

Solving a Special Type of Optimal Transport Problem by a Modified Hungarian Algorithm

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Abstract

Computing the empirical Wasserstein distance in the Wasserstein-distance-based independence test is an optimal transport (OT) problem with a special structure. This observation inspires us to study a special type of OT problem and propose *a modified Hungarian algorithm* to solve it *exactly*. For the OT problem involving two marginals with m and n atoms ($m \geq n$), respectively, the computational complexity of the proposed algorithm is $\mathcal{O}(m^2n)$. Computing the empirical Wasserstein distance in the independence test requires solving this special type of OT problem, where $m = n^2$. The associated computational complexity of the proposed algorithm is $\mathcal{O}(n^5)$, while the order of applying the classic Hungarian algorithm is $\mathcal{O}(n^6)$. In addition to the aforementioned special type of OT problem, it is shown that the modified Hungarian algorithm could be adopted to solve a wider range of OT problems. Broader applications of the proposed algorithm are discussed—solving the one-to-many assignment problem and the many-to-many assignment problem. We conduct numerical experiments to validate our theoretical results. The experiment results demonstrate that the proposed modified Hungarian algorithm compares favorably with the Hungarian algorithm and the well-known Sinkhorn algorithm.

1 Introduction

One appealing application of optimal transport (OT) and Wasserstein distance (Villani, 2009; Peyré et al., 2019) is the independence test. The Wasserstein distance between two distributions μ_1, μ_2 on Z is defined as:

$$W(\mu_1, \mu_2) := \inf \left\{ \int_{Z^2} d(z, z') d\gamma(z, z') : \gamma \text{ is a distribution with marginals } \mu_1 \text{ and } \mu_2 \right\},$$

where (Z, d) is a metric space (w.l.o.g., 1-Wasserstein distance is considered in this paper). The Wasserstein distance is a metric on probability measures (Villani, 2009). To test the independence between the variables $Y \sim \nu_1$ and $Z \sim \nu_2$, people utilize the Wasserstein distance between the joint distribution of Y, Z and the product distribution of Y, Z , i.e., $W(\gamma(\nu_1, \nu_2), \nu_1 \otimes \nu_2)$. While the statistical properties of this approach have been intensely investigated (Nies et al., 2021; Mordant & Segers, 2022; Wiesel, 2022), no existing literature focuses on the computational aspect. In this paper, we discuss the following:

How to compute the empirical Wasserstein distance in the independence test?

In practice, given n i.i.d. samples $\{(y_1, z_1), \dots, (y_n, z_n)\}$ generated from (Y, Z) , one can build the statistic— $W(\gamma(\hat{\nu}_1, \hat{\nu}_2), \hat{\nu}_1 \otimes \hat{\nu}_2)$, where $\hat{\nu}$ denotes the corresponding empirical distribution—to test the independence. Computing $W(\gamma(\hat{\nu}_1, \hat{\nu}_2), \hat{\nu}_1 \otimes \hat{\nu}_2)$ is equivalent to solving the following optimization problem: (more details are presented in Section 5.)

$$\min_{X^\circ \in \Pi^\circ} \sum_{i,j,k=1}^n d((y_i, z_j), (y_k, z_k)) X_{ij;k}^\circ, \quad \Pi^\circ = \left\{ X_{ij;k}^\circ \geq 0 \mid \sum_{k=1}^n X_{ij;k}^\circ = \frac{1}{n^2}, \sum_{i,j=1}^n X_{ij;k}^\circ = \frac{1}{n} \right\}, \quad (1)$$

where the metric d is usually chosen as $d((y_i, z_j), (y_k, z_l)) = \|y_i - y_k\|_p + \|z_j - z_l\|_p$, and $\|\cdot\|_p$ denotes the l_p norm.

Problem (1) is an OT problem involving two marginals. One marginal is uniform with n atoms (i.e., we have $\sum_{i,j=1}^n X_{ij;k}^\circ = 1/n, \forall k, 1 \leq k \leq n$), and the other marginal is uniform with n^2 atoms (i.e., we have $\sum_{k=1}^n X_{ij;k}^\circ = 1/n^2, \forall i, j, 1 \leq i, j \leq n$). Motivated by this structure, we study the following special OT problem:

$$\min_{X' \in \mathcal{U}'} \sum_{i=1}^m \sum_{j=1}^n X'_{ij} C_{ij}, \quad \mathcal{U}' = \left\{ X'_{ij} \geq 0 \mid \sum_{j=1}^n X'_{ij} = \frac{1}{m}, \sum_{i=1}^m X'_{ij} = \frac{m_j}{m}, \sum_{j=1}^n m_j = m \right\}. \quad (2)$$

where $0 < n \leq m$, m_j 's are positive integers. One marginal of this OT problem is n -dimensional where the probability of each component is prescribed as m_j/m (i.e., we have $\sum_{i=1}^m X'_{ij} = m_j/m, \forall j, 1 \leq j \leq n$), and the other marginal is uniform with m atoms (i.e., we have $\sum_{j=1}^n X'_{ij} = 1/m, \forall i, 1 \leq i \leq m$). In essence, problem (1) is a special case of problem (2), where $m_j = n, m = n^2$.

Problem (2) is a linear programming (LP) problem with $\mathcal{O}(mn)$ variables and $\mathcal{O}(m+n)$ constraints, and then could be solved *exactly* by LP solvers. The well-known simplex method has exponential worst-case complexity (Klee & Minty, 1972). To the best of our knowledge, there are no known polynomial-time algorithms to get *exact* solutions to LP with *real*-valued inputs. Throughout this paper, we consider real-valued entries in the cost matrix and later propose a strongly polynomial-time algorithm to solve problem (2) precisely.

Per Birkhoff's theorem (Birkhoff, 1946), the solution to problem (2) is a vertex (whose coordinates are zeros and ones). Thus, problem (2) is equivalent to the following optimization problem: (the proof is relegated to the Appendix.)

$$\min_{X \in \mathcal{U}} \sum_{i=1}^m \sum_{j=1}^n \frac{1}{m} X_{ij} C_{ij}, \quad \mathcal{U} = \left\{ X_{ij} \in \{0, 1\} \mid \sum_{j=1}^n X_{ij} = 1, \sum_{i=1}^m X_{ij} = m_j, \sum_{j=1}^n m_j = m \right\}. \quad (3)$$

One may recall the assignment problem, where the permutation matrix is the solution matrix. $X \in \mathcal{U}$ is similar but different from the permutation matrix: $X \in \mathcal{U}$ is an $m \times n$ matrix instead of a square matrix and has *multiple* entries of 1 in each column instead of only one entry. In this case, we are not able to directly apply algorithms for the assignment problem, such as the Hungarian algorithm (Munkres, 1957). An approach to obtain the precise solution to problem (3) is first to duplicate the columns of C and X , then apply the Hungarian algorithm. The computational complexity of this approach is $\mathcal{O}(m^3)$. In this paper, a *modified Hungarian algorithm* is proposed. The algorithm specializes in solving the special type of OT problem (3), which is equivalent to problem (2), with a provable lower order— $\mathcal{O}(m^2n)$.

Back to the Wasserstein-distance-based independence test problem (1), the resulting computational complexity of applying the proposed algorithm is $\mathcal{O}(n^5)$ while the order of applying the classic Hungarian algorithm is $\mathcal{O}(n^6)$. In this sense, the proposed algorithm is faster.

In addition to the application in the Wasserstein independence test, broader applications of the modified Hungarian algorithm, including solving the one-to-many assignment problem and the many-to-many assignment problem (Zhu et al., 2011; 2016), are investigated. Two practical assignment problems involving the soccer game and agent-task assignment serve as examples to illustrate how to apply the proposed algorithm.

1.1 Related work:

Approximation algorithms: Our modified Hungarian algorithm is an exact OT solver. There are also a bunch of approximation algorithms (Cuturi, 2013; Altschuler et al., 2017; Dvurechensky et al., 2018; Lin et al., 2019; Guo et al., 2020; Xie et al., 2022). Notably, precise solutions are needed in some scenarios, and Dong et al. (2020) demonstrates the favorable numerical performance of the exact solutions over the approximate solutions. Therefore, the development of efficient exact algorithms is of much interest. Numerical experiments are conducted to compare the modified Hungarian algorithm with the most widely-used approximation algorithm—the Sinkhorn algorithm, highlighting the efficiency of our exact algorithm.

