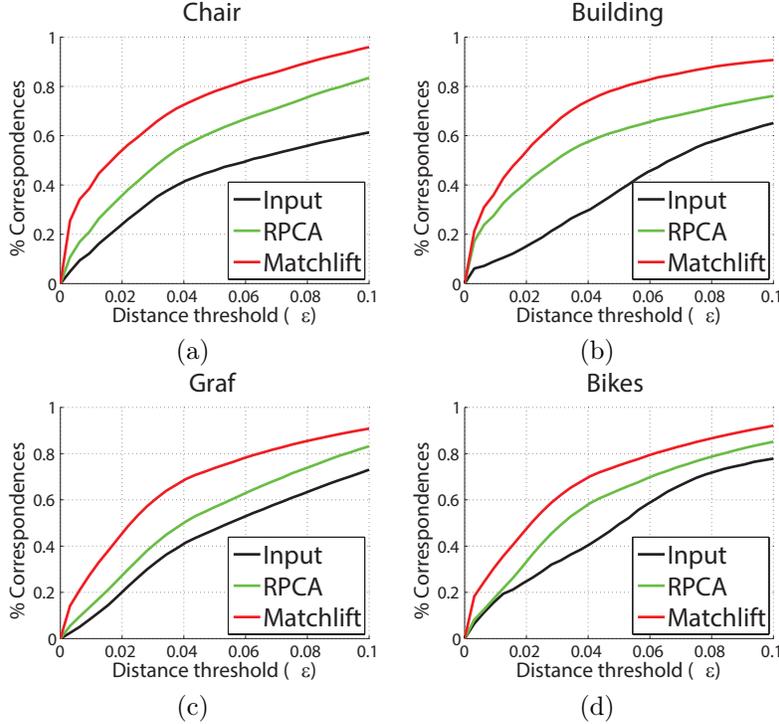


Figure 6: Percentages of ground-truth correspondences, whose distances to a map collection are below a varying threshold ϵ .

knowledge of data-driven effect, where large object collections possess stronger self-correction power than small object collections.

6 Conclusions

This paper delivers some encouraging news: given a few noisy object matches computed in isolation, a collection of partially similar objects can be accurately matched via semidefinite relaxation – an approach which provably works under dense errors. The proposed algorithm is essentially parameter-free, and can be solved by ADMM achieving remarkable efficiency and accuracy, with the assistance of a greedy rounding strategy.

The proposed algorithm achieves near-optimal error-correction ability, as it is guaranteed to work even when a dominant fraction of inputs are corrupted. This in turn underscore the importance of joint object matching: however low the quality of input sources is, perfect matching is achievable as long as we obtain sufficiently many instances. Also, while partial matching may incur much more severe input errors than those occurring in full-similarity matching, in many situations, the recovery ability of our algorithm is nearly the same as that in the full-similarity case (up to some constant factor). In a broader sense, our findings suggest that a large class of combinatorial / integer programming problems might be solved perfectly by semidefinite relaxation.

A Alternating Direction Method of Multipliers (ADMM)

This section presents the procedure for the ADMM algorithm. For notational simplicity, we represent the convex program as follows:

$$\begin{array}{ll}
 \underset{\mathbf{X}}{\text{minimize}} & \langle \mathbf{W}, \overline{\mathbf{X}} \rangle & \text{dual variable} \\
 \text{subject to} & \mathcal{A}(\overline{\mathbf{X}}) = \mathbf{b}, & \mathbf{y}_{\mathcal{A}} \\
 & \overline{\mathbf{X}} \succeq \mathbf{0}, & \mathbf{Z} \succeq \mathbf{0} \\
 & \overline{\mathbf{X}} \succeq \mathbf{0}, & \mathbf{S} \succeq \mathbf{0}
 \end{array}$$

where we denote $\overline{\mathbf{X}} := \begin{bmatrix} m & \mathbf{1}^\top \\ \mathbf{1} & \mathbf{X} \end{bmatrix}$. The matrices and operators are defined as follows

- (i) \mathbf{W} encapsulate all block coefficient matrices \mathbf{W}_{ij} for all $(i, j) \in \mathcal{G}$;
- (ii) $\mathcal{A}(\overline{\mathbf{X}}) = \mathbf{b}$ represents the constraint that $\mathbf{X}_{ii} = \mathbf{I}_{m_i}$ ($1 \leq i \leq n$) and the constraint $\overline{\mathbf{X}} = \begin{bmatrix} m & \mathbf{1}^\top \\ \mathbf{1} & \mathbf{X} \end{bmatrix}$;
- (iii) The variables on the right hand, i.e., $\mathbf{y}_{\mathcal{A}}$, \mathbf{Z} and \mathbf{S} , represent dual variables associated with respective constraints.

The Lagrangian associated with the convex program can be given as follows

$$\begin{aligned}
 \mathcal{L} &= \langle \mathbf{W}, \overline{\mathbf{X}} \rangle + \langle \mathbf{y}_{\mathcal{A}}, \mathcal{A}(\overline{\mathbf{X}}) - \mathbf{b} \rangle - \langle \mathbf{Z}, \overline{\mathbf{X}} \rangle - \langle \mathbf{S}, \overline{\mathbf{X}} \rangle \\
 &= \langle \mathbf{W} + \mathcal{A}^*(\mathbf{y}_{\mathcal{A}}) - \mathbf{Z} - \mathbf{S}, \overline{\mathbf{X}} \rangle - \langle \mathbf{b}, \mathbf{y}_{\mathcal{A}} \rangle.
 \end{aligned}$$

where \mathcal{A}^* denotes the conjugate operator w.r.t. an operator \mathcal{A} . The augmented Lagrangian for the convex program can now be written as

$$\begin{aligned}
 \mathcal{L}_{1/\mu} &= \langle \mathbf{b}, \mathbf{y}_{\mathcal{A}} \rangle + \langle \mathbf{Z} + \mathbf{S} - \mathbf{W} - \mathcal{A}^*(\mathbf{y}_{\mathcal{A}}), \overline{\mathbf{X}} \rangle \\
 &\quad + \frac{1}{2\mu} \|\mathbf{Z} + \mathbf{S} - \mathbf{W} - \mathcal{A}^*(\mathbf{y}_{\mathcal{A}})\|_{\mathbb{F}}^2.
 \end{aligned}$$

Here, the linear terms above represent the negative standard Lagrangian, whereas the quadratic parts represent the augmenting terms. μ is the penalty parameter that balances the standard Lagrangian and the augmenting terms. The ADMM then proceeds by alternately optimizing each primal and dual variable with others fixed, which results in closed-form solution for each subproblem. Denote by superscript k the iteration number, then we can present the ADMM iterative update procedures as follows

$$\begin{aligned}
 \mathbf{y}_{\mathcal{A}}^{(k+1)} &= (\mathcal{A}\mathcal{A}^*)^{-1} \left\{ \mathcal{A} \left(-\mathbf{W} + \mathbf{S}^{(k)} + \mu \overline{\mathbf{X}}^{(k)} + \mathbf{Z}^{(k)} \right) - \mu \mathbf{b} \right\}, \\
 \mathbf{Z}^{(k+1)} &= \left(\mathbf{W} + \mathcal{A}^* \left(\mathbf{y}_{\mathcal{A}}^{(k+1)} \right) - \mathbf{S}^{(k)} - \mu \overline{\mathbf{X}}^{(k)} \right)_+, \\
 \mathbf{S}^{(k+1)} &= \mathcal{P}_{\text{psd}} \left(\mathbf{W} + \mathcal{A}^* \left(\mathbf{y}_{\mathcal{A}}^{(k+1)} \right) - \mathbf{Z}^{(k+1)} - \mu \overline{\mathbf{X}}^{(k)} \right), \tag{11}
 \end{aligned}$$

$$\overline{\mathbf{X}}^{(k+1)} = \overline{\mathbf{X}}^{(k)} + \frac{1}{\mu} \left(\mathbf{Z}^{(k+1)} + \mathbf{S}^{(k+1)} - \mathbf{W} - \mathcal{A}^* \left(\mathbf{y}_{\mathcal{A}}^{(k+1)} \right) \right) \tag{12}$$

$$= -\frac{1}{\mu} \mathcal{P}_{\text{nsd}} \left(\mathbf{W} + \mathcal{A}^* \left(\mathbf{y}_{\mathcal{A}}^{(k+1)} \right) - \mathbf{Z}^{(k+1)} - \mu \overline{\mathbf{X}}^{(k)} \right). \tag{13}$$

Here, the operator \mathcal{P}_{psd} (resp. \mathcal{P}_{nsd}) denotes the projection onto the positive (resp. negative) semidefinite cone, and $(\cdot)_+$ operator projects all entries of a vector / matrix to non-negative values. Within a reasonable amount of time, ADMM typically returns moderately acceptable results.

B Proof of Theorem 1

The key step to the proof of Theorem 1 is to show that the set of outliers, even when they account for a dominant portion of the input matrix, behave only as a small perturbation to the spectrum of the non-corrupted components. Under the randomized model described in Section 4.1, it can be easily seen that

the trimming procedure is not invoked with high probability. Consequently, Theorem 1 can be established through the following lemma.

Lemma 1. *Given any set of n permutation matrices $\mathbf{P}_i \in \mathbb{R}^{m \times m}$ ($1 \leq i \leq n$), generate a random matrix \mathbf{M} via the following procedure.*

1. *Generate a symmetric block matrix $\mathbf{A} = [\mathbf{A}_{ij}]_{1 \leq i, j \leq n}$ such that*

$$\mathbf{A}_{ii} = \mathbf{I}, \quad 1 \leq i \leq n$$

and for all $i < j$,

$$\mathbf{A}_{ij} = \begin{cases} \mathbf{0}, & \text{if } \mu_{ij} = 0, \\ \mathbf{P}_i \mathbf{P}_j^\top, & \text{if } \nu_{ij} = 1 \text{ and } \mu_{ij} = 1, \\ \mathbf{U}_{ij}, & \text{else,} \end{cases} \quad (14)$$

where $\nu_{ij} \sim \text{Bernoulli}(p)$ and $\mu_{ij} \sim \text{Bernoulli}(\tau)$ are independent binary variables, and $\mathbf{U}_{ij} \in \mathbb{R}^{m \times m}$ are independent random permutation matrices obeying $\mathbb{E} \mathbf{U}_{ij} = \frac{1}{m} \mathbf{1}_m \cdot \mathbf{1}_m^\top$.

2. *\mathbf{M} is a principal minor of \mathbf{A} from rows / columns at indices from a set $I \subseteq \{1, 2, \dots, mn\}$, where each $1 \leq i \leq mn$ is contained in I independently with probability q .*

Then there exist absolute constants $c_1, c_2 > 0$ such that if $p \geq c_1 \frac{\log^2(mn)}{q\sqrt{\tau n}}$, one has

$$\begin{cases} \lambda_i(\mathbf{M}) \geq \left(1 - \frac{1}{\log(mn)}\right) \tau p q n, & \text{if } 1 \leq i \leq m \\ \lambda_i(\mathbf{M}) \leq c_2 \sqrt{\tau n} \log(mn) < \frac{\tau p q n}{\log(mn)}, & \text{if } i > m \end{cases} \quad (15)$$

with probability exceeding $1 - \frac{1}{m^5 n^5}$. Here, $\lambda_i(\mathbf{M})$ represents the i th largest eigenvalue of \mathbf{M} .

