SWIFT: Scalable Wasserstein Factorization for Sparse Nonnegative Tensors

Ardavan Afshar, Kejing Yin, Sherry Yan, Cheng Qian, Joyce Ho, Haesun Park, Jineng Sun

1 Georgia Institute of Technology 2 Hong Kong Baptist University 3 Sutter Health 4 IQVIA
5 Emory University 6 University of Illinois at Urbana-Champaign

aafshar8@gatech.edu, cskjyin@comp.hkbu.edu.hk, YanSX@sutterhealth.org, alextoqc@gmail.com, joyce.c.ho@emory.edu, hpark@cc.gatech.edu, jineng@illinois.edu

Abstract

Existing tensor factorization methods assume that the input tensor follows some specific distribution (i.e., Poisson, Bernoulli, and Gaussian), and solve the factorization by minimizing some empirical loss functions defined based on the corresponding distribution. However, it suffers from several drawbacks: 1) In reality, the underlying distributions are complicated and unknown, making it infeasible to be approximated by a simple distribution. 2) The correlation across dimensions of the input tensor is not well utilized, leading to sub-optimal performance. Although heuristics were proposed to incorporate such correlation as side information under Gaussian distribution, they can not easily be generalized to other distributions. Thus, a more principled way of utilizing the correlation in tensor factorization models is still an open challenge. Without assuming any explicit distribution, we formulate the tensor factorization as an optimal transport problem with Wasserstein distance, which can handle non-negative inputs.

We introduce SWIFT, which minimizes the Wasserstein distance that measures the distance between the input tensor and that of the reconstruction. In particular, we define the N-th order tensor Wasserstein loss for the widely used tensor CP factorization and derive the optimization algorithm that minimizes it. By leveraging sparsity structure and different equivalent formulations for optimizing computational efficiency, SWIFT is as scalable as other well-known CP algorithms. Using the factor matrices as features, SWIFT achieves up to 9.65% and 11.31% relative improvement over baselines for downstream prediction tasks. Under the noisy conditions, SWIFT achieves up to 15% and 17% relative improvements over the best competitors for the prediction tasks.

1 Introduction

Tensor factorization techniques are effective and powerful tools for analyzing multi-modal data and have been shown tremendous success in a wide range of applications including spatio-temporal analysis (Bahadori, Yu, and Liu 2014; Fanacae-T and Gama 2016), graph analysis (Gujral, Parsicha, and Papalexakis 2020), and health informatics (Yin et al. 2019; He, Henderson, and Ho 2019) applications. Many constraints such as non-negativity (Kim, He, and Park 2014), sparsity (Henderson et al. 2017), orthogonality (Wang et al. 2015), and smoothness (Afshar et al. 2018) are imposed on tensor methods in order to improve the performance both quantitatively and qualitatively. Moreover, depending on the nature of input data, tensor methods fit different distributions on data including Gaussian (Bader and Kolda 2007), Poisson (Chi and Kolda 2012), Bernoulli (Hong, Kolda, and Duersch 2020) distributions and minimize various empirical loss functions such as sum square loss, KL-divergence, and log-loss. However, there are several limitations with these techniques. 1) Existing factorization models often assume some specific data distributions, yet in practice, the underlying distributions are complicated and often unknown. 2) The nature of these factorization models neglects correlation relations within each tensor mode (such as external knowledge about similarity among those features). Although there are several extensions to tensor factorization approaches that consider these similarity matrices as side information (Acar, Kolda, and Dunlavy 2011; Kim et al. 2017), they are derived under Gaussian distribution and are not directly generalizable to unknown distributions.

Recent success of Wasserstein distance or loss (a.k.a. earth mover’s distance or optimal transport distance) shows its potential as a better measure of the difference between two distributions (Arjovsky, Chintala, and Bottou 2017; Cuturi 2013; Froger et al. 2015). This distance metric provides a natural measure of the distance between two distributions via a ground metric of choice and can be defined as the cost of the optimal transport plan for moving the mass in the source distribution to match that in the target one. Recently, Wasserstein distance has been applied to matrix factorization and dictionary learning problems with great success (Sandler and Lindenbaum 2009; Rolet, Cuturi, and Peyré 2016; Qian et al. 2016; Schmitz et al. 2018; Varol, Nejabakhsh, and McGrory 2019). To the best of our knowledge, its extension to tensor factorization was never studied and is in fact non-trivial due to the following challenges:

• Wasserstein loss is not well-defined for tensors: The Wasserstein loss is originally defined over vectors, where each entry of the vector represents one physical location and the cost of transporting from one location to another is used. Existing matrix-based Wasserstein loss are defined by the sum of the vector-based Wasserstein loss over the columns of the matrices (Rolet, Cuturi, and Peyré 2016). Unfortunately, this definition is not applicable to tensors of multiple modes.
• Wasserstein loss is difficult to scale: Learning with Wasserstein loss generally requires solving the optimal transport problem in each iteration, which is extremely time-consuming.

• Large and sparse nonnegative input: Existing works on Wasserstein matrix factorization are developed based on dense input with relatively small size. In reality, tensors can often be very large and are often sparse. Efficient learning algorithms are possible only when the sparsity structures of the large input tensor are properly identified and fully utilized.