Independence criteria: There are some other independence criteria based on OT or the Wasserstein distance (Shi et al., 2020; Deb & Sen, 2021; Liu et al., 2022). We mainly focus on the formulation (1).

1.2 Our contributions:

We propose a modified Hungarian algorithm to solve a special type of OT problem (2). The modification enables us to deal with the scenario where two marginals have different sizes of atoms, and the atoms in one of the marginals have multiple assignments. Further, the proposed modified Hungarian algorithm could be extended to solve more general OT problems. Moreover, the applications of the proposed algorithm are explored: adopting the modified Hungarian algorithm to solve the Wasserstein independence test problem (1), the one-to-many assignment problem and the many-to-many assignment problem. Finally, several numerical experiments are carried out to show the favorability of our algorithm over both the classic Hungarian algorithm and the Sinkhorn algorithm.

1.3 Organization

The remainder of this paper is organized as follows. In Section 2, we introduce some basics of graph theory. In Section 3, we propose the modified Hungarian algorithm and compute its computational complexity. In Section 4, we explain how to generalize the proposed algorithm to solve a wider range of OT problems. In Section 5, we apply the modified Hungarian algorithm to the Wasserstein-distance-based independence test problem. In Section 6, we apply the modified Hungarian algorithm to the one-to-many assignment problem and the many-to-many assignment problem. In Section 7, we carry out various numerical experiments on both synthetic data and real data to validate our theoretical results and show the favorability of our algorithm. We discuss some future work in Section 8.

2 Preliminaries

Some definitions related to combinatorial optimization and graph theory (Suri, 2006; Burkard et al., 2012) are introduced. They will be needed in the rest of this paper.

Definition 1 (Assignment problem). *Given an $k \times k$ cost matrix with components $c_{ij} \geq 0, i, j \in [k] = \{1, \dots, k\}$, the assignment problem is to solve $\min_{\phi} \sum_{i=1}^k c_{i\phi(i)}$, where ϕ is the permutation of set $[k]$.*

Definition 2 (Bipartite graph). *A graph $G = (V, E)$ is bipartite if there exists a partition $V = V_1 \cup V_2$ such that $V_1 \cap V_2 = \emptyset$ and $E \subset V_1 \times V_2$. W.L.O.G., we always assume $E = V_1 \times V_2$.*

Definition 3 (Matching and perfect matching). *$M \subset E$ in the bipartite graph $G = (V, E)$ is a matching if every node of G coincides with at most one edge of M . W.L.O.G., it is assumed that the number of nodes in V_1 is the same as V_2 . In this case, if every node of G coincides with exactly an edge of M , M is called a perfect matching*

Definition 4 (Weighted bipartite graph). *A bipartite graph $G = (V, E)$ is a weighted bipartite graph if there is a weight $w(\cdot) \geq 0$ for each edge $e \in E$. The weight of a matching M is defined as $\sum_{e \in M} w(e)$.*

Definition 5 (Labeling and feasible labeling). *For a weighted bipartite graph $G = (V, E)$, a labeling is a function $l : V \rightarrow \mathbb{R}$. A feasible labeling is one labeling such that $l(v_1) + l(v_2) \geq w(v_1, v_2), \forall v_1 \in V_1, v_2 \in V_2$.*

Definition 6 (Equality graph and neighbor). *The equality graph w.r.t. labeling l is $G' = (V, E_l)$ where $E_l = \{(v_1, v_2) : l(v_1) + l(v_2) = w(v_1, v_2)\}$. Define the neighbor of $v_2 \in V_2$ and $S \subset V_2$ as $N_l(v_2) = \{v_1 : (v_1, v_2) \in E_l\}$ and $N_l(S) = \cup_{v_2 \in S} N_l(v_2)$, respectively.*

Definition 7 (Alternating and augmenting path). *Let M be a matching of the bipartite graph $G = (V, E)$. A path is alternating if its edges alternate between M and $E - M$. An alternating path is augmenting if both endpoints do not coincide with any edges in M .*

3 Modified Hungarian algorithm

In this section, we propose a modified Hungarian algorithm to solve the special type of OT problem (2), which is equivalent to problem (3).



Figure 1: pseudo-matching (left) and perfect pseudo-matching (right), where $n = 3, m = 9, m_1 = 2, m_2 = 3, m_3 = 4$.

3.1 Review

We first review the Hungarian algorithm (Munkres, 1957). Recall that the assignment problem is to solve $\min_{\phi} \sum_{i=1}^m c_{i\phi(i)}$. If we negate the costs and add the maximum of the costs to each component, solving the assignment problem is equivalent to finding a maximum weighted matching in the weighted bipartite graph with weights $w(i, j) = \max_{i,j} c_{ij} - c_{ij}$. The Kuhn-Munkres theorem (Munkres, 1957) shows that finding a maximum weighted matching is equivalent to finding a perfect matching on the equality graph associated with some feasible labeling in the bipartite graph. In this regard, the Hungarian algorithm solves the assignment problem by identifying a perfect matching on some equality graph in the weighted bipartite graph.

3.2 Pseudo-matching

In problem (3), $X \in \mathcal{U}$ has one entry of 1 in each row, multiple entries of 1 in each column, and 0's elsewhere. Since a permutation matrix corresponds to a (perfect) matching in the bipartite graph, we define ‘pseudo-matching’ in the bipartite graph $G = (V_1 \cup V_2, E)$ to describe X . V_1 has m nodes representing the rows of X while V_2 has n nodes representing the columns of X . Notice that we usually have $m > n$. In this case, each node in V_1 coincides with at most one edge, while multiple edges are allowed to connect with nodes in V_2 . See the formal definition in Definition 8.

Definition 8 (pseudo-matching, perfect pseudo-matching). *In the bipartite graph G , where $|V_1| = m, |V_2| = n$. $PM \subset E$ is a pseudo-matching if every node of V_1 coincides with at most one edge of PM , and j th node of V_2 coincides with at most m_j edges of PM , where $\sum_{j=1}^n m_j = m$. Furthermore, if every node of V_1 coincides with exactly one edge of PM and j th node of V_2 coincides with exactly m_j edges of PM , PM is called a perfect pseudo-matching.*

Figure 1 is an example of (perfect) pseudo-matching, where $n = 3, m = 9, m_1 = 2, m_2 = 3, m_3 = 4$. Under this setting, each node in the left-hand side of the graph can coincide with at most one edge, while each node in right-hand side can coincide with at most 2 edges, 3 edges, and 4 edges, respectively.

3.3 Our algorithm

Solving problem (3) is equivalent to looking for a maximum weighted pseudo-matching in the bipartite graph. We develop a modified Kuhn-Munkres theorem based on the pseudo-matching. See Theorem 1. (The proof can be found in the Appendix.) It demonstrates that we only need to find a perfect pseudo-matching on some equality graph to solve problem (3).

Theorem 1 (Modified Kuhn-Munkres theorem). *If l is a feasible labeling on the weighted bipartite graph $G = (V, E)$, and $PM \subset E_l$ is a perfect pseudo-matching on the corresponding equality graph $G' = (V, E_l)$, PM is a maximum weighted pseudo-matching.*

Equipped with the modified Kuhn-Munkres theorem, we design a modified Hungarian algorithm (Algorithm 1). The definitions used in the algorithm are specified in Definition 9. The modified Hungarian algorithm improves either the feasible labeling (adding edges to the associated equality graph) or the pseudo-matching until the pseudo-matching is perfect on some equality graph w.r.t. some feasible labeling. The algorithm

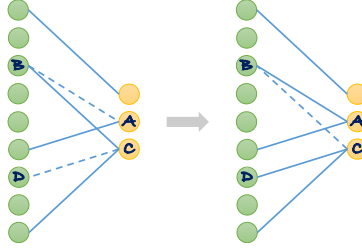


Figure 2: pseudo-augmenting process, where $n = 3, m = 9, m_1 = 2, m_2 = 3, m_3 = 4$.

improves the pseudo-matching by generating pseudo-augmenting paths and then exchanging the edge status along the paths. This process is called the pseudo-augmenting process. Also, we force the pseudo-augmenting paths emanating from V_2 , which have a lower order of nodes.