Proof of Lemma 1. Without loss of generality, we assume that $\mathbf{P}_i = \mathbf{I}_m$ for all $1 \leq i \leq n$, since rearranging rows / columns of \mathbf{A} does not change its eigenvalues. For convenience of presentation, we write $\mathbf{A} = \mathbf{Y} + \mathbf{Z}$ such that

$$\mathbf{Y}_{ii} = \tau \left(\frac{(1-p)}{m} \mathbf{1}_m \cdot \mathbf{1}_m^\top + p \mathbf{I}_m \right), \quad 1 \leq i \leq n$$

and for all $1 \leq i \leq j \leq n$:

$$\mathbf{Y}_{ij} = \begin{cases} \mathbf{0}, & \text{if } \mu_{ij} = 0, \\ \mathbf{I}_m, & \text{if } \nu_{ij} = 1 \text{ and } \mu_{ij} = 1, \\ \mathbf{U}_{ij}, & \text{else.} \end{cases} \quad (16)$$

This means that

$$\mathbf{Z}_{ij} = \begin{cases} \mathbf{I}_m - \mathbf{Y}_{ii}, & \text{if } i = j, \\ \mathbf{0}, & \text{else.} \end{cases} \quad (17)$$

Apparently, \mathbf{Z} is a block diagonal matrix satisfying

$$\|\mathbf{Z}\| \leq 2, \quad (18)$$

which is only a mild perturbation of \mathbf{Y} . This way we have reduced to the case where all blocks (including diagonal blocks) are i.i.d., which is slightly more convenient to analyze.

Decompose \mathbf{Y} into 2 components $\mathbf{Y} = \mathbf{Y}^{\text{mean}} + \mathbf{Y}^{\text{var}}$ such that

$$\forall 1 \leq i \leq j \leq n: \quad \mathbf{Y}_{ij}^{\text{mean}} = \tau \left(\frac{(1-p)}{m} \mathbf{1}_m \cdot \mathbf{1}_m^\top + p \mathbf{I}_m \right), \quad (19)$$

$$\forall 1 \leq i \leq n: \quad \mathbf{Y}_{ii}^{\text{var}} = \mathbf{0}, \quad (20)$$

and

$$\forall 1 \leq i < j \leq n: \quad \mathbf{Y}_{ij}^{\text{var}} = \begin{cases} -\tau \left(\frac{(1-p)}{m} \mathbf{1}_m \cdot \mathbf{1}_m^\top + p \mathbf{I}_m \right), & \text{if } \mu_{ij} = 0, \\ (1 - \tau p) \mathbf{I}_m - \frac{(1-p)}{m} \mathbf{1}_m \cdot \mathbf{1}_m^\top, & \text{if } \nu_{ij} = 1 \text{ and } \mu_{ij} = 1, \\ \mathbf{U}_{ij} - \tau \left(\frac{(1-p)}{m} \mathbf{1}_m \cdot \mathbf{1}_m^\top + p \mathbf{I}_m \right), & \text{else.} \end{cases} \quad (21)$$

In other words, \mathbf{Y}^{mean} represents the mean component of \mathbf{Y} , while \mathbf{Y}^{var} comprises all variations around the mean component. It is straightforward to check that

$$\mathbf{Y}^{\text{mean}} \succeq \mathbf{0}, \quad \text{rank}(\mathbf{Y}^{\text{mean}}) \leq m + 1.$$

If we denote by $\mathbf{Y}_I^{\text{mean}}$ the principal minor coming from the rows and columns of \mathbf{Y} at indices from I , then from Weyl's inequality one can easily see that

$$\lambda_i(\mathbf{M}) \geq \lambda_i(\mathbf{Y}_I^{\text{mean}}) - \|\mathbf{Y}^{\text{var}}\| - \|\mathbf{Z}\| \geq \lambda_i(\mathbf{Y}_I^{\text{mean}}) - \|\mathbf{Y}^{\text{var}}\| - 2, \quad 1 \leq i \leq m \quad (22)$$

and

$$\lambda_i(\mathbf{M}) \leq \lambda_i(\mathbf{Y}_I^{\text{mean}}) + \|\mathbf{Y}^{\text{var}}\| + \|\mathbf{Z}\| \leq \lambda_i(\mathbf{Y}_I^{\text{mean}}) + \|\mathbf{Y}^{\text{var}}\| + 2, \quad i > m. \quad (23)$$

In light of this, it suffices to evaluate $\|\mathbf{Y}^{\text{var}}\|$ as well as the eigenvalues of $\mathbf{Y}_I^{\text{mean}}$.

We are now in position to quantify the eigenvalues of $\mathbf{Y}_I^{\text{mean}}$. Without affecting its eigenvalue distribution, one can rearrange the rows / columns of $\mathbf{Y}_I^{\text{mean}}$ so that

$$\mathbf{Y}_I^{\text{mean}} \stackrel{\text{(permutation)}}{=} \tau p \begin{bmatrix} \mathbf{1}_{n_1} \cdot \mathbf{1}_{n_1}^\top & & \\ & \ddots & \\ & & \mathbf{1}_{n_m} \cdot \mathbf{1}_{n_m}^\top \end{bmatrix} + \frac{\tau(1-p)}{m} \mathbf{1}_N \cdot \mathbf{1}_N^\top. \quad (24)$$

Here, n_i ($1 \leq i \leq m$) denotes the cardinality of a set I_i generated by independently sampling n elements each with probability q , and we set $N := n_1 + \dots + n_m$ for simplicity. From Bernstein inequality, there exist universal constants $c_5, c_6 > 0$ such that if $q > \frac{c_5 \log(mn)}{n}$, then

$$|n_i - nq| \leq c_6 \sqrt{nq \log(mn)}, \quad 1 \leq i \leq m \quad (25)$$

holds with probability exceeding $1 - (mn)^{-10}$.

Since $\mathbf{Y}_I^{\text{mean}}$ is positive semidefinite, from (24) one can easily check that all non-zero eigenvalues of $\mathbf{Y}_I^{\text{mean}}$ are also eigenvalues of the following $(m+1) \times (m+1)$ matrix

$$\bar{\mathbf{Y}}_I^{\text{mean}} := \tau \begin{bmatrix} \sqrt{p} \mathbf{1}_{n_1}^\top & & & \\ & \sqrt{p} \mathbf{1}_{n_2}^\top & & \\ & & \ddots & \\ & & & \sqrt{p} \mathbf{1}_{n_m}^\top \\ & & & & \sqrt{\frac{1-p}{m}} \mathbf{1}_N^\top \end{bmatrix} \begin{bmatrix} \sqrt{p} \mathbf{1}_{n_1} & & & & \\ & \sqrt{p} \mathbf{1}_{n_2} & & & \\ & & \ddots & & \\ & & & \sqrt{p} \mathbf{1}_{n_m} & \\ & & & & \sqrt{\frac{1-p}{m}} \mathbf{1}_N \end{bmatrix} \quad (26)$$

$$= \tau \begin{bmatrix} pn_1 & & & & \\ & pn_2 & & & \\ & & \ddots & & \\ & & & pn_m & \\ \sqrt{\frac{p(1-p)}{m}} n_1 & \sqrt{\frac{p(1-p)}{m}} n_2 & \dots & \sqrt{\frac{p(1-p)}{m}} n_m & \frac{1-p}{m} N \end{bmatrix} \begin{bmatrix} \sqrt{\frac{p(1-p)}{m}} n_1 \\ \sqrt{\frac{p(1-p)}{m}} n_2 \\ \vdots \\ \sqrt{\frac{p(1-p)}{m}} n_m \\ \frac{1-p}{m} N \end{bmatrix} \quad (27)$$

$$= \tau q n \underbrace{\begin{bmatrix} p & & & \sqrt{\frac{p(1-p)}{m}} \\ & \ddots & & \vdots \\ & & p & \sqrt{\frac{p(1-p)}{m}} \\ \sqrt{\frac{p(1-p)}{m}} & \dots & \sqrt{\frac{p(1-p)}{m}} & 1-p \end{bmatrix}}_{\bar{\mathbf{Y}}_{I,0}} + \tau \underbrace{\begin{bmatrix} p \Delta_1 & & & \sqrt{\frac{p(1-p)}{m}} \Delta_1 \\ & \ddots & & \vdots \\ & & p \Delta_m & \sqrt{\frac{p(1-p)}{m}} \Delta_m \\ \sqrt{\frac{p(1-p)}{m}} \Delta_1 & \dots & \sqrt{\frac{p(1-p)}{m}} \Delta_m & \frac{1-p}{m} \Delta_N \end{bmatrix}}_{\bar{\mathbf{Y}}_{I,\Delta}}, \quad (27)$$

where

$$\Delta_i = n_i - nq \quad 1 \leq i \leq m,$$

and

$$\Delta_N = N - qnm,$$

which satisfies $|\Delta_N| \leq m \max_{1 \leq i \leq m} |\Delta_i|$.

By Schur complement condition for positive definite matrices [46], if $\begin{bmatrix} \mathbf{C} & \mathbf{B} \\ \mathbf{B}^\top & \mathbf{D} \end{bmatrix} \succ \mathbf{0}$, then $\mathbf{C} \succ \mathbf{0}$ and $\mathbf{D} - \mathbf{B}^\top \mathbf{C}^{-1} \mathbf{B} \succ \mathbf{0}$. Applying this condition to $\bar{\mathbf{Y}}_{I,0}$ suggests that $\bar{\mathbf{Y}}_{I,0} \succ \mathbf{0}$ can only hold when

$$(1-p) - \frac{p(1-p)}{m} \frac{1}{p} \mathbf{1}_m^\top \cdot \mathbf{1}_m > 0,$$

which however cannot be satisfied since $(1-p) - \frac{p(1-p)}{m} \frac{1}{p} \mathbf{1}_m^\top \cdot \mathbf{1}_m = 0$. Thus, $\bar{\mathbf{Y}}_{I,0}$ is rank deficient.