Definition 9 (Free, matched, pseudo-matched, pseudo-alternating path, pseudo-augmenting path). *Let PM be a pseudo-matching of $G = (V, E)$.*

- *If the node v is in V_1 , it is pseudo-matched if it is an endpoint of some edge in PM ; if the node v is the j th node in V_2 , it is pseudo-matched if it is an endpoint of m_j edges in PM . Otherwise, the node is free.*
- *If the node $v \in V$, we say it is matched if it is an endpoint of some edge in PM .*
- *A path is pseudo-alternating if its edge alternates between PM and $E - PM$. A pseudo-alternating path is pseudo-augmenting if both its endpoints are free.*

An example of the pseudo-augmenting process is given in Figure 2. The solid line means that the edge belongs to the pseudo-matching. The dashed line means that the edge belongs to the equality graph but does not belong to the pseudo-matching. Node B and node C are pseudo-matched. Edge A-B and edge C-D are not in the pseudo-matching. In this sense, A-B-C-D is a pseudo-alternating path. Because node A and node D are free, A-B-C-D is a pseudo-augmenting. The pseudo-augmenting process is to exchange the status of the edges: delete B-C from the pseudo-matching and enter A-B, C-D into the pseudo-matching. The pseudo-matching has been improved in this way.

3.4 Computational complexity

We now analyze the computational complexity of Algorithm 1. Similar to the Hungarian algorithm (Suri, 2006), we keep track of $slack_{v_1} = \min_{v_2 \in S} \{l(v_1) + l(v_2) - w(v_1, v_2)\}$, $\forall v_1 \notin T$. The computational cost increases when computing α_l via *slacks*, updating the values of *slacks*, and calculating the labeling.

The number of edges of the pseudo-matching increases by 1 after one loop, so $\mathcal{O}(m)$ loops is needed to form a perfect pseudo-matching. There are two subroutines in each loop: the first is to update the feasible labeling (Step 2), and the second is to improve the pseudo-matching (Step 3). In the procedure of updating the feasible labeling, since there are n nodes in V_2 , the improvement occurs $\mathcal{O}(n)$ times to build a pseudo-alternating tree. In each time, computing α_l , updating the *slacks*, and calculating the labeling cost $\mathcal{O}(m)$. In the procedure of improving the pseudo-matching, when a new node has been added to S , it costs $\mathcal{O}(m)$ to update *slacks*, and $\mathcal{O}(n)$ nodes could be added. On the other hand, when a node has been added to T , we just remove the corresponding $slack_{v_1}$. We conclude that each loop costs $\mathcal{O}(mn)$, so the total computational complexity of Algorithm 1 to solve problem (3) is $\mathcal{O}(m^2n)$. We summarize the analysis above in Theorem 2.

Theorem 2. *The computational complexity of applying the modified Hungarian algorithm to solve problem (3) is $\mathcal{O}(m^2n)$.*

Note that the adoption of the Hungarian algorithm has an order of $\mathcal{O}(m^3)$. Hence, the proposed modified Hungarian algorithm will outperform, especially when $m \gg n$.

Algorithm 1: Modified Hungarian Algorithm

Generate an initial feasible labeling l : $\forall v_2 \in V_2, l(v_2) = 0; \forall v_1 \in V_1, l(v_1) = \max_{v_2 \in V_2} \{w(v_1, v_2)\}$ and initialize a pseudo-matching M in E_l ;

```

1 if  $M$  is a perfect pseudo-matching then
  | Stop
else
  | Pick up a free node  $v_{\text{free}} \in V_2$ . Set  $S = \{v_{\text{free}}\}, T = \emptyset$ ;
  | for  $\mathbf{v}_1 \in V_1$  is matched to  $v_{\text{free}}$  do
  |   |  $T = T \cup \mathbf{v}_1$ 
  | end
end
2 if  $N_l(S) - T = \emptyset$  then
  | update labeling such that forcing  $N_l(S) - T \neq \emptyset$ :
  |
  | 
$$\alpha_l = \min_{v_1 \notin T, v_2 \in S} \{l(v_1) + l(v_2) - w(v_1, v_2)\}, \quad l(v) = \begin{cases} l(v) - \alpha_l & v \in S \\ l(v) + \alpha_l, & v \in T \\ l(v), & \text{otherwise} \end{cases}.$$

  |
end
3 if  $N_l(S) - T \neq \emptyset$  then
  | pick  $v_1 \in N_l(S) - T$ ;
  | if  $v_1$  is free then
  |   |  $v_{\text{free}} \rightarrow v_1$  is a pseudo-augmenting path. Pseudo-augment the pseudo-matching  $M$ . Go to Step 1;
  | end
  | if  $v_1$  is pseudo-matched to  $z$  then
  |   | extend the pseudo-matching tree:  $S = S \cup \{z\}, T = T \cup \{v_1\}$ ;
  |   | for  $\mathbf{v}_1 \in V_1$  is matched to  $z$  do
  |   |   |  $T = T \cup \mathbf{v}_1$ ;
  |   | end
  | end
  | Go to Step 2.
end

```

4 Generalization

In this section, we discuss how to adopt the modified Hungarian algorithm to solve a class of more general OT problems.

For the special type of OT problem we discuss earlier, we require that one marginal of the OT problem should be uniform. The next step is to relax the ‘uniform’ requirement:

What if each component probability of two marginals is prescribed as n_i/M and m_j/M , respectively?

In other words, we are interested in solving the following optimization problem:

$$\min_{X^* \in \mathcal{U}^*} \sum_{i=1}^m \sum_{j=1}^n X_{ij}^* C_{ij}, \quad \mathcal{U}^* = \left\{ X_{ij}^* \geq 0 \mid \sum_{j=1}^n X_{ij}^* = \frac{n_i}{M}, \sum_{i=1}^m X_{ij}^* = \frac{m_j}{M}, \sum_{j=1}^n m_j = \sum_{i=1}^m n_i = M \right\}, \quad (4)$$

where $0 < n \leq m$, n_i ’s, m_j ’s are positive integers. One possible method is first to rewrite problem (4) as the formulation of the special type of OT problem (2), seeing Proposition 1 (the proof is relegated to the Appendix).

Proposition 1. *Problem (4) is equivalent to the following optimization problem:*

$$\min_{X^\dagger \in \mathcal{U}^\dagger} \sum_{i=1}^M \sum_{j=1}^n X_{ij}^\dagger C_{ij}^\dagger, \quad \mathcal{U}^\dagger = \left\{ X_{ij}^\dagger \geq 0 \mid \sum_{j=1}^n X_{ij}^\dagger = \frac{1}{M}, \sum_{i=1}^M X_{ij}^\dagger = \frac{m_j}{M} \right\}. \quad (5)$$

where C^\dagger is an $M \times n$ matrix generated by duplicating the i th row of C n_i times:

$$C_{tj}^\dagger = \begin{cases} C_{1j} & 1 \leq t \leq n_1 \\ C_{ij} & n_1 + \dots + n_{i-1} + 1 \leq t \leq n_1 + \dots + n_i, 2 \leq i \leq m \end{cases}.$$

Problem (5) belongs to the special type of OT problem (2). We could apply the proposed modified Hungarian algorithm to problem (5) and get the exact solution to problem (4). The resulting computational complexity is $\mathcal{O}(M^2n)$.

5 Application to the independence test using the Wasserstein distance

In this section, we apply the modified Hungarian algorithm to the Wasserstein-distance-based independence test, which originally motivates us to study the special type of OT problem (2).

Suppose that there are n i.i.d. samples $\{(y_1, z_1), \dots, (y_n, z_n)\}$, where $(y_i, z_i) \sim (Y, Z)$, $Y \sim \nu_1$, $Z \sim \nu_2$. One could prove the following equivalence:

$$Y \perp Z \iff \nu_1 \otimes \nu_2 = \gamma(\nu_1, \nu_2) \iff W(\nu_1 \otimes \nu_2, \gamma(\nu_1, \nu_2)) = 0,$$

which follows from the fact that the Wasserstein distance is a valid metric between probability measures. Given the empirical data, we utilize the statistic $W(\gamma(\hat{\nu}_1, \hat{\nu}_2), \hat{\nu}_1 \otimes \hat{\nu}_2)$ to test the independence between Y and Z , where $\hat{\nu}$ denotes the empirical distribution and has the following expressions:

$$\gamma(\hat{\nu}_1, \hat{\nu}_2) = \begin{pmatrix} \frac{1}{n} & & 0 \\ & \dots & \\ 0 & & \frac{1}{n} \end{pmatrix}, \quad \hat{\nu}_1 \otimes \hat{\nu}_2 = \begin{pmatrix} \frac{1}{n^2} & \dots & \frac{1}{n^2} \\ \dots & \dots & \dots \\ \frac{1}{n^2} & \dots & \frac{1}{n^2} \end{pmatrix}.$$

Plug in the Wasserstein distance formula, the resulting optimization problem is:

$$\min_{X \in \Pi} \sum_{i,j,k,l=1}^n d((y_i, z_j), (y_k, z_l)) X_{ij;kl}, \quad (6)$$

where

$$\Pi = \left\{ X_{ij;kl} \geq 0 \mid \sum_{k,l=1}^n X_{ij;kl} = \frac{1}{n^2}, \sum_{i,j=1}^n X_{ij;kl} = \begin{cases} \frac{1}{n} & k = l \\ 0 & k \neq l \end{cases}, \forall i, j, k, l = 1, \dots, n. \right\}.$$

It is worth noting that $X_{ij;kl} = 0, k \neq l$. If we let $X_{ij;k}^\circ := \sum_{l=1}^n X_{ij;kl}$, problem (6) can be simplified as problem (1). Problem (1) belongs to the special type of OT problem, where $m_j = n, \forall j, 1 \leq j \leq n, m = n^2$. Adopting the Hungarian algorithm to problem (1) costs $\mathcal{O}(n^6)$, while adopting the proposed Hungarian algorithm directly costs $\mathcal{O}(n^5)$.

6 Application to the one-to-many assignment problem and the many-to-many assignment problem

In this section, we proceed to explain how to apply the modified Hungarian algorithm to solve the one-to-many assignment problem and the many-to-many assignment problem. The applications are shown based on two practical examples.

Example 1 (one-to-many assignment problem): An assignment problem involving the soccer ball game mentioned by Zhu et al. (2011) is considered here. Suppose a coach is tasked to choose players from a soccer team with m_1 players (a_1, \dots, a_{m_1}) . There are 4 roles (r_1 goalkeeper, r_2 backs, r_3 midfielders, r_4 forwards). It is assumed that $m_1 > 4$. Suppose each player's performance evaluation of each role is known. The overall performance evaluation of the team is the sum of each selected player's performance evaluation

of its assigned role. The optimal strategy is to maximize the overall performance evaluation of the team. The coach should solve the following optimization problem:

$$\max_{A \in \mathcal{A}} \sum_{i=1}^{m_1} \sum_{j=1}^4 A_{ij} P_{ij}, \quad \mathcal{A} = \left\{ A_{ij} = \{0, 1\} \mid \sum_{j=1}^4 A_{ij} \leq 1, \sum_{i=1}^{m_1} A_{ij} = r_j \right\},$$

where $P_{ij} \geq 0$ denotes player a_i 's performance evaluation of role j , $A_{ij} = 1$ means player a_i is selected as role j while $A_{ij} = 0$ means the player is not selected as role j .

If $m_1 = \sum_{j=1}^4 r_j$, the optimization problem above belongs to the special type of OT problem (3), where $n = 4, m = \sum_{j=1}^4 r_j$. The modified Hungarian algorithm could be applied to find the optimal strategy, and the resulting computational complexity is $\mathcal{O}(4m_1^2)$.

If $m_1 > \sum_{j=1}^4 r_j$, we can't apply the modified Hungarian algorithm directly. To make the problem tractable, we create one more role, and each player's performance evaluation of this role is 0. Players who are not selected are 'assigned' to this role by default. In this scenario, our goal is to solve the following optimization problem:

$$\max_{A^\dagger \in \mathcal{A}^\dagger} \sum_{i=1}^{m_1} \sum_{j=1}^5 A_{ij}^\dagger P_{ij}^\dagger, \quad \mathcal{A}^\dagger = \left\{ A_{ij}^\dagger = \{0, 1\} \mid \sum_{j=1}^5 A_{ij}^\dagger = 1, \sum_{i=1}^{m_1} A_{ij}^\dagger = \begin{cases} r_j & 1 \leq j \leq 4 \\ m_1 - \sum_{j=1}^4 r_j & j = 5 \end{cases} \right\},$$

where we append P by adding one more column of zeros to get P^\dagger . It belongs to the special type of OT problem, where $n = 5, m = m_1$. Then, we could apply the modified Hungarian algorithm to solve the problem, and the resulting computational complexity is $\mathcal{O}(5m_1^2)$.

Note that the computational order of applying the algorithm developed by Zhu et al. (2011) is $\mathcal{O}(m_1^3)$, which is worse than the proposed modified Hungarian algorithm.

Example 2 (many-to-many assignment problem): The following example is an agent-task assignment problem mentioned by Zhu et al. (2016). Assume there are m_2 tasks (t_1, \dots, t_{m_2}) and n_1 agents (a_1, \dots, a_{n_1}) in total. It is assumed that $n_1 < m_2$. Each task should be undertaken by many agents, and each agent can perform many tasks. To be more specific, task t_i must be assigned to l_i agents, agent a_j can perform at most s_j tasks. Suppose the performance evaluation of each agent performing each task is known. The optimal assignment plan is to maximize the overall performance. The resulting optimization problem is as follows:

$$\max_{A' \in \mathcal{A}'} \sum_{i=1}^{m_2} \sum_{j=1}^{n_1} A'_{ij} P'_{ij}, \quad \mathcal{A}' = \left\{ A'_{ij} = \{0, 1\} \mid \sum_{j=1}^{n_1} A'_{ij} = l_i, \sum_{i=1}^{m_2} A'_{ij} \leq s_j \right\},$$

where $P'_{ij} \geq 0$ denotes agent a_j 's performance evaluation on task t_i , $A'_{ij} = 1$ means that agent a_j is assigned to perform task t_i while $A'_{ij} = 0$ means that agent a_j is not assigned to perform task t_i .

If $\sum_{i=1}^{m_2} l_i = \sum_{j=1}^{n_1} s_j$, the optimization problem follows the formulation of the problem in Section 4, where $n = n_1, M = \sum_{j=1}^{n_1} s_j$. We could apply the modified Hungarian algorithm to find the optimal assignment plan, and the resulting computational complexity is $\mathcal{O}(n_1(\sum_{j=1}^{n_1} s_j)^2)$.

If $\sum_{i=1}^{m_2} l_i < \sum_{j=1}^{n_1} s_j$, we create one more task which must be performed by $(\sum_{j=1}^{n_1} s_j - \sum_{i=1}^{m_2} l_i)$ agents, and each agent's performance of this new task equals 0. This reformulation promises that each agent performs the maximum amount of tasks. Accordingly, we need to solve the following optimization problem:

$$\max_{A^\ddagger \in \mathcal{A}^\ddagger} \sum_{i=1}^{m_2+1} \sum_{j=1}^{n_1} A_{ij}^\ddagger P_{ij}^\ddagger, \quad \mathcal{A}^\ddagger = \left\{ A_{ij}^\ddagger = \{0, 1\} \mid \sum_{j=1}^{n_1} A_{ij}^\ddagger = \begin{cases} l_i & 1 \leq i \leq m_2 \\ \sum_{j=1}^{n_1} s_j - \sum_{i=1}^{m_2} l_i & j = m_2 + 1 \end{cases}, \sum_{i=1}^{m_2+1} A_{ij}^\ddagger = s_j \right\},$$

where we append P' by adding one more row of zeros to get P^\ddagger . It follows the formulation of problem (4), where $n = n_1, M = \sum_{j=1}^{n_1} s_j$. We could adopt the method introduced in Section 4, and the resulting computation complexity is $\mathcal{O}(n_1(\sum_{j=1}^{n_1} s_j)^2)$.