In fact, all non-zero eigenvalues of $\bar{\mathbf{Y}}_{I,0}$ can be quantified as well. Specifically, for any vector

$$\mathbf{z}_i := \mathbf{e}_i - \frac{1}{m} \begin{bmatrix} \mathbf{1}_m \\ 0 \end{bmatrix}, \quad 1 \leq i \leq m-1,$$

one can compute

$$\bar{\mathbf{Y}}_{I,0} \cdot \mathbf{z}_i = (\tau qpn) \mathbf{z}_i, \quad 1 \leq i \leq m-1. \quad (28)$$

That said, τqpn is an eigenvalue of $\bar{\mathbf{Y}}_{I,0}$ with multiplicity $m-1$. On the other hand, we have

$$\begin{cases} \bar{\mathbf{Y}}_{I,0} \cdot \begin{bmatrix} \mathbf{1}_m \\ \sqrt{\frac{(1-p)m}{p}} \end{bmatrix} = \tau qn \begin{bmatrix} \mathbf{1}_m \\ \sqrt{\frac{(1-p)m}{p}} \end{bmatrix}, \\ \mathbf{z}_i^\top \cdot \begin{bmatrix} \mathbf{1}_m \\ \sqrt{\frac{(1-p)m}{p}} \end{bmatrix} = 0, \end{cases} \quad (29)$$

indicating that τqn is another eigenvalue of $\bar{\mathbf{Y}}_{I,0}$. Putting these together yields

$$\lambda_i(\bar{\mathbf{Y}}_{I,0}) = \begin{cases} \tau qn, & i = 1 \\ \tau pqn, & 2 \leq i \leq m, \\ 0, & i > m. \end{cases} \quad (30)$$

Furthermore, the residual component $\bar{\mathbf{Y}}_{I,\Delta}$ can be bounded as follows

$$\begin{aligned} \|\bar{\mathbf{Y}}_{I,\Delta}\| &\leq \tau \left\| \begin{bmatrix} p\Delta_1 & & & \\ & \ddots & & \\ & & p\Delta_m & \\ & & & \frac{1-p}{m} \Delta_N \end{bmatrix} \right\| + \tau \left\| \begin{bmatrix} 0 & & & \sqrt{\frac{p(1-p)}{m}} \Delta_1 \\ & \ddots & & \vdots \\ & & 0 & \sqrt{\frac{p(1-p)}{m}} \Delta_m \\ \sqrt{\frac{p(1-p)}{m}} \Delta_1 & \dots & \sqrt{\frac{p(1-p)}{m}} \Delta_m & 0 \end{bmatrix} \right\|_{\mathbb{F}} \\ &\leq \tau \max \left\{ p \max_{1 \leq i \leq m} |\Delta_i|, \frac{1-p}{m} |\Delta_N| \right\} + \tau \sqrt{2p(1-p)} \max_{1 \leq i \leq m} |\Delta_i| \\ &\leq 2\tau \max_{1 \leq i \leq m} |\Delta_i| \leq 2c_6 \tau \sqrt{nq \log(mn)}, \end{aligned}$$

where the last inequality follows from (25). This taken collectively with (27) and (30) yields that: when $p > \frac{2c_6 \log^2(mn)}{\sqrt{nq}}$ or, equivalently, when $2c_6 \sqrt{nq \log(mn)} < \frac{1}{\log^{1.5}(mn)} npq$, one has

$$\begin{cases} \lambda_i(\mathbf{Y}_I^{\text{mean}}) \geq \left(1 - \frac{1}{\log^{\frac{3}{2}}(mn)}\right) \tau pqn, & 1 \leq i \leq m, \\ \lambda_i(\mathbf{Y}_I^{\text{mean}}) \leq 2c_6 \tau \sqrt{nq \log(mn)} \leq \frac{1}{\log^{\frac{3}{2}}(mn)} \tau pqn, & i > m. \end{cases} \quad (31)$$

Furthermore, observe that $\mathbb{E}\mathbf{Y}_{ij}^{\text{var}} = 0$, $\mathbb{E}\left\|\frac{1}{2\sqrt{\tau}}\mathbf{Y}_{ij}^{\text{var}}\right\|^2 \leq 1$, and $\frac{1}{2\sqrt{\tau}}\|\mathbf{Y}_{ij}^{\text{var}}\| \leq \frac{1}{\sqrt{\tau}}$. When $\tau > \frac{1}{n}$, Lemma 3 yields that

$$\|\mathbf{Y}^{\text{var}}\| \leq 2c_0\sqrt{\tau n} \log(mn) \quad (32)$$

with probability at least $1 - (mn)^{-5}$. Hence, $\|\mathbf{Y}^{\text{var}}\| = o(\tau pqn)$ if $p > \frac{c_{10} \log^2 n}{q\sqrt{\tau n}}$ for some constant $c_{10} > 0$.

Finally, the claim follows by substituting (31) and (32) into (22) and (23). \square

C Proof of Theorem 2

To prove Theorem 2, we first analyze the Karush–Kuhn–Tucker (KKT) condition for exact recovery, which provides a sufficient and almost necessary condition for uniqueness and optimality. Valid dual certificates are then constructed to guarantee exact recovery.

C.1 Preliminaries and Notations

Without loss of generality, we can treat \mathbf{X}^{gt} as a sub-matrix of an augmented square matrix $\mathbf{X}_{\text{sup}}^{\text{gt}}$ such that

$$\mathbf{X}_{\text{sup}}^{\text{gt}} := \mathbf{1} \cdot \mathbf{1}^\top \otimes \mathbf{I}_n, \quad (33)$$

and

$$\mathbf{X}^{\text{gt}} := \begin{bmatrix} \mathbf{\Pi}_1 & & & \\ & \mathbf{\Pi}_2 & & \\ & & \ddots & \\ & & & \mathbf{\Pi}_n \end{bmatrix} \mathbf{X}_{\text{sup}}^{\text{gt}} \begin{bmatrix} \mathbf{\Pi}_1^\top & & & \\ & \mathbf{\Pi}_2^\top & & \\ & & \ddots & \\ & & & \mathbf{\Pi}_n^\top \end{bmatrix}, \quad (34)$$

where the matrices $\mathbf{\Pi}_i \in \mathbb{R}^{|\mathcal{S}_i| \times m}$ are defined such that $\mathbf{\Pi}_i$ denotes the submatrix of \mathbf{I}_m coming from its rows at indices from \mathcal{S}_i . For instance, if $\mathcal{S}_i = \{2, 3\}$, then one has

$$\mathbf{\Pi}_i = \begin{bmatrix} 0 & 1 & 0 & \cdots & 0 \\ 0 & 0 & 1 & \cdots & 0 \end{bmatrix}.$$

With this notation, $\mathbf{\Pi}_i \mathbf{M} \mathbf{\Pi}_j^\top$ represents a submatrix of $\mathbf{M} \in \mathbb{R}^{m \times m}$ coming from the rows at indices from \mathcal{S}_i and columns at indices from \mathcal{S}_j . Conversely, for any matrix $\tilde{\mathbf{M}} \in \mathbb{R}^{|\mathcal{S}_i| \times |\mathcal{S}_j|}$, the matrix $\mathbf{\Pi}_i^\top \tilde{\mathbf{M}} \mathbf{\Pi}_j$ converts $\tilde{\mathbf{M}}$ to an $m \times m$ matrix space via zero padding.

With this notation, we can represent \mathbf{X}^{in} as a submatrix of $\mathbf{X}_{\text{sup}}^{\text{in}}$, which is a corrupted version of $\mathbf{X}_{\text{sup}}^{\text{gt}}$ and obeys

$$\mathbf{X}_{ij}^{\text{in}} := \mathbf{\Pi}_i (\mathbf{X}_{\text{sup}}^{\text{in}})_{ij} \mathbf{\Pi}_j^\top. \quad (35)$$

For notational simplicity, we set

$$\mathbf{W}_{ij} := \begin{cases} -\mathbf{X}_{ij}^{\text{in}} + \lambda \mathbf{1} \cdot \mathbf{1}^\top, & \text{if } (i, j) \in \mathcal{G}, \\ \lambda \mathbf{1} \cdot \mathbf{1}^\top, & \text{else.} \end{cases} \quad (36)$$

Before continuing to the proof, it is convenient to introduce some notations that will be used throughout. Denote by Ω_{gt} and Ω_{gt}^\perp the support of \mathbf{X}^{gt} and its complement support, respectively, and let $\mathcal{P}_{\Omega_{\text{gt}}}$ and $\mathcal{P}_{\Omega_{\text{gt}}^\perp}$ represent the orthogonal projection onto the linear space of matrices supported on Ω_{gt} and its complement support Ω_{gt}^\perp , respectively. Define T_{gt} to be the tangent space at \mathbf{X}^{gt} w.r.t. all symmetric matrices of rank at most m , i.e. the space of symmetric matrices of the form

$$T_{\text{gt}} := \left\{ \begin{bmatrix} \mathbf{\Pi}_1 \\ \mathbf{\Pi}_2 \\ \vdots \\ \mathbf{\Pi}_n \end{bmatrix} \mathbf{M} + \mathbf{M}^\top \begin{bmatrix} \mathbf{\Pi}_1^\top & \mathbf{\Pi}_2^\top & \cdots & \mathbf{\Pi}_n^\top \end{bmatrix} : \mathbf{M} \in \mathbb{R}^{m \times N} \right\}, \quad (37)$$

and denote by T_{gt}^\perp its orthogonal complement. We then denote by $\mathcal{P}_{T_{\text{gt}}}$ (resp. $\mathcal{P}_{T_{\text{gt}}^\perp}$) the orthogonal projection onto T_{gt} (resp. T_{gt}^\perp). In passing, if we define

$$\Sigma := \text{Diag} \left\{ \left[\frac{n}{n_1}, \dots, \frac{n}{n_m} \right] \right\}, \quad (38)$$

then the columns of

$$\mathbf{U} := \frac{1}{\sqrt{n}} \begin{bmatrix} \mathbf{\Pi}_1 \\ \mathbf{\Pi}_2 \\ \vdots \\ \mathbf{\Pi}_n \end{bmatrix} \Sigma^{\frac{1}{2}} \quad (39)$$

form the set of eigenvectors of \mathbf{X}^{gt} , and for any symmetric matrix \mathbf{M} ,

$$\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{M}) = (\mathbf{I} - \mathbf{U}\mathbf{U}^\top) \mathbf{M} (\mathbf{I} - \mathbf{U}\mathbf{U}^\top). \quad (40)$$

Furthermore, we define a vector \mathbf{d} to be

$$\mathbf{d} := \begin{bmatrix} \mathbf{\Pi}_1 \\ \mathbf{\Pi}_2 \\ \vdots \\ \mathbf{\Pi}_n \end{bmatrix} \Sigma \mathbf{1}_m. \quad (41)$$

Put another way, if any row index j of \mathbf{X}^{gt} is associated with the element $s \in [m]$, then $\mathbf{d}_j = \frac{n}{n_s}$. One can then easily verify that

$$\left\langle \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{X}^{\text{gt}} - \frac{1}{m} \mathbf{1} \cdot \mathbf{1}^\top \right\rangle = \left\langle \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{X}^{\text{gt}} \right\rangle - \frac{1}{m} (\mathbf{1}^\top \cdot \mathbf{d})^2 = 0. \quad (42)$$

In fact, when n_i 's are sufficiently close to each other, $\mathbf{d} \cdot \mathbf{d}^\top$ is a good approximation of $\mathbf{1} \cdot \mathbf{1}^\top$, as claimed in the following lemma.

Lemma 2. *Consider a set of Bernoulli random variables $\nu_i \sim \text{Bernoulli}(p)$ ($1 \leq i \leq n$), and set $s := \sum_{i=1}^n \nu_i$. Let n_i ($1 \leq i \leq m$) be independent copies of s , and denote $N = n_1 + \dots + n_m$. If $p > \frac{c_7 \log^2(mn)}{n}$, then the matrix*

$$\mathbf{A} := (np)^2 \begin{bmatrix} \frac{1}{n_1} \mathbf{1}_{n_1} \\ \frac{1}{n_2} \mathbf{1}_{n_2} \\ \vdots \\ \frac{1}{n_m} \mathbf{1}_{n_m} \end{bmatrix} \left[\frac{1}{n_1} \mathbf{1}_{n_1}^\top \quad \frac{1}{n_2} \mathbf{1}_{n_2}^\top \quad \dots \quad \frac{1}{n_m} \mathbf{1}_{n_m}^\top \right] \quad (43)$$

satisfies

$$\left\| \frac{1}{m} \mathbf{A} - \frac{1}{m} \mathbf{1}_N \cdot \mathbf{1}_N^\top \right\| \leq c_8 \sqrt{np \log(mn)} \quad (44)$$

and

$$\|\mathbf{A} - \mathbf{1}_N \cdot \mathbf{1}_N^\top\|_\infty \leq c_9 \sqrt{\frac{\log(mn)}{np}} \quad (45)$$

with probability exceeding $1 - \frac{1}{m^5 n^5}$, where c_7, c_8, c_9 are some universal constants.