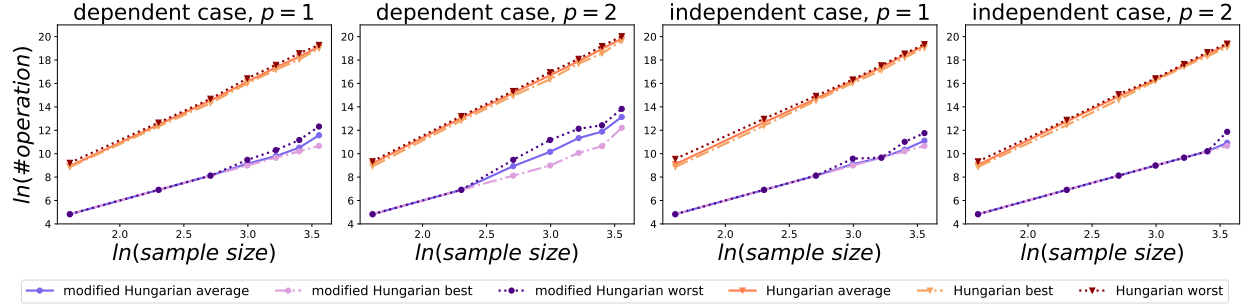


Figure 3: Comparison with the Hungarian algorithm on synthetic data

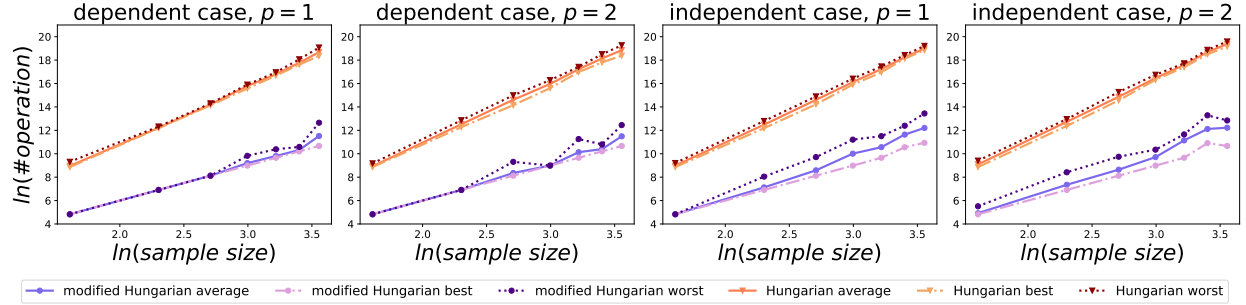


Figure 4: Comparison with the Hungarian algorithm on CIFAR10

Note that the computational order of applying the algorithm developed by Zhu et al. (2016) is $\mathcal{O}((\sum_{j=1}^{n_1} s_j)^3)$, which has a higher computational burden than our proposed method.

7 Numerical experiments

In this section, we carry out experiments on the Wasserstein independence test problem on a synthetic dataset, CIFAR10¹ (Krizhevsky et al., 2009) and Wisconsin breast cancer dataset² (Dua & Graff, 2017). We compare the proposed modified Hungarian algorithm with the exact algorithm—Hungarian algorithm, and the approximation algorithm—Sinkhorn algorithm. The numerical results validate the computational complexity of the modified Hungarian algorithm, and show the favorability of applying the proposed algorithm over the Hungarian algorithm and Sinkhorn algorithm. The performance is evaluated by the number of *numerical operations* here. We relegate the results of the *running time* to the Appendix.

7.1 Experiment setting

Our algorithm is adaptive to any metric. The foregoing experiments are based on the l_p norm-based metric: $d((x_i, y_j), (x_k, y_l)) = \|x_i - x_k\|_p + \|y_j - y_l\|_p$. More specifically, we examine how the modified Hungarian algorithm, the Hungarian algorithm and the Sinkhorn algorithm perform when $p = 1$ and $p = 2$. We create one dependent case and one independent case with different sample sizes for each dataset and run the algorithms on each case 10 times. We plot the worst, best and average number of numerical operations for each case.

Synthetic data: Suppose that there are independent variables $X \sim N(5\mathbf{1}_{10}, 30I_{10})$, where $\mathbf{1}_{10}$ is a 10-dimensional vector with all ones and I_{10} is the identity matrix; and $Y = (Y_1, \dots, Y_{25})^T$, where Y_i 's are independent and follow $\text{Unif}(10, 20)$. We calculate the empirical Wasserstein distance in (1) *independent*

¹<https://www.cs.toronto.edu/~kriz/cifar.html>

²[https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+\(Diagnostic\)](https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+(Diagnostic))

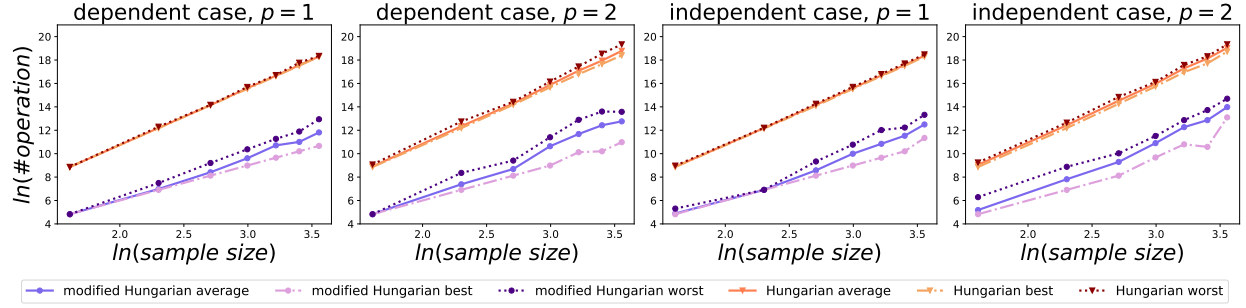


Figure 5: Comparison with the Hungarian algorithm on Wisconsin breast cancer data

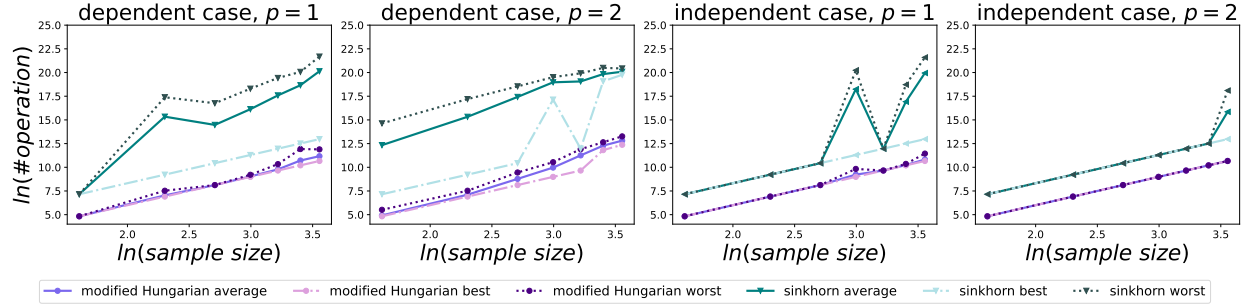


Figure 6: Comparison with the Sinkhorn algorithm on synthetic data

case: between X and Y ; (2) *dependent case*: between X and Z (where $Z = X_1 + Y_1$, X_1 is the first 5 coordinates of X , Y_1 is the first 5 coordinates of Y).

CIFAR10: Each image in CIFAR10 contains 32×32 pixels, and each pixel is composed of 3 color channels. Each image is essentially a 3072-dimensional vector. Then, we rescale the vector components to $[0, 1]$. Suppose $X \in \mathbb{R}^{3072}$ is the distribution generated uniformly from the images of classes: airplane, automobile, bird, cat, and deer; $Y \in \mathbb{R}^{3072}$ is the distribution generated uniformly from the images of other five classes. We calculate the empirical Wasserstein distance in (1) *independent case*: between X and Y_1 (where Y_1 is the first 1536 coordinates of Y); (2) *dependent case*: between X and Z (where $Z = X_2/2 + Y_1/2$, X_2 is the last 1536 coordinates of X , Y_1 is the first 1536 coordinates of Y).

Breast cancer data: There are 569 instances, and each instance possesses 30 features. Each instance is a 30-dimensional vector, and we rescale the components to $[0, 1]$. There are two classes of instances: benign and malignant. Let $X \in \mathbb{R}^{30}$ be the distribution generated uniformly from the benign class, and $Y \in \mathbb{R}^{30}$ be the distribution generated uniformly from the malignant class. We calculate empirical Wasserstein distance in (1) *independent case*: between X_1 and Y_2 (where X_1 is the first 5 coordinates of X , Y_2 the last 25 coordinates of Y); (2) *dependent case*: between X and Z (where $Z = X_1 * Y_1$, X_1 is the first 5 coordinates of X , Y_1 is the first 5 coordinates of Y , $*$ means the coordinate-wise product).

7.2 Comparison with the Hungarian algorithm

We compare the modified Hungarian algorithm with the classic Hungarian algorithm. The results are presented in Figure 3, 4, 5. The figures illustrate that the proposed algorithm gains a factor n in computational complexity when solving the proposed special type of OT problem. To be more specific, notice that the slope of $\ln(\text{number of numerical operations})$ over $\ln(\text{sample size})$ indicates the order of the associated algorithm, and the slope of our algorithm is around 5 while the slope of the Hungarian algorithm is around 6. This observation implies that the order of applying our algorithm is $\mathcal{O}(n^5)$ while the order of applying the Hungarian algorithm is $\mathcal{O}(n^6)$. Such observations are consistent with our theoretical results.

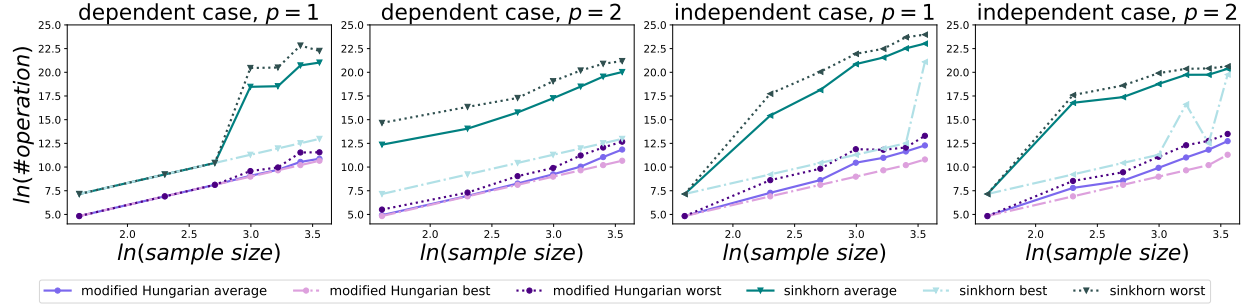


Figure 7: Comparison with the Sinkhorn algorithm on CIFAR10

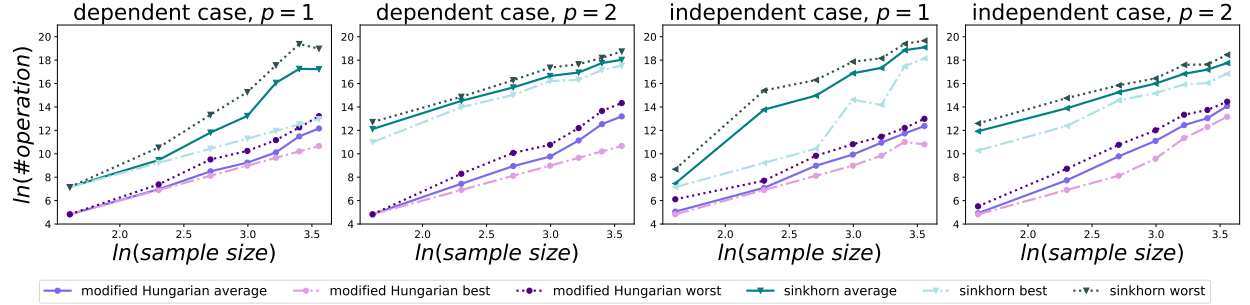


Figure 8: Comparison with the Sinkhorn algorithm on Wisconsin breast cancer data

7.3 Comparison with the Sinkhorn algorithm

We compare the modified Hungarian algorithm with the Sinkhorn algorithm. Among the state-of-the-art approximation solvers for OT problems, the first-order approximation algorithms (Dvurechensky et al., 2018; Lin et al., 2019; Guo et al., 2020) are mainly employed to solve the balanced case ($m = n$). The Sinkhorn algorithm could deal with the unbalanced scenario ($m \neq n$) and is widely used in all kinds of OT-related models. Therefore, we choose the Sinkhorn algorithm as the baseline and then investigate the performance of the Sinkhorn algorithm in the Wasserstein-distance-based independence test problem. When we implement the Sinkhorn algorithm, we set the regularization parameter as 0.1 and the accuracy as 0.0001.

The results are presented in Figure 6, 7, 8. One may observe that for almost all instances, the total number of numerical operations of the modified Hungarian algorithm is less than the Sinkhorn algorithm. Moreover, the performance of the modified Hungarian algorithm has a lower variance than the Sinkhorn algorithm. In conclusion, the modified Hungarian algorithm outperforms the Sinkhorn algorithm.

8 Discussion

A modified Hungarian algorithm is developed to efficiently solve a wide range of OT problems. Theoretical analysis and numerical experiments demonstrate that the proposed algorithm compares favorably with the Hungarian algorithm and the Sinkhorn algorithm. In addition to the computational aspects, broad applications are explored, including the Wasserstein-distance-based independence test, the one-to-many assignment problem and the many-to-many assignment problem. The many-to-many assignment problem closely relates to practical problems involving service assignment problems (Ng et al., 2008), sensor networks (Bhardwaj & Chandrakasan, 2002), and access control (Ahn & Hu, 2007). Future work along this line is to apply the proposed algorithm to problems involving engineering and control. Also, there is some possibility of applying the proposed algorithm to some unsupervised learning problems. For example, the clustering problem could be formulated as an OT problem (Genevay et al., 2019). Assume that there are n clusters and m samples in total, and each cluster has m_j samples. If we want to identify the cluster assignment to minimize the

‘distance’ between cluster ‘centers’ and the associated assigned samples, we are solving the special type of OT problem in this paper. The future work along this line may be to find a scheme to determine ‘distance’ and ‘centers’ to promise desirable model performances.

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A Appendix

A.1 Proof of proposition in Section 1

Proposition 2. *The optimization problem (2) is equivalent to the optimization problem (3).*

Proof. We first consider the following two optimization problems (7), (8):

$$\min_{X^1 \in \mathcal{U}^1} \sum_{i=1}^m \sum_{j=1}^m X_{ij}^1 C_{ij}^\dagger, \quad \mathcal{U}^1 = \left\{ X_{ij}^1 \geq 0 \mid \sum_{j=1}^m X_{ij}^1 = \frac{1}{m}, \sum_{i=1}^m X_{ij}^1 = \frac{1}{m} \right\}. \quad (7)$$

where C^\dagger is an $m \times m$ matrix generated by duplicating the j th column of C m_j times:

$$C_{it}^\dagger = \begin{cases} C_{i1} & 1 \leq t \leq m_1, \\ C_{ij} & m_1 + \dots + m_{j-1} + 1 \leq t \leq m_1 + \dots + m_j, 2 \leq j \leq n \end{cases}.$$

$$\min_{X^\dagger \in \mathcal{U}^\dagger} \sum_{i=1}^m \sum_{j=1}^m \frac{1}{m} X_{ij}^\dagger C_{ij}^\dagger, \quad \mathcal{U}^\dagger = \left\{ X_{ij}^\dagger \in \{0, 1\} \mid \sum_{j=1}^m X_{ij}^\dagger = 1, \sum_{i=1}^m X_{ij}^\dagger = 1 \right\}, \quad (8)$$

Then, we denote the objective functions of problems (2), (3), (7) and (8) by $f'(X')$, $f(X)$, $f^1(X^1)$, and $f^\dagger(X^\dagger)$, respectively.

Firstly, we prove (2) \iff (7).

On one hand, for any $X^1 \in \mathcal{U}^1$, if we let

$$X'_{ij} = \sum_{t=1}^{m_1} X^1_{it}, \quad j = 1,$$

$$X'_{ij} = \sum_{t=m_1+\dots+m_{j-1}+1}^{m_1+\dots+m_j} X^1_{it}, \quad 2 \leq j \leq n,$$

then we have

$$X'_{ij} \geq 0,$$

$$\sum_{j=1}^n X'_{ij} = \sum_{t=1}^{m_1} X^1_{it} + \sum_{t=m_1+\dots+m_{j-1}+1}^{m_1+\dots+m_j} X^1_{it} = \sum_{t=1}^m X^1_{it} = \frac{1}{m},$$

$$\sum_{i=1}^m X'_{ij} = \sum_{i=1}^m \sum_{t=1}^{m_1} X^1_{it} = \frac{m_1}{m}, \quad j = 1,$$

$$\sum_{i=1}^m X'_{ij} = \sum_{i=1}^m \sum_{t=m_1+\dots+m_{j-1}+1}^{m_1+\dots+m_j} X^1_{it} = \frac{m_j}{m}, \quad 2 \leq j \leq n.$$

Thus, $X' \in \mathcal{U}'$.