Proof. See Appendix D.1. □

Since $p^2 \mathbf{d} \cdot \mathbf{d}^\top$ is equivalent to \mathbf{A} defined in (43) up to row / column permutation, Lemma 2 reveals that

$$\left\| \frac{p^2}{m} \mathbf{d} \cdot \mathbf{d}^\top - \frac{1}{m} \mathbf{1}_N \cdot \mathbf{1}_N^\top \right\| \leq c_8 \sqrt{np \log(mn)}$$

with high probability.

The following bound on the operator norm of a random block matrix is useful for deriving our main results.

Lemma 3. Let $\mathbf{M} = [\mathbf{M}_{ij}]_{1 \leq i, j \leq n}$ be a symmetric block matrix, where \mathbf{M}_{ij} 's are jointly independent $m_i \times m_j$ matrices satisfying

$$\mathbb{E}\mathbf{M}_{ij} = \mathbf{0}, \quad \mathbb{E}\|\mathbf{M}_{ij}\|^2 \leq 1, \quad \text{and} \quad \|\mathbf{M}_{ij}\| \leq \sqrt{n}, \quad (1 \leq i, j \leq n). \quad (46)$$

Besides, $m_i \leq m$ holds for all $1 \leq i \leq n$. Then there exists an absolute constant $c_0 > 0$ such that

$$\|\mathbf{M}\| \leq c_0 \sqrt{n} \log(mn)$$

holds with probability exceeding $1 - \frac{1}{m^5 n^5}$.

Proof. See Appendix D.2. □

Additionally, the second smallest eigenvalue of the Laplacian matrix of a random Erdős–Rényi graph can be bounded below by the following lemma.

Lemma 4. Consider an Erdős–Rényi graph $\mathcal{G} \sim \mathcal{G}(n, p)$ and any positive integer m , and let $\mathbf{L} \in \mathbb{R}^{n \times n}$ represent its (unnormalized) Laplacian matrix. There exist absolute constants $c_3, c_4 > 0$ such that if $p > c_3 \log^2(mn)/n$, then the algebraic connectivity $a(\mathcal{G})$ of \mathcal{G} (i.e. the second smallest eigenvalue of \mathbf{L}) satisfies

$$a(\mathcal{G}) \geq np - c_4 \sqrt{np} \log(mn) \quad (47)$$

with probability exceeding $1 - \frac{2}{(mn)^5}$.

Proof. See Appendix D.3. □

Finally, if we denote by n_s (resp. $n_{s,t}$) the number of sets \mathcal{S}_i ($1 \leq i \leq n$) containing the element s (resp. containing s and t simultaneously), then these quantities sharply concentrate around their mean values, as stated in the following lemma.

Lemma 5. There are some universal constants $c_8, c_9 > 0$ such that if $p_{\text{set}}^2 > \frac{\log(mn)}{n}$, then

$$\begin{aligned} |n_s - np_{\text{set}}| &\leq \sqrt{c_8 np_{\text{set}} \log(mn)}, \quad \forall 1 \leq s \leq m, \\ |n_{s,t} - np_{\text{set}}^2| &\leq \sqrt{c_8 np_{\text{set}}^2 \log(mn)}, \quad \forall 1 \leq s < t \leq m, \end{aligned}$$

hold with probability exceeding $1 - \frac{1}{(mn)^{10}}$.

Proof. In passing, the claim follows immediately from the Bernstein inequality that

$$\mathbb{P}\left(\left|\sum_{i=1}^n \nu_i - np\right| > t\right) \leq 2 \exp\left(-\frac{\frac{1}{2}t^2}{np(1-p) + \frac{1}{3}t}\right)$$

where $\nu_i \sim \text{Bernoulli}(p)$ are i.i.d. random variables. Interested readers are referred to [47] for a tutorial. □

C.2 Optimality and Uniqueness Condition

Recall that $n_i := |\mathcal{I}_i|$ denotes the number of sets \mathcal{S}_j containing the element i . The convex relaxation is exact if one can construct valid dual certificates, as summarized in the following lemma.

Lemma 6. Suppose that there exist dual certificates $\alpha > 0$, $\mathbf{Z} = [\mathbf{Z}_{ij}]_{1 \leq i, j \leq n} \in \mathbb{S}^{N \times N}$ and $\mathbf{Y} = [\mathbf{Y}_{ij}]_{1 \leq i, j \leq n} \in \mathbb{S}^{N \times N}$ obeying

$$\mathbf{Y} - \alpha \mathbf{d} \mathbf{d}^\top \succeq \mathbf{0}, \quad (48)$$

$$\mathcal{P}_{\Omega_{\text{gt}}}(\mathbf{Z}) = \mathbf{0}, \quad \mathcal{P}_{\Omega_{\text{gt}}^\perp}(\mathbf{Z}) \succeq \mathbf{0}, \quad (49)$$

$$\mathbf{Y}_{ij} = \mathbf{W}_{ij} - \mathbf{Z}_{ij}, \quad 1 \leq i < j \leq n, \quad (50)$$

$$\mathbf{Y} - \alpha \mathbf{d} \mathbf{d}^\top \in T_{\text{gt}}^\perp. \quad (51)$$

Then \mathbf{X}^{gt} is the unique solution to MatchLift if either of the following two conditions is satisfied:
i) All entries of \mathbf{Z}_{ij} ($\forall i \neq j$) within the support Ω_{gt}^\perp are strictly positive;
ii) For all \mathbf{M} satisfying $\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{M}) \succ \mathbf{0}$,

$$\langle \mathbf{Y} - \alpha \mathbf{d} \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{M}) \rangle > 0, \quad (52)$$

and, additionally,

$$\frac{n}{n_i} + \frac{n}{n_j} \neq \frac{n^2}{n_i n_j}, \quad 1 \leq i, j \leq m. \quad (53)$$

Proof. See Appendix D.4. \square

That said, to prove Theorem 2, it is sufficient (under the hypotheses of Theorem 2) to generate, with high probability, valid dual certificates \mathbf{Y} , \mathbf{Z} and $\alpha > 0$ obeying the optimality conditions of Lemma 6. This is the objective of the next subsection.

C.3 Construction of Dual Certificates

Decompose the input \mathbf{X}^{in} into two components $\mathbf{X}^{\text{in}} = \mathbf{X}^{\text{false}} + \mathbf{X}^{\text{true}}$, where

$$\mathbf{X}^{\text{true}} = \mathcal{P}_{\Omega_{\text{gt}}}(\mathbf{X}^{\text{in}}), \quad \text{and} \quad \mathbf{X}^{\text{false}} = \mathcal{P}_{\Omega_{\text{gt}}^\perp}(\mathbf{X}^{\text{in}}). \quad (54)$$

That said, \mathbf{X}^{true} (resp. $\mathbf{X}^{\text{false}}$) consists of all correct (resp. incorrect) correspondences (i.e. non-zero entries) encoded in \mathbf{X}^{in} . This allows us to write

$$\mathbf{W}_{ij} = \begin{cases} -\mathbf{X}_{ij}^{\text{false}} + \lambda \mathbf{E}_{ij} - \mathbf{X}_{ij}^{\text{true}} + \lambda \mathbf{E}_{ij}^\perp, & \text{if } (i, j) \in \mathcal{G}, \\ \lambda \mathbf{E}_{ij} + \lambda \mathbf{E}_{ij}^\perp, & \text{else,} \end{cases} \quad (55)$$

where \mathbf{E} and \mathbf{E}^\perp are defined to be

$$\mathbf{E} := \mathcal{P}_{\Omega_{\text{gt}}}(\mathbf{1} \cdot \mathbf{1}^\top), \quad \text{and} \quad \mathbf{E}^\perp := \mathbf{1} \cdot \mathbf{1}^\top - \mathbf{E}. \quad (56)$$

We propose constructing the dual certificate \mathbf{Y} by producing three symmetric matrix components $\mathbf{Y}^{\text{true},1}$, $\mathbf{Y}^{\text{true},2}$, and \mathbf{Y}^{L} separately, as follows.

1. *Construction of \mathbf{Z}^{m} and \mathbf{R}^{m} .* For any $\beta \geq 0$, define α_β to be

$$\alpha_\beta := \arg \min_{\alpha: \beta \mathbf{1} \cdot \mathbf{1}^\top - \alpha \mathbf{d} \cdot \mathbf{d}^\top \geq \mathbf{0}} \left\| \beta \mathbf{1} \cdot \mathbf{1}^\top - \alpha \mathbf{d} \cdot \mathbf{d}^\top \right\|_\infty. \quad (57)$$

By setting $\beta_0 := \lambda - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} - \sqrt{\frac{c_{10}p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}}$, we produce \mathbf{Z}^{m} and \mathbf{R}^{m} as follows

$$\mathbf{Z}^{\text{m}} = \mathcal{P}_{\Omega_{\text{gt}}^\perp} \left(\left(\lambda - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \right) \mathbf{1} \cdot \mathbf{1}^\top - \alpha_{\beta_0} \mathbf{d} \cdot \mathbf{d}^\top \right) \quad (58)$$

and

$$\mathbf{R}^{\text{m}} = \mathcal{P}_{\Omega_{\text{gt}}} \left(\left(\lambda - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \right) \mathbf{1} \cdot \mathbf{1}^\top - \alpha_{\beta_0} \mathbf{d} \cdot \mathbf{d}^\top \right) \quad (59)$$

for some sufficiently large constant $c_{10} > 0$.

2. *Construction of $\mathbf{Y}^{\text{true},1}$ and $\mathbf{Y}^{\text{true},2}$.* We set

$$\mathbf{Y}_{ij}^{\text{true},1} = \begin{cases} -\mathbf{X}_{ij}^{\text{true}} + \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{E}_{ij}, & \text{if } i < j, \\ \left(\sum_{j=1}^n \left(\mathbf{X}_{ij}^{\text{true}} - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{E}_{ij} \right) \right) \mathbf{\Pi}_j \mathbf{\Pi}_i^\top, & \text{if } i = j, \end{cases}$$

and

$$\mathbf{Y}_{ij}^{\text{true},2} = \begin{cases} \mathbf{R}_{ij}^{\text{m}}, & \text{if } i < j, \\ -\sum_{j=1}^n \mathbf{R}_{ij}^{\text{m}} \mathbf{\Pi}_j \mathbf{\Pi}_i^\top, & \text{if } i = j. \end{cases}$$

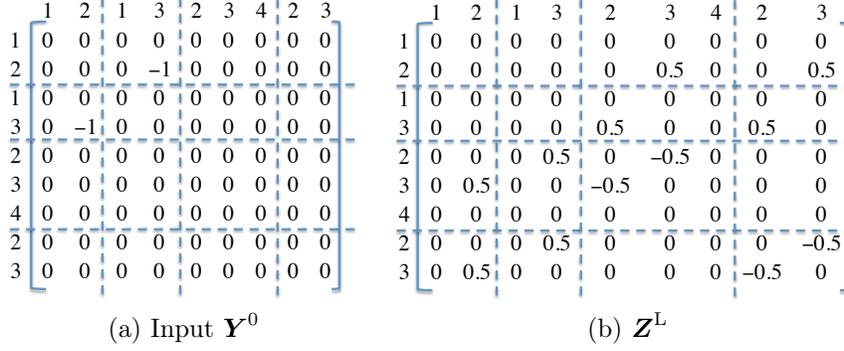


Figure 7: A toy example for constructing \mathbf{Z}^L , where 4 shapes $\mathcal{S}_1 = \{1, 2\}$, $\mathcal{S}_2 = \{1, 3\}$, $\mathcal{S}_3 = \{2, 3, 4\}$, and $\mathcal{S}_4 = \{2, 3\}$ are considered. The input incorrectly maps point 1 to 3 between \mathcal{S}_1 and \mathcal{S}_2 , and both points are contained in \mathcal{S}_3 and \mathcal{S}_4 . One can check that $\mathbf{Y}^0 + \mathbf{Z}^L \in T_{\text{gt}}^\perp$ in this example.