For the objective functions, we have the following:

$$f'(X') = \sum_{i=1}^m \sum_{j=1}^n X'_{ij} C_{ij} = \sum_{i=1}^m \left(\sum_{t=1}^{m_1} X^1_{it} C^1_{it} + \sum_{t=m_1+\dots+m_{j-1}+1}^{m_1+\dots+m_j} X^1_{it} C^1_{it} \right) = \sum_{i=1}^m \sum_{t=1}^m X^1_{it} C^1_{it} = f^1(X^1).$$

On the other hand, for any $X' \in \mathcal{U}$, if we let

$$X^1_{it} = \begin{cases} X'_{i1}/m_1 & 1 \leq t \leq m_1 \\ X'_{ij}/m_j & m_1 + \dots + m_{j-1} + 1 \leq t \leq m_1 + \dots + m_j, 2 \leq j \leq n, \end{cases}$$

then we have

$$X^1_{it} \geq 0,$$

$$\sum_{t=1}^m X^1_{it} = \sum_{j=1}^n \frac{X'_{ij}}{m_j} m_j = \sum_{j=1}^n X'_{ij} = \frac{1}{m},$$

$$\sum_{i=1}^m X^1_{it} = \sum_{i=1}^n \frac{X'_{ij}}{m_j} = \frac{1}{m_j} \sum_{i=1}^n X'_{ij} = \frac{1}{m}.$$

Thus, $X^1 \in \mathcal{U}^1$.

For the objective functions, we have the following:

$$f^1(X^1) = \sum_{i=1}^m \sum_{t=1}^m X^1_{it} C^1_{it} = \sum_{i=1}^m \sum_{j=1}^n \frac{X'_{ij}}{m_j} C_{it} m_j = f'(X').$$

Hence, (2) \iff (7).

By Birkhoff's theorem, we know (7) \iff (8). Therefore, we have (2) \iff (8).

Similarly, for any $X^\dagger \in \mathcal{U}^\dagger$, if we let

$$X_{ij} = \sum_{t=1}^{m_1} X^\dagger_{it}, \quad j = 1,$$

$$X_{ij} = \sum_{t=m_1+\dots+m_{j-1}+1}^{m_1+\dots+m_j} X_{it}^\dagger, \quad 2 \leq j \leq n,$$

then we have $X \in \mathcal{U}$ and $f^\dagger(X^\dagger) = f(X)$.

For any $X \in \mathcal{U}$, if we let

$$X_{it}^\dagger = \begin{cases} X_{i1}/m_1, & 1 \leq t \leq m_1 \\ X_{ij}/m_j & m_1 + \dots + m_{j-1} + 1 \leq t \leq m_1 + \dots + m_j, 2 \leq j \leq n \end{cases}$$

then we have $X^\dagger \in \mathcal{U}^\dagger$ and $f^\dagger(X^\dagger) = f(X)$.

Therefore, (3) \iff (8).

In conclusion, we have (3) \iff (8) \iff (2). □

A.2 Proof of Theorem 1

Proof. Denote the edge $e \in E$ by $e = (e_{v_1}, e_{v_2})$. Let PM' be any perfect pseudo-matching in G (not necessarily in the equality graph E_l). And $v_1^i, i = 1, \dots, m; v_2^j, j = 1, \dots, n$ are nodes from V_1 and V_2 , respectively. Since $v_1^i \in V_1$ is covered exactly once by PM' , and $v_2^j \in V_2$ is covered exactly m_j times by PM' , we have

$$w(PM') = \sum_{e \in PM'} w(e) \leq \sum_{e \in PM'} (l(e_{v_1}) + l(e_{v_2})) = \sum_{i=1}^m l(v_1^i) + \sum_{j=1}^n m_j l(v_2^j),$$

where the first inequality comes from the definition of feasible labeling.

Thus, $\sum_{i=1}^m l(v_1^i) + \sum_{j=1}^n m_j l(v_2^j)$ is the upper bound of the weight of any perfect pseudo-matching. Then let PM be a perfect pseudo-matching in the equality graph E_l , we have

$$w(PM) = \sum_{e \in PM} w(e) = \sum_{i=1}^m l(v_1^i) + \sum_{j=1}^n m_j l(v_2^j).$$

Hence $w(PM') \leq w(PM)$, and PM is the maximum weighted pseudo-matching. □

A.3 Proof of Proposition 1

Proof. We denote the objective functions of problems (4) and (5) by $g^*(X^*)$, and $g^\dagger(X^\dagger)$, respectively.

On one hand, for any $X^\dagger \in \mathcal{U}^\dagger$, if we let

$$X_{ij}^* = \sum_{t=1}^{n_1} X_{tj}^\dagger, \quad i = 1,$$

$$X_{ij}^* = \sum_{t=n_1+\dots+n_{i-1}+1}^{n_1+\dots+n_i} X_{tj}^\dagger, \quad 2 \leq i \leq m,$$

then we have

$$X_{ij}^* \geq 0,$$

$$\sum_{i=1}^m X_{ij}^* = \sum_{t=1}^{n_1} X_{tj}^\dagger + \sum_{t=n_1+\dots+n_{i-1}+1}^{n_1+\dots+n_i} X_{tj}^\dagger = \sum_{t=1}^M X_{tj}^\dagger = \frac{m_j}{M},$$

$$\sum_{j=1}^n X_{ij}^* = \sum_{j=1}^n \sum_{t=1}^{n_1} X_{tj}^\dagger = \frac{n_1}{M}, \quad i = 1,$$

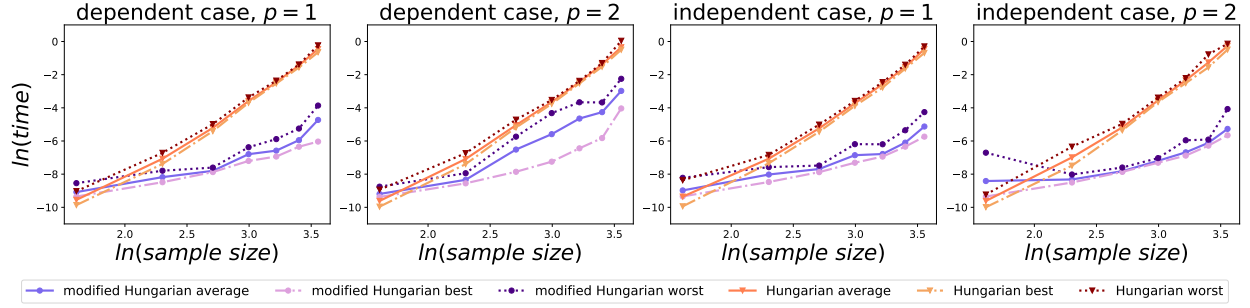


Figure 9: Comparison with the Hungarian algorithm on synthetic data w.r.t. time

$$\sum_{i=1}^m X_{ij}^* = \sum_{i=1}^m \sum_{t=n_1+\dots+n_{i-1}+1}^{n_1+\dots+n_i} X_{it}^\dagger = \frac{n_i}{m}, \quad 2 \leq i \leq m.$$

Thus, $X^* \in \mathcal{U}^*$.

For the objective function, we have the following:

$$g^*(X^*) = \sum_{i=1}^M \sum_{j=1}^n X_{ij}^* C_{ij} = \sum_{j=1}^n \left(\sum_{t=1}^{n_1} X_{tj}^\dagger C_{tj} + \sum_{t=n_1+\dots+n_{i-1}+1}^{n_1+\dots+n_i} X_{it}^\dagger C_{it} \right) = \sum_{j=1}^n \sum_{t=1}^M X_{it}^\dagger C_{it} = g^\dagger(X^\dagger).$$

On the other hand, for any $X^* \in \mathcal{U}^*$, if we let

$$X_{tj}^\dagger = \begin{cases} X_{1j}^*/n_1, & 1 \leq t \leq n_1 \\ X_{ij}^*/n_i & n_1 + \dots + n_{i-1} + 1 \leq t \leq n_1 + \dots + n_i, 2 \leq i \leq m \end{cases}$$

then we have

$$\begin{aligned} X_{ij}^\dagger &\geq 0, \\ \sum_{j=1}^n X_{ij}^\dagger &= \sum_{j=1}^n \frac{X_{ij}^*}{n_i} = \frac{n_i}{M}, \\ \sum_{i=1}^M X_{ij}^\dagger &= \sum_{i=1}^M \frac{X_{ij}^*}{n_i} n_i = \sum_{i=1}^m X_{ij}^* = \frac{m_j}{M}. \end{aligned}$$

Thus, $X^\dagger \in \mathcal{U}^\dagger$.