3. *Construction of \mathbf{Y}^L and \mathbf{Z}^L via an iterative procedure.* Next, we generate \mathbf{Y}^L via the following iterative procedure. Here, for any matrix M , we let $M_{ij}(s, s')$ represent the entry in the (i, j) th block M_{ij} that encodes the correspondence from s to s' .

Construction of a dual certificate \mathbf{Y}^L .

1. **initialize:** Set the symmetric matrix $\mathbf{Y}^{L,0}$ such that

$$\mathbf{Y}_{ij}^{L,0} = \begin{cases} -\mathbf{X}_{ij}^{\text{false}} + \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{E}_{ij}^\perp, & \text{if } i < j, \\ \mathbf{0}, & \text{if } i = j, \end{cases}$$

and start with $\mathbf{Z}^L = \mathbf{0}$.

2. **for** each *non-zero* entry $\mathbf{Y}_{ij}^{L,0}(s, s')$:
3. Set $a = \mathbf{Y}_{ij}^{L,0}(s, s')$, $B_{i,j,s,s'} = \{l \notin \{i, j\} \mid (s, s') \in \mathcal{S}_l\}$ and $n_{i,j}^{s,s'} = |B_{i,j,s,s'}|$.
4. **for** each set $l \in B_{i,j,s,s'}$: perform

$$\begin{cases} \mathbf{Z}_{il}^L(s, s') \leftarrow \mathbf{Z}_{il}^L(s, s') - \frac{a}{n_{i,j}^{s,s'}}, & \mathbf{Z}_{li}^L(s', s) \leftarrow \mathbf{Z}_{li}^L(s', s) - \frac{a}{n_{i,j}^{s,s'}}, \\ \mathbf{Z}_{lj}^L(s, s') \leftarrow \mathbf{Z}_{lj}^L(s, s') - \frac{a}{n_{i,j}^{s,s'}}, & \mathbf{Z}_{jl}^L(s', s) \leftarrow \mathbf{Z}_{jl}^L(s', s) - \frac{a}{n_{i,j}^{s,s'}}, \\ \mathbf{Z}_{ll}^L(s, s') \leftarrow \mathbf{Z}_{ll}^L(s, s') + \frac{a}{n_{i,j}^{s,s'}}, & \mathbf{Z}_{ll}^L(s', s) \leftarrow \mathbf{Z}_{ll}^L(s', s) + \frac{a}{n_{i,j}^{s,s'}}. \end{cases}$$

5. **output:** $\mathbf{Y}^L = \mathbf{Y}^{L,0} + \mathbf{Z}^L$.
-

4. *Construction of \mathbf{Y} and \mathbf{Z} :* define \mathbf{Y} and \mathbf{Z} such that

$$\mathbf{Y} = \mathbf{Y}^{\text{true},1} + \mathbf{Y}^{\text{true},2} + \mathbf{Y}^L + \alpha_{\beta_0} \mathbf{d} \cdot \mathbf{d}^\top, \quad (60)$$

$$\mathbf{Z}_{ij} = \begin{cases} \mathbf{Z}_{ij}^m - \mathbf{Z}_{ij}^L, & \text{if } i \neq j, \\ \mathbf{0}, & \text{if } i = j. \end{cases} \quad (61)$$

Remark 5. Below is a toy example to illustrate the proposed procedure for constructing \mathbf{Z}^L . Consider three sets $\mathcal{S}_1 = \{1, 2\}$, $\mathcal{S}_2 = \{1, 3\}$, $\mathcal{S}_3 = \{2, 3, 4\}$, and $\mathcal{S}_4 = \{1, 3\}$. Suppose that $\mathbf{Y}^{L,0}$ only contains two non-zero entries that incorrectly maps elements 1 to 3 in $\mathbf{Y}_{12}^{L,0}$, as illustrated in Fig. 7(a). The resulting \mathbf{Z}^L is shown in Fig. 7(b). Clearly, $\mathbf{Y}^{L,0} + \mathbf{Z}^L$ obeys $\mathbf{Y}^{L,0} + \mathbf{Z}^L \in T_{\text{gt}}^\perp$.

With the above construction procedure, one can easily verify that:

- (1) $\mathbf{Y}^{\text{true},1}$, $\mathbf{Y}^{\text{true},2}$ and \mathbf{Y}^{L} are all contained in the space T_{gt}^\perp ;
- (2) $\mathcal{P}_{\Omega^{\text{st}}}(\mathbf{Z}) = \mathbf{0}$;
- (3) If we set $\mathbf{M}^{\text{m}} := \alpha_{\beta_0} \mathbf{d} \cdot \mathbf{d}^\top$, then for any $i \neq j$,

$$\begin{aligned}
\mathbf{Y}_{ij} &= \mathbf{Y}_{ij}^{\text{true},1} + \mathbf{Y}_{ij}^{\text{true},2} + \mathbf{Y}_{ij}^{\text{L}} + \mathbf{M}_{ij}^{\text{m}} \\
&= -\mathbf{X}_{ij}^{\text{true}} + \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{E}_{ij} + \mathbf{R}_{ij}^{\text{m}} - \mathbf{X}_{ij}^{\text{false}} + \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{E}_{ij}^\perp + \mathbf{Z}_{ij}^{\text{L}} + \mathbf{M}_{ij}^{\text{m}} \\
&= -\mathbf{X}_{ij}^{\text{true}} - \mathbf{X}_{ij}^{\text{false}} + \lambda \mathbf{1} \cdot \mathbf{1}^\top - \left(\left(\lambda - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \right) \mathbf{1} \cdot \mathbf{1}^\top - \mathbf{R}_{ij}^{\text{m}} \right) + \mathbf{Z}_{ij}^{\text{L}} + \mathbf{M}_{ij}^{\text{m}} \\
&= \mathbf{W}_{ij} - \left(\mathbf{Z}_{ij}^{\text{m}} - \mathbf{Z}_{ij}^{\text{L}} \right).
\end{aligned} \tag{62}$$

Furthermore, from Lemma 2 one can obtain

$$\left\| \mathbf{d} \cdot \mathbf{d}^\top - \mathbf{1} \cdot \mathbf{1}^\top \right\|_\infty = O \left(\sqrt{\frac{\log(mn)}{np_{\text{set}}}} \right).$$

This taken collectively with (57) and the assumption (8) ensures that

$$\alpha_{\beta_0} = \lambda - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} - O \left(\sqrt{\frac{c_{10}p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}} \right) > 0 \tag{63}$$

as long as $p_{\text{set}}^3 > \frac{c_{15} \log(mn)}{n}$ for some constant $c_{15} > 0$.

Consequently, we will establish that \mathbf{Y} and \mathbf{Z} are valid dual certificates if they satisfy

$$\begin{cases} \text{all entries of } \mathbf{Z}_{ij}^{\text{m}} - \mathbf{Z}_{ij}^{\text{L}} \text{ (}\forall i \neq j\text{) within } \Omega_{\text{gt}}^\perp \text{ are strictly positive;} \\ \mathbf{Y}^{\text{true},1} + \mathbf{Y}^{\text{true},2} + \mathbf{Y}^{\text{L}} \geq \mathbf{0}. \end{cases} \tag{64}$$

Such conditions will be established through the following lemmas.

Lemma 7. *There are some universal constants $c_0, c_1 > 0$ such that*

$$\left\| \mathbf{Y}^{\text{L}} \right\| \leq c_0 \sqrt{\frac{np_{\text{obs}} \log(mn)}{p_{\text{set}}^2}}$$

and

$$\left\| \mathbf{Z}_{ij}^{\text{L}} \right\|_\infty \leq \sqrt{\frac{c_1 p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}}, \quad 1 \leq i < j \leq n$$

with probability exceeding $1 - \frac{1}{(mn)^4}$.

Proof. See Appendix D.5. □

Lemma 8. *There are some universal constants $c_5, c_6, c_7 > 0$ such that if $p_{\text{true}} p_{\text{obs}} p_{\text{set}} > \frac{c_7 \log^2(mn)}{n}$ and $\lambda < \frac{\sqrt{p_{\text{obs}} \log(mn)}}{p_{\text{set}}}$, then with probability exceeding $1 - \frac{1}{(mn)^{10}}$, one has*

$$\left\| \mathbf{Y}^{\text{true},2} \right\| \leq c_5 \sqrt{\frac{np_{\text{obs}}}{p_{\text{set}}}} \log(mn),$$

and

$$\langle \mathbf{v} \mathbf{v}^\top, \mathbf{Y}^{\text{true},1} \rangle \geq \frac{1}{2} np_{\text{set}} p_{\text{true}} p_{\text{obs}} - c_6 \sqrt{np_{\text{set}} p_{\text{obs}}} \log(mn)$$

for all unit vector \mathbf{v} satisfying $\mathbf{v} \mathbf{v}^\top \in T_{\text{gt}}^\perp$.

Proof. See Appendix D.6. □

Combining Lemmas 7 and 8 yields that there exists an absolute constant $c_0 > 0$ such that if

$$p_{\text{true}} > c_0 \frac{\log^2(mn)}{\sqrt{np_{\text{obs}} p_{\text{set}}^4}},$$

then

$$\mathbf{Y} = \mathbf{Y}^{\text{true},1} + \mathbf{Y}^{\text{true},2} + \mathbf{Y}^{\text{L}} \succeq \mathbf{0}.$$

On the other hand, observe that all entries of the non-negative matrix \mathbf{Z}^{m} lying in the index set $\Omega_{\text{gt}}^{\perp}$ are *bounded below* in magnitude by $\sqrt{\frac{c_{10} p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}}$. For sufficiently large c_{10} , one can conclude that all entries of $\mathbf{Z}_{il}^{\text{m}} - \mathbf{Z}_{il}^{\text{L}}$ outside Ω_{gt} are strictly positive.

So far we have justified that \mathbf{Y} and \mathbf{Z} satisfy (64), thereby certifying that the proposed algorithm correctly recovers the ground-truth matching.

D Proofs of Auxiliary Lemmas

D.1 Proof of Lemma 2

Denote by $\bar{\mathbf{A}} := \mathbf{1}_N \cdot \mathbf{1}_N^T$. From Bernstein inequality, n_i sharply concentrates around np such that if $p > \frac{c_6 \log^2(mn)}{n}$

$$|n_i - np| \leq c_5 \sqrt{np \log(mn)}, \quad \forall 1 \leq i \leq m \tag{65}$$

with probability exceeding $1 - (mn)^{-10}$, where $c_5, c_6 > 0$ are some absolute constants.

The bound (65) also implies that

$$\begin{aligned} \left\| \mathbf{I} - \begin{bmatrix} \frac{np}{n_1} & & & \\ & \frac{np}{n_2} & & \\ & & \ddots & \\ & & & \frac{np}{n_m} \end{bmatrix} \right\| &\leq \max_{1 \leq i \leq m} \frac{|n_i - np|}{n_i} \leq \frac{c_5 \sqrt{np \log(mn)}}{np - c_5 \sqrt{np \log(mn)}} \\ &\leq 2c_5 \sqrt{\frac{\log(mn)}{np}}. \end{aligned}$$

Similarly, one has

$$|N - nmp| \leq c_5 \sqrt{pmn \log(mn)}$$

with probability exceeding $1 - (mn)^{-10}$, which implies that

$$\|\bar{\mathbf{A}}\| = N \leq nmp + c_5 \sqrt{pmn \log(mn)} < 2nmp.$$

Rewrite \mathbf{A} as

$$\mathbf{A} := \begin{bmatrix} \frac{np}{n_1} \text{Diag}(\mathbf{1}_{n_1}) & & & \\ & \ddots & & \\ & & \frac{np}{n_m} \text{Diag}(\mathbf{1}_{n_m}) & \\ & & & \ddots \end{bmatrix} \cdot \bar{\mathbf{A}} \cdot \begin{bmatrix} \frac{np}{n_1} \text{Diag}(\mathbf{1}_{n_1}) & & & \\ & \ddots & & \\ & & \frac{np}{n_m} \text{Diag}(\mathbf{1}_{n_m}) & \\ & & & \ddots \end{bmatrix}.$$

This allows us to bound the deviation of \mathbf{A} from $\overline{\mathbf{A}}$ as follows

$$\begin{aligned}
\|\mathbf{A} - \overline{\mathbf{A}}\| &\leq \left\| \mathbf{A} - \begin{bmatrix} \frac{np}{n_1} \text{Diag}(\mathbf{1}_{n_1}) & & \\ & \ddots & \\ & & \frac{np}{n_m} \text{Diag}(\mathbf{1}_{n_m}) \end{bmatrix} \overline{\mathbf{A}} \right\| + \left\| \begin{bmatrix} \frac{np}{n_1} \text{Diag}(\mathbf{1}_{n_1}) & & \\ & \ddots & \\ & & \frac{np}{n_m} \text{Diag}(\mathbf{1}_{n_m}) \end{bmatrix} \overline{\mathbf{A}} - \overline{\mathbf{A}} \right\| \\
&\leq \left(\left\| \begin{bmatrix} \frac{np}{n_1} \text{Diag}(\mathbf{1}_{n_1}) & & \\ & \ddots & \\ & & \frac{np}{n_m} \text{Diag}(\mathbf{1}_{n_m}) \end{bmatrix} \right\| + 1 \right) \|\overline{\mathbf{A}}\| \left\| \mathbf{I} - \begin{bmatrix} \frac{np}{n_1} \text{Diag}(\mathbf{1}_{n_1}) & & \\ & \ddots & \\ & & \frac{np}{n_m} \text{Diag}(\mathbf{1}_{n_m}) \end{bmatrix} \right\| \\
&\leq \left(1 + c_5 \sqrt{\frac{\log(mn)}{np}} + 1 \right) 2nmp \cdot 2c_5 \sqrt{\frac{\log(mn)}{np}} \\
&\leq c_6 m \sqrt{np \log(mn)}
\end{aligned}$$

for some universal constant $c_6 > 0$.

On the other hand, it follows immediately from (65) that

$$\begin{aligned}
\|\mathbf{A} - \mathbf{1} \cdot \mathbf{1}^\top\|_\infty &= \max_{1 \leq i, j \leq m} \left| \frac{(np)^2}{n_i n_j} - 1 \right| = \max_{1 \leq i, j \leq m} \left| \frac{pn(pn - n_j) + (pn - n_i)n_j}{n_i n_j} \right| \\
&\leq \max_{1 \leq i, j \leq m} \frac{|pn + c_5 \sqrt{np \log(mn)}|}{(pn - c_5 \sqrt{np \log(mn)})^2} c_5 \sqrt{np \log(mn)} \\
&\leq c_9 \sqrt{\frac{\log(mn)}{np}}
\end{aligned}$$

for some absolute constant $c_9 > 0$.

D.2 Proof of Lemma 3

The norm of \mathbf{M} can be bounded via the moment method, which attempts to control $\text{tr}(\mathbf{M}^k)$ for some even integer k . See [48, Section 2.3.4] for a nice introduction.

Specifically, observe that $\mathbb{E} \text{tr}(\mathbf{M}^k)$ can be expanded as follows

$$\mathbb{E} \text{tr}(\mathbf{M}^k) = \sum_{1 \leq i_1, \dots, i_k \leq n} \mathbb{E} \text{tr}(\mathbf{M}_{i_1 i_2} \mathbf{M}_{i_2 i_3} \cdots \mathbf{M}_{i_k i_1}),$$

a trace sum over all k -cycles in the vertex set $\{1, \dots, n\}$. Note that (i, i) are also treated as valid edges. For each term $\mathbb{E} \text{tr}(\mathbf{M}_{i_1 i_2} \mathbf{M}_{i_2 i_3} \cdots \mathbf{M}_{i_k i_1})$, if there exists an edge occurring exactly once, then the term vanishes due to the independence assumption. Thus, it suffices to examine the terms in which each edge is repeated at least twice. Consequently, there are at most $k/2$ relevant edges, which span at most $k/2 + 1$ distinct vertices. We also need to assign vertices to $k/2$ edges, which adds up to no more than $(k/2)^k$ different choices.

By following the same procedure and notation as adopted in [48, Page 119], we divide all non-vanishing k -cycles into $(k/2)^k$ classes based on the above labeling order; each class is associated with j ($1 \leq j \leq k/2$) edges e_1, \dots, e_j with multiplicities a_1, \dots, a_j , where $(e_1, \dots, a_1, \dots, a_j)$ determines the class of cycles and $a_1 + \dots + a_j = k$. Since there are at most n^{j+1} distinct vertices, one can see that no more than n^{j+1} cycles falling within this particular class. For notational simplicity, set $K = \sqrt{n}$, and hence $\|\mathbf{M}_{ij}\| \leq K$. By assumption (46), one has

$$\begin{aligned}
\mathbb{E} \text{tr}(\mathbf{M}_{i_1 i_2} \mathbf{M}_{i_2 i_3} \cdots \mathbf{M}_{i_k i_1}) &\leq m \mathbb{E} (\|\mathbf{M}_{e_1}\|^{a_1} \cdots \|\mathbf{M}_{e_j}\|^{a_j}) \\
&\leq m \mathbb{E} \|\mathbf{M}_{e_1}\|^2 \cdots \mathbb{E} \|\mathbf{M}_{e_j}\|^2 K^{a_1 - 2} \cdots K^{a_j - 2} \\
&\leq m K^{k - 2j}.
\end{aligned}$$

Thus, the total contribution of this class does not exceed

$$mn^{j+1}K^{k-2j} = mn^{\frac{k}{2}+1}.$$

By summing over all classes one obtains the crude bound

$$\mathbb{E}\text{tr}(\mathbf{M}^k) \leq m \left(\frac{k}{2}\right)^k n^{\frac{k}{2}+1},$$

which follows that

$$\mathbb{E}\|\mathbf{M}\|^k \leq \mathbb{E}\text{tr}(\mathbf{M}^k) \leq m \left(\frac{k}{2}\right)^k n^{\frac{k}{2}+1}.$$

If we set $k = \log(mn)$, then from Markov's inequality we have

$$\mathbb{P}\left(\|\mathbf{M}\| \geq \frac{k}{2}n^{\frac{1}{2}+\frac{1}{k}}(mn)^{\frac{5}{k}}m^{\frac{1}{k}}\right) \leq \frac{\mathbb{E}\|\mathbf{M}\|^k}{\left(\frac{k}{2}n^{\frac{1}{2}+\frac{1}{k}}(mn)^{\frac{5}{k}}m^{\frac{1}{k}}\right)^k} \leq \frac{m\left(\frac{k}{2}\right)^k n^{\frac{k}{2}+1}}{m\left(\frac{k}{2}\right)^k n^{\frac{k}{2}+1}(mn)^5} \leq \frac{1}{(mn)^5}.$$

Since $n^{\frac{1}{\log n}} = O(1)$, there exists a constant $c_0 > 0$ such that

$$\mathbb{P}\left(\|\mathbf{M}\| \geq c_0n^{\frac{1}{2}}\log(mn)\right) \leq \frac{1}{m^5n^5},$$

which completes the proof.

D.3 Proof of Lemma 4

When $\mathcal{G} \sim \mathcal{G}(n, p)$, the adjacency matrix \mathbf{A} consists of independent Bernoulli components (except for diagonal entries), each with mean p and variance $p(1-p)$. Lemma 3 immediately implies that if $p > \frac{2\log(mn)}{n}$, then

$$\frac{1}{\sqrt{p(1-p)}}\|\mathbf{A} - p\mathbf{1}_n \cdot \mathbf{1}_n^\top\| \leq c_0\sqrt{n}\log(mn) + 1 \quad (66)$$

with probability at least $1 - (mn)^{-5}$. That said, there exists an absolute constant $c_1 > 0$ such that

$$\|\mathbf{A} - p\mathbf{1}_n \cdot \mathbf{1}_n^\top\| \leq c_1\sqrt{pn}\log(mn) \quad (67)$$

with probability exceeding $1 - (mn)^{-5}$.

On the other hand, from Bernstein inequality, the degree of each vertex exceeds

$$d_{\min} := pn - c_2\sqrt{pn\log(mn)} \quad (68)$$

with probability at least $1 - (mn)^{-10}$, where c_2 is some constant. When $p > \frac{2\log(mn)}{n}$, \mathcal{G} is connected, and hence the least eigenvalue of \mathbf{L} is zero with the eigenvector $\mathbf{1}_n$. This taken collectively with (67) and (68) suggests that when $p > \frac{c_3^2\log^2(mn)}{n}$, one has

$$a(\mathcal{G}) \geq d_{\min} - \|\mathbf{A} - p\mathbf{1}_n \cdot \mathbf{1}_n^\top\| \geq pn - c_3\sqrt{pn}\log(mn)$$

with high probability.

D.4 Proof of Lemma 6

Suppose that $\mathbf{X}^{\text{gt}} + \mathbf{H}$ is the solution to MatchLift for some perturbation $\mathbf{H} \neq \mathbf{0}$. By Schur complement condition for positive definiteness, the feasibility constraint $\begin{bmatrix} m & \mathbf{1}^\top \\ \mathbf{1} & \mathbf{X}^{\text{gt}} + \mathbf{H} \end{bmatrix} \succeq \mathbf{0}$ is equivalent to

$$\begin{cases} \mathbf{X}^{\text{gt}} + \mathbf{H} & \succeq \mathbf{0}, \\ \mathbf{X}^{\text{gt}} + \mathbf{H} - \frac{1}{m}\mathbf{1} \cdot \mathbf{1}^\top & \succeq \mathbf{0}, \end{cases}$$

which immediately yields

$$\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) = (\mathbf{I} - \mathbf{U}\mathbf{U}^\top)(\mathbf{X}^{\text{gt}} + \mathbf{H})(\mathbf{I} - \mathbf{U}\mathbf{U}^\top) \succeq \mathbf{0}, \quad (69)$$

and

$$\langle \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{H} \rangle = \left\langle \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{X}^{\text{gt}} - \frac{1}{m} \mathbf{1} \cdot \mathbf{1}^\top + \mathbf{H} \right\rangle \geq 0. \quad (70)$$

The above inequalities follow from the facts $\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{X}^{\text{gt}}) = \mathbf{0}$ and $\langle \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{X}^{\text{gt}} - \frac{1}{m} \mathbf{1} \cdot \mathbf{1}^\top \rangle = 0$.