For the objective function, we have the following:

$$g^\dagger(X^\dagger) = \sum_{i=1}^M \sum_{j=1}^n X_{ij}^\dagger C_{ij} = \sum_{j=1}^n \sum_{i=1}^m \frac{X_{ij}^*}{n_i} C_{it} n_i = g^*(X^*).$$

Hence, (4) \iff (5). □

A.4 Experiment results w.r.t. running time

We run the Hungarian algorithm code from package ‘scipy’ in Python, modify the Sinkhorn algorithm code according to package ‘POT’, and implement the modified Hungarian algorithm. We record the running time. Figure 9, 10, 11, 12, 13, 14 are experiment results w.r.t. the running time.

When the sample size is small, our algorithm may be slower than the Hungarian algorithm. However, as the sample size increases, our algorithm becomes more efficient than the Hungarian algorithm. In some

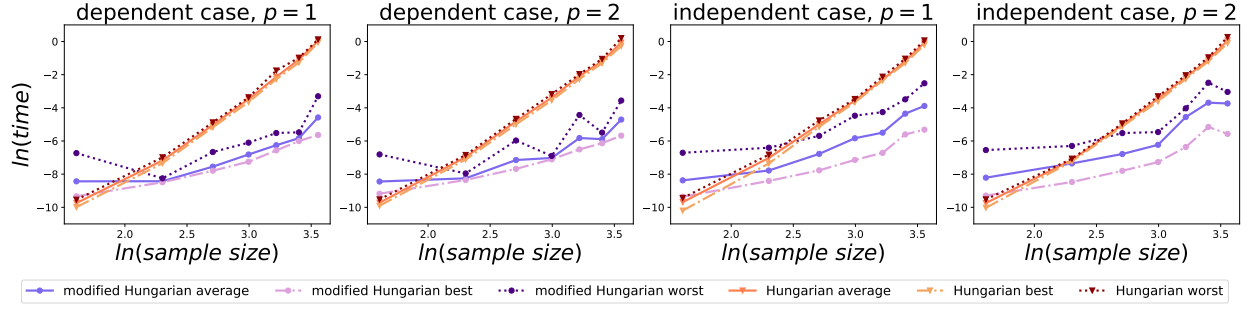


Figure 10: Comparison with the Hungarian algorithm on CIFAR10 w.r.t. time

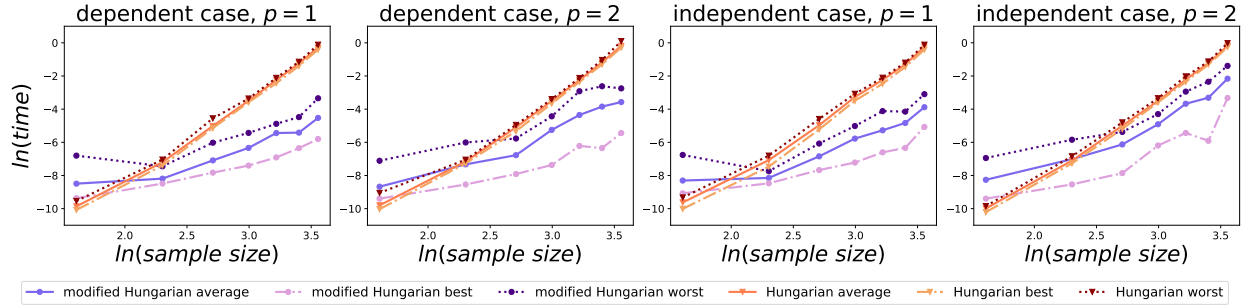


Figure 11: Comparison with the Hungarian algorithm on Wisconsin cancer data w.r.t. time

cases, the Sinkhorn algorithm may be faster than the modified Hungarian algorithm. However, the modified Hungarian algorithm still outperforms the Sinkhorn algorithm on average w.r.t. running time.

In theory, our proposed algorithm will compare more favorably if all algorithms are implemented in a low-level language, such as C++.

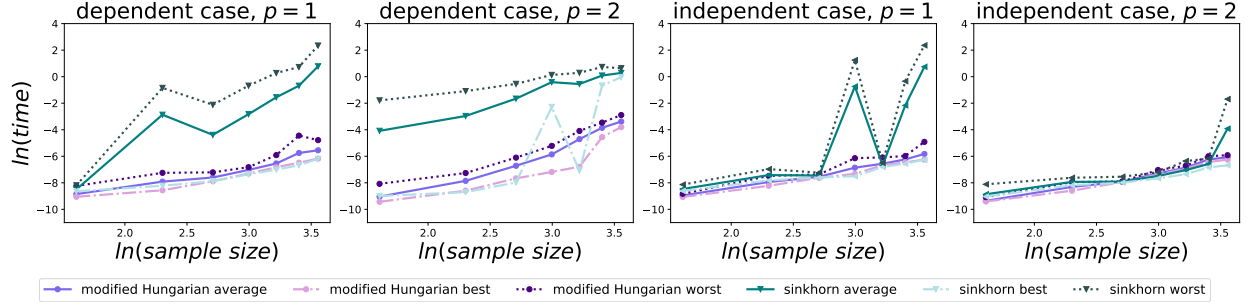


Figure 12: Comparison with the Sinkhorn algorithm on synthetic w.r.t. time

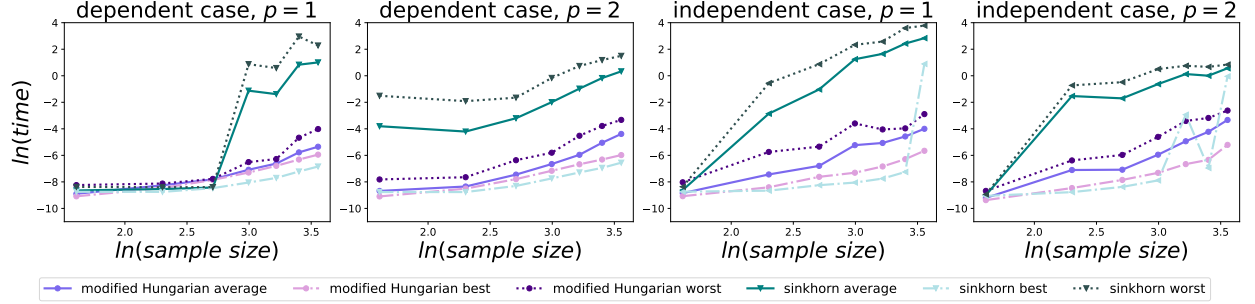


Figure 13: Comparison with the Sinkhorn algorithm on CIFAR10 w.r.t. time

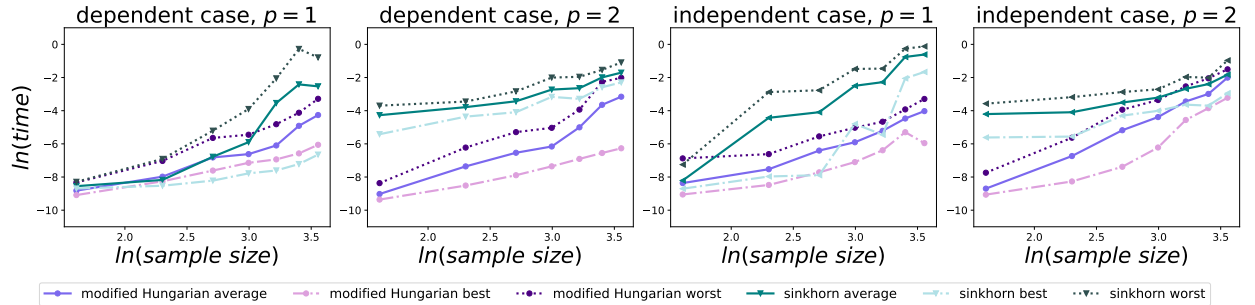


Figure 14: Comparison with the Sinkhorn algorithm on Wisconsin cancer data w.r.t. time