From Assumption (51), one can derive

$$\begin{aligned} \langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle + \langle \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{H} \rangle &= \langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{H} \rangle + \langle \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{H} \rangle \\ &= \langle \mathbf{Y}, \mathbf{H} \rangle = \sum_{i \neq j} \langle \mathbf{Y}_{ij}, \mathbf{H}_{ij} \rangle. \end{aligned} \quad (71)$$

This allows us to bound

$$\begin{aligned} &\langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle + \sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle \\ &\leq \langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle + \langle \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{H} \rangle + \sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle \end{aligned} \quad (72)$$

$$= \sum_{i \neq j} \langle \mathbf{Y}_{ij}, \mathbf{H}_{ij} \rangle + \sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle \quad (73)$$

$$= \sum_{i \neq j} \langle \mathbf{W}_{ij}, \mathbf{H}_{ij} \rangle, \quad (74)$$

where the first inequality follows from (70), and the last equality follows from Assumption (50).

In order to preclude the possibility that $\mathbf{X}^{\text{gt}} + \mathbf{H}$ is the solution to MatchLift, we need to show that $\sum_{i \neq j} \langle \mathbf{W}_{ij}, \mathbf{H}_{ij} \rangle > 0$. From (74) it suffices to establish that

$$\langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle + \sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle > 0 \quad (75)$$

for any feasible $\mathbf{H} \neq \mathbf{0}$. In fact, since $\mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top$ and $\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H})$ are both positive semidefinite, one must have

$$\langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle \geq 0. \quad (76)$$

On the other hand, the constraints

$$\text{supp}(\mathbf{Z}) \subseteq \Omega_{\text{gt}}^\perp, \quad \mathcal{P}_{\Omega_{\text{gt}}^\perp}(\mathbf{Z}) \geq \mathbf{0}, \quad \text{and} \quad \mathcal{P}_{\Omega_{\text{gt}}^\perp}(\mathbf{H}) \geq \mathbf{0}$$

taken together imply that

$$\sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle \geq 0. \quad (77)$$

Putting (76) and (77) together gives

$$\langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle + \sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle \geq 0.$$

Comparing this with (75), we only need to establish either $\langle \mathbf{Y} - \alpha \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle > 0$ or $\sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle > 0$.

i) Suppose first that all entries of \mathbf{Z}_{ij} ($\forall i \neq j$) in the support Ω_{gt}^\perp are strictly positive. If the identity $\sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle = 0$ holds, then the strict positivity assumption of \mathbf{Z}_{ij} on Ω_{gt}^\perp as well as the constraint $\mathcal{P}_{\Omega_{\text{gt}}^\perp}(\mathbf{H}) \geq \mathbf{0}$ immediately leads to

$$\mathcal{P}_{\Omega_{\text{gt}}^\perp}(\mathbf{H}) = \mathbf{0}.$$

Besides, the feasibility constraint requires that $\mathcal{P}_{\Omega_{\text{gt}}}(\mathbf{H}_{ij}) \leq \mathbf{0}$. If $\mathcal{P}_{\Omega_{\text{gt}}}(\mathbf{H}_{ij}) \neq \mathbf{0}$, then all non-zero entries of \mathbf{H}_{ij} are *negative*, and hence

$$\langle \mathbf{d} \cdot \mathbf{d}^\top, \mathbf{H} \rangle = \langle \mathbf{d} \cdot \mathbf{d}^\top, \mathcal{P}_{\Omega_{\text{gt}}}(\mathbf{H}) \rangle < 0,$$

which follows since all entries of \mathbf{d} are strictly positive. This contradicts with (70). Consequently, we must either have $\mathbf{H} = \mathbf{0}$ or $\sum_{i \neq j} \langle \mathbf{Z}_{ij}, \mathbf{H}_{ij} \rangle > 0$. This together with (75) establishes the claim.

ii) Next, we prove the claim under Assumptions (52) and (53). In fact, Assumption (52) together with (69) asserts that $\langle \mathbf{Y}, \mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) \rangle \leq 0$ can only occur if $\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) = \mathbf{0}$. This necessarily leads to $\mathbf{H} = \mathbf{0}$, as claimed by Lemma 9.

Lemma 9. *Suppose that $\mathbf{X}^{\text{gt}} + \mathbf{H}$ is feasible for MatchLift, and assume that*

$$\frac{n}{n_i} + \frac{n}{n_j} \neq \frac{n^2}{n_i n_j}, \quad \forall 1 \leq i, j \leq m. \quad (78)$$

If $\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) = \mathbf{0}$, then one has $\mathbf{H} = \mathbf{0}$.

Proof. See Appendix D.7. □

In summary, we can conclude that \mathbf{X}^{gt} is the unique optimizer in both cases.

D.5 Proof of Lemma 7

First, we would like to bound the operator norm of \mathbf{Y}^{L} . Since each random matrix $\mathbf{X}_{ij}^{\text{in}} \mathbb{I}_{\{\mathbf{X}_{ij}^{\text{in}} \text{ is observed and corrupted}\}}$ is independently drawn with mean $\frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{1} \cdot \mathbf{1}^\top$, it is straightforward to see that

$$\mathbb{E} \mathbf{Y}^{\text{L},0} = \mathbb{E} \left(-\mathbf{X}^{\text{false}} + \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \mathbf{E}^\perp \right) = \mathbf{0}.$$

By observing that \mathbf{Z}^{L} is constructed as a linear transform of $\mathbf{Y}^{\text{L},0}$, one can also obtain

$$\mathbb{E} \mathbf{Z}^{\text{L}} = \mathbf{0}, \quad \Rightarrow \quad \mathbb{E} \mathbf{Y}^{\text{L}} = \mathbb{E} \mathbf{Z}^{\text{L}} + \mathbb{E} \mathbf{Y}^{\text{L},0} = \mathbf{0}.$$

Thus, it suffices to examine the deviation of $\|\mathbf{Y}^{\text{L}}\|$ incurred by the uncertainty of $\mathbf{X}^{\text{false}}$.

Denote by $\mathbf{A}^{i,j} \in \mathbb{R}^{N \times N}$ the component of \mathbf{Z}^{L} generated due to the $(i, j)^{\text{th}}$ block $-\mathbf{X}_{ij}^{\text{false}}$, which clearly satisfies

$$\mathbf{Z}^{\text{L}} = \mathbf{A}^{i,j} - \mathbb{E} \mathbf{A}^{i,j}.$$

For each non-zero entry of $\mathbf{X}_{ij}^{\text{false}}$, if it encodes an incorrect correspondence between elements s and t , then it will affect no more than $6n_{s,t}$ entries in $\mathbf{A}^{i,j}$, where each of these entries are affected in magnitude by an amount at most $\frac{1}{n_{s,t}}$. Recall that $n_{s,t}$ represents the number of sets \mathcal{S}_i ($1 \leq i \leq n$) containing s and t simultaneously, which sharply concentrates within $\left[np_{\text{set}}^2 \pm O\left(\sqrt{np_{\text{set}}^2 \log(mn)}\right) \right]$ as asserted in Lemma 5. As a result, the *sum of squares* of these affected entries is bounded by

$$\frac{6n_{s,t}}{n_{s,t}^2} = O\left(\frac{1}{n_{s,t}}\right). \quad (79)$$

Moreover, since each row / column of $\mathbf{X}_{ij}^{\text{false}}$ can have at most one non-zero entry, we can rearrange $\mathbf{A}^{i,j}$ with row / column permutation such that $\mathbf{A}^{i,j}$ becomes a block-diagonal matrix, where the components

affected by different entries of $\mathbf{X}_{ij}^{\text{false}}$ are separated into distinct diagonal blocks. This together with (79) leads to

$$\|\mathbf{A}^{i,j}\| \leq \|\mathbf{A}^{i,j}\|_{\text{F}} \leq \max_{s \neq t} \sqrt{\frac{8}{n_{s,t}}},$$

and hence

$$\|\mathbb{E}\mathbf{A}^{i,j} (\mathbf{A}^{i,j})^{\top}\| \leq p_{\text{obs}} \left(\max_{s \neq t} \sqrt{\frac{8}{n_{s,t}}} \right)^2 \leq \frac{c_{16} p_{\text{obs}}}{n p_{\text{set}}^2}$$

for some absolute constant $c_{16} > 0$, where the last inequality follows from Lemma 5.

Observe that $\mathbf{A}^{i,j} - \mathbb{E}\mathbf{A}^{i,j}$ ($i \neq j$) are independently generated with mean zero, whose operator norm is bounded above by $2 \max_{s \neq t} \sqrt{\frac{8}{n_{s,t}}}$. Applying the matrix Bernstein inequality [49, Theorem 1.4] suggests that there exist universal constants $c_5, c_6 > 0$ such that for any $t = O(\sqrt{n} \text{poly} \log(mn))$,

$$\mathbb{P} \left(\left\| \sum_{(i,j) \in \mathcal{G}} \mathbf{A}^{i,j} - \mathbb{E}\mathbf{A}^{i,j} \right\| > t \right) \leq n^2 \exp \left(- \frac{\frac{1}{2} t^2}{n^2 \left(\frac{c_{16} p_{\text{obs}}}{n p_{\text{set}}^2} \right) + \frac{2 \max_{s \neq t} \sqrt{\frac{8}{n_{s,t}}}}{3}} \right).$$

Put in another way, there exists a universal constant $c_6 > 0$ such that

$$\|\mathbf{Z}^{\text{L}}\| = \left\| \sum_{i \neq j} \mathbf{A}^{i,j} - \mathbb{E}\mathbf{A}^{i,j} \right\| < c_6 \sqrt{\frac{n p_{\text{obs}}}{p_{\text{set}}^2} \log(mn)} \quad (80)$$

holds with probability exceeding $1 - \frac{1}{(mn)^{10}}$. This follows from Lemma 5.

Additionally, observe that $\mathbb{E}\mathbf{Y}_{ij}^{\text{L},0} = \mathbf{0}$ and

$$\left\| \frac{1}{\sqrt{p_{\text{obs}}}} \mathbf{Y}_{ij}^{\text{L},0} \right\| \leq \sqrt{n}$$

as long as $p_{\text{obs}} > \frac{1}{n}$. Applying Lemma 3 suggests that

$$\|\mathbf{Y}^{\text{L},0}\| < c_0 \sqrt{n p_{\text{obs}} \log(mn)}$$

with probability at least $1 - \frac{1}{(mn)^5}$. This combined with (80) yields

$$\|\mathbf{Y}^{\text{L}}\| \leq \|\mathbf{Y}^{\text{L},0}\| + \|\mathbf{Z}^{\text{L}}\| < c_{11} \sqrt{\frac{n p_{\text{obs}} \log(mn)}{p_{\text{set}}^2}}$$

with probability at least $1 - \frac{3}{(mn)^5}$, where c_{11} is some universal constant.

On the other hand, for each (s, t) entry of $\mathbf{Z}_{il}^{\text{L}}$ ($i \neq l$), it can only be affected by those *observed* blocks $\mathbf{X}_{ij}^{\text{false}}$ (or $\mathbf{X}_{jl}^{\text{false}}$) satisfying $t \in \mathcal{S}_j$ (or $s \in \mathcal{S}_j$). Consequently, each entry of $\mathbf{Z}_{il}^{\text{L}}$ can be expressed as a sum of $\Theta(np_{\text{set}} p_{\text{obs}})$ zero-mean independent variables, each of them being bounded in magnitude by $\frac{1}{(\min_{s \neq t} n_{s,t})}$. From Hoeffding's inequality one can derive

$$\mathbb{P} \left(\left\| \mathbf{Z}_{il}^{\text{L}} \right\|_{\infty} > t \right) \leq m^2 \mathbb{P} \left(- \frac{t^2}{c_7 n p_{\text{set}} p_{\text{obs}} \frac{1}{\left(\min_{s \neq t} n_{s,t} \right)^2}} \right) \leq m^2 \mathbb{P} \left(- \frac{t^2}{\tilde{c}_7 p_{\text{obs}} \frac{1}{n p_{\text{set}}^3}} \right)$$

for some constants $c_7, \tilde{c}_7 > 0$, indicating that

$$\left\| \mathbf{Z}_{il}^{\text{L}} \right\|_{\infty} \leq \sqrt{\frac{c_8 p_{\text{obs}} \log(mn)}{n p_{\text{set}}^3}}, \quad \forall i \neq l$$

with probability exceeding $1 - \frac{1}{(mn)^{10}}$.

D.6 Proof of Lemma 8

By construction of $\mathbf{Y}^{\text{true},1}$, one can see that all non-zero entries lie within the support Ω_{gt} . One important feature of $\mathbf{X}_{ij}^{\text{gt}}$ is that it can be converted, via row / column permutation, into a block diagonal matrix that consists of m all-one blocks, where the i^{th} block is of size n_i ($1 \leq i \leq m$). From Lemma 5, one has

$$n_i \in \left[np_{\text{set}} \pm \sqrt{c_8 np_{\text{set}} \log(mn)} \right], \quad 1 \leq i \leq m$$

with high probability. Thus, $\mathbf{Y}^{\text{true},1}$ can also be rearranged such that its non-zero entries form m disjoint diagonal blocks. We will quantify the eigenvalues of $\mathbf{Y}^{\text{true},1}$ by bounding the spectrum of each of these matrix blocks.

We first decompose the matrix $\mathbf{Y}^{\text{true},1}$ into two parts $\bar{\mathbf{Y}}^{\text{true},1}$ and $\tilde{\mathbf{Y}}^{\text{true},1}$ such that

$$\forall i \neq j, \quad \bar{\mathbf{Y}}_{ij}^{\text{true},1} = \begin{cases} -\mathbf{X}_{ij}^{\text{in}}, & \text{if } \mathbf{X}_{ij}^{\text{in}} \text{ is observed and not corrupted,} \\ \mathbf{0}, & \text{else;} \end{cases}$$

and

$$\forall i \neq j, \quad \tilde{\mathbf{Y}}_{ij}^{\text{true},1} = \begin{cases} -\mathbf{X}_{ij}^{\text{in}} + \frac{(1-p_{\text{true}})p_{\text{obs}}}{m}, & \text{if } \mathbf{X}_{ij}^{\text{in}} \text{ is observed and corrupted,} \\ \frac{(1-p_{\text{true}})p_{\text{obs}}}{m}, & \text{else.} \end{cases}$$

That said, $\bar{\mathbf{Y}}_{ij}^{\text{true},1}$ consists of all non-corrupted components, while $\tilde{\mathbf{Y}}^{\text{true},1}$ consists of all ‘‘debiased’’ random outliers.

By Lemma 4, one can verify that for all unit vector \mathbf{v} such that $\mathbf{v}\mathbf{v}^\top \in T_{\text{gt}}^\perp$,

$$\begin{aligned} \langle \mathbf{v}\mathbf{v}^\top, \bar{\mathbf{Y}}^{\text{true},1} \rangle &\geq \min_{1 \leq s \leq m} (n_s p_{\text{true}} p_{\text{obs}} - c_4 \sqrt{n_s p_{\text{obs}}} \log(mn)) \\ &\geq \frac{1}{2} np_{\text{set}} p_{\text{true}} p_{\text{obs}} - c_5 \sqrt{np_{\text{set}} p_{\text{obs}}} \log(mn) \end{aligned} \quad (81)$$

for some absolute constant $c_5 > 0$, where the second inequality follows from the concentration result stated in Lemma 5.

In addition, each entry of $\tilde{\mathbf{Y}}_{ij}^{\text{true},1}$ ($i \neq j$) lying in the support Ω_{gt} has mean zero and variance $\frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \left(1 - \frac{(1-p_{\text{true}})p_{\text{obs}}}{m} \right)$

Lemma 3 then suggests that the norm of each non-zero block of $\tilde{\mathbf{Y}}^{\text{true},1}$ (the ones with size n_i) is bounded above by $O(\sqrt{p_{\text{obs}} n_i} \log(nm))$. As a result,

$$\left\| \tilde{\mathbf{Y}}^{\text{true},1} \right\| \leq c_{15} \max_{1 \leq s \leq m} \sqrt{p_{\text{obs}} n_s} \log(nm) < \tilde{c}_{15} \sqrt{np_{\text{set}} p_{\text{obs}}} \log(nm).$$

This taken collectively with (81) yields that

$$\langle \mathbf{v}\mathbf{v}^\top, \mathbf{Y}^{\text{true},1} \rangle \geq \frac{1}{2} np_{\text{set}} p_{\text{true}} p_{\text{obs}} - (c_5 + \tilde{c}_{15}) \sqrt{np_{\text{set}} p_{\text{obs}}} \log(mn). \quad (82)$$

On the other hand, we know from the construction procedure and Lemma 2 that

$$\begin{aligned} \|\mathbf{R}^{\text{m}}\|_\infty &\leq \sqrt{\frac{c_{10} p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}} + \lambda \left\| \mathbf{d} \cdot \mathbf{d}^\top - \mathbf{1} \cdot \mathbf{1}^\top \right\|_\infty \\ &\leq \tilde{c}_{10} \left(\sqrt{\frac{p_{\text{obs}} \log(mn)}{np_{\text{set}}^3}} + \frac{\sqrt{p_{\text{obs}} \log(mn)}}{p_{\text{set}}} \sqrt{\frac{\log(mn)}{np_{\text{set}}}} \right) \\ &\leq 2\tilde{c}_{10} \sqrt{\frac{p_{\text{obs}}}{np_{\text{set}}^3}} \log(mn) \end{aligned}$$

for some constants $c_{10}, \tilde{c}_{10} > 0$. Since $\mathbf{R}^m \in \Omega_{\text{gt}}$, we can also rearrange \mathbf{R}^m into m diagonal blocks each of size n_i ($1 \leq i \leq m$). Hence, a crude upper bound yields

$$\begin{aligned} \|\mathbf{Y}^{\text{true},2}\| &\leq \|\mathbf{Y}^{\text{true},2}\|_1 \leq \left(\max_{1 \leq i \leq m} n_i \right) \left(2\tilde{c}_{10} \sqrt{\frac{p_{\text{obs}}}{np_{\text{set}}^3}} \log(mn) \right) \leq c_{11} np_{\text{set}} \sqrt{\frac{np_{\text{obs}}}{p_{\text{set}}^3}} \log(mn) \\ &= c_{11} n \sqrt{\frac{np_{\text{obs}}}{p_{\text{set}}}} \log(mn) \end{aligned}$$

for some universal constant $c_{11} > 0$.

D.7 Proof of Lemma 9

Define an augmented matrix \mathbf{H}^{sup} such that

$$\mathbf{H}_{ij}^{\text{sup}} = \boldsymbol{\Pi}_i^\top \mathbf{H}_{ij} \boldsymbol{\Pi}_j. \quad (83)$$

Recall that n_i denotes the number of sets containing element i , and that

$$\boldsymbol{\Sigma} := \begin{bmatrix} \frac{n}{n_1} & & & \\ & \frac{n}{n_2} & & \\ & & \ddots & \\ & & & \frac{n}{n_m} \end{bmatrix}.$$

The assumption that $\mathcal{P}_{T_{\text{gt}}^\perp}(\mathbf{H}) = \mathbf{0}$ can be translated into

$$\left(\mathbf{I} - \frac{1}{n} (\mathbf{1}_n \otimes \mathbf{I}_m) \boldsymbol{\Sigma} (\mathbf{1}_n \otimes \mathbf{I}_m) \right) \mathbf{H}^{\text{sup}} \left(\mathbf{I} - \frac{1}{n} (\mathbf{1}_n \otimes \mathbf{I}_m) \boldsymbol{\Sigma} (\mathbf{1}_n \otimes \mathbf{I}_m) \right) = \mathbf{0}.$$

We can easily compute that

$$\mathbf{H}_{ii}^{\text{sup}} - \boldsymbol{\Sigma} \overline{\mathbf{H}}_{\cdot i}^{\text{sup}} - \overline{\mathbf{H}}_{i \cdot}^{\text{sup}} \boldsymbol{\Sigma} + \boldsymbol{\Sigma} \overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} \boldsymbol{\Sigma} = \mathbf{0}, \quad 1 \leq i \leq n,$$

where

$$\begin{cases} \overline{\mathbf{H}}_{\cdot i}^{\text{sup}} &:= \frac{1}{n} \sum_{j=1}^n \mathbf{H}_{ji}^{\text{sup}}, \\ \overline{\mathbf{H}}_{i \cdot}^{\text{sup}} &:= \frac{1}{n} \sum_{j=1}^n \mathbf{H}_{ij}^{\text{sup}}, \\ \overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} &:= \frac{1}{n^2} \sum_{i=1}^n \sum_{j=1}^n \mathbf{H}_{ij}^{\text{sup}}. \end{cases}$$

This combined with the identity $\mathbf{H}_{ii} = \mathbf{0}$ (and hence $\mathbf{H}_{ii}^{\text{sup}} = \mathbf{0}$) yields

$$\boldsymbol{\Sigma} \overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} \boldsymbol{\Sigma} = \boldsymbol{\Sigma} \overline{\mathbf{H}}_{\cdot i}^{\text{sup}} + \overline{\mathbf{H}}_{i \cdot}^{\text{sup}} \boldsymbol{\Sigma}, \quad 1 \leq i \leq n.$$

Summing over all i leads to

$$\boldsymbol{\Sigma} \overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} \boldsymbol{\Sigma} = \boldsymbol{\Sigma} \left(\frac{1}{n} \sum_{i=1}^n \overline{\mathbf{H}}_{\cdot i}^{\text{sup}} \right) + \left(\frac{1}{n} \sum_{i=1}^n \overline{\mathbf{H}}_{i \cdot}^{\text{sup}} \right) \boldsymbol{\Sigma} = \boldsymbol{\Sigma} \overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} + \overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} \boldsymbol{\Sigma}.$$

Expanding it yields

$$\frac{n^2}{n_i n_j} \left(\overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} \right)_{i,j} = \left(\frac{n}{n_i} + \frac{n}{n_j} \right) \left(\overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} \right)_{i,j}, \quad 1 \leq i, j \leq m.$$

From our assumption that $\frac{n^2}{n_i n_j} \neq \frac{n}{n_i} + \frac{n}{n_j}$, we can derive

$$\overline{\mathbf{H}}_{\cdot \cdot}^{\text{sup}} = \mathbf{0}. \quad (84)$$

Due to the feasibility constraint, all diagonal entries of $\mathbf{H}_{ij}^{\text{sup}}$ are non-positive, and all off-diagonal entries of $\mathbf{H}_{ij}^{\text{sup}}$ are non-negative. These conditions together with (84) establish that $\mathbf{H} = \mathbf{0}$.

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