To overcome these challenges, we propose SWIFT, a tensor factorization method which efficiently minimizes Wasserstein distance for sparse nonnegative tensors. The main contributions of SWIFT include:

• Defining Optimal Transport for Tensors that Handles Nonnegative Input: SWIFT is the first technique that minimizes optimal transport (OT) distance for tensor factorization approaches. The benefit of OT is that it does not assume any specific distribution in the data. SWIFT is able to handle nonnegative inputs such as binary, counts and probability measures, which are common input in real-world tensor data.

• Full Utilization of Data Sparsity and Parallelism: By fully exploring and utilizing the sparsity structure of the input data, SWIFT significantly reduces the number of times required to compute OT and enables parallelism. SWIFT obtains up to 16× faster OT computation than direct implementation of Wasserstein tensor factorization.

• Efficient Computation: SWIFT reduces the amount of computations by smartly rearranging the objective function for solving each factor matrix. Scalability of SWIFT is comparable with well-known CP algorithms. Moreover, SWIFT achieves up to 921× speed up over a direct implementation of Wasserstein tensor factorization without our speedup strategies, as shown in the appendix.

2 Notations and Background

2.1 Basic Notations and Tensor Operations

We denote vectors by bold lowercase letters (e.g. \( \mathbf{u} \)), matrices by bold uppercase letters (e.g. \( \mathbf{A} \)), and tensors by Euler script letters (e.g. \( \hat{X} \)). The entropy \( E \) for a nonnegative matrix \( \mathbf{A} \in \mathbb{R}^{M \times N} \) is defined as \( E(\mathbf{A}) = -\sum_{i,j=1}^{M,N} \mathbf{A}(i,j) \log(\mathbf{A}(i,j)) \).

KL(\( \mathbf{A} \| \mathbf{B} \)) is the general KL-divergence between two matrices \( \mathbf{A}, \mathbf{B} \in \mathbb{R}^{M \times N} \) is defined as \( KL(\mathbf{A} \| \mathbf{B}) = \sum_{i,j=1}^{M,N} \mathbf{A}(i,j) \log(\frac{\mathbf{A}(i,j)}{\mathbf{B}(i,j)}) - \mathbf{A}(i,j) + \mathbf{B}(i,j) \).

Mode-\( n \) Matricization. Matricization (Kolda and Bader 2009) is the process of reordering the entries of a tensor into a matrix. Specifically, the mode-\( n \) matricization is the concatenation of all the mode-\( n \) fibers obtained by fixing the indices for every but the \( n \)-th mode. It transforms the tensor \( \mathbf{X} \in \mathbb{R}^{I_1 \times I_2 \times \ldots \times I_N} \) into matrix \( \mathbb{X}(n) \), and the size of the resulting matrix is \( I_n \) by \( I_1 \ldots I_{n-1} I_{n+1} \ldots I_N \). To ease the notation, we define \( I_{(n)} = I_1 \ldots I_{n-1} I_{n+1} \ldots I_N \).

Khatri-Rao Product. The Khatri-Rao product (Kolda and Bader 2009) of two matrices \( \mathbf{A} \in \mathbb{R}^{I \times R} \) and \( \mathbf{B} \in \mathbb{R}^{J \times R} \) is the column-wise Kronecker product \( \mathbf{C} = \mathbf{A} \circ \mathbf{B} = \{a_1 \odot b_1, a_2 \odot b_2, \ldots, a_R \odot b_R\} \), where \( a_i, b_i \) are the column-\( i \) of matrices \( \mathbf{A} \) and \( \mathbf{B} \), denotes the Khatri-Rao product, and \( \mathbf{C} \in \mathbb{R}^{I \times J \times R} \). We denote the Khatri-Rao product of all factor matrices except the \( n \)-th mode as \( \mathbf{A}^{(-n)} = (\mathbf{A}_1 \odot \ldots \odot \mathbf{A}_{n+1} \odot \mathbf{A}_{n-1} \odot \ldots \odot \mathbf{A}_1) \in \mathbb{R}^{(I_{(n)} \times R)} \), (1) where \( \mathbf{A}_n \in \mathbb{R}^{I_n \times R} \) indicates the \( n \)-th factor matrix.

Canonical/Polyadic (CP) decomposition. The CP factorization (Kolda and Bader 2009) approximates a tensor \( \mathbf{X} \) as the sum of rank-one tensors \( \mathbf{X} = (\mathbf{A}^{(1)}, \mathbf{A}^{(2)}, \ldots, \mathbf{A}^{(N)}) = \sum_{r=1}^{R} a^{(1)} \odot a^{(2)} \odot \ldots \odot a^{(N)} \), where \( \mathbf{A}_n \) is a reconstructed tensor, \( a^{(n)} \) is the \( r \)-th column of factor matrix \( \mathbf{A}^{(n)} \), denotes the outer product of vectors, and \( R \) is the number of the rank-one tensors to approximate the input tensor, i.e., the target rank. The mode-\( n \) matricization of reconstructed tensor \( \hat{X}^{(n)} \) is \( \hat{X}^{(n)} = \mathbf{A}_n(\mathbf{A}_n^{(-n)})^T \in \mathbb{R}^{I_n \times I_{(n)}} \).

2.2 Preliminaries & Related Work

Wasserstein Distance and Optimal Transport. Wasserstein distance (a.k.a. earth mover’s distance or optimal transport distance) computes the distance between two probability vectors\(^1\). Given two vectors \( \mathbf{a}, \mathbf{b} \in \mathbb{R}^n \) and cost matrix \( \mathbf{C} \in \mathbb{R}^{n \times n} \), the Wasserstein distance between \( \mathbf{a}, \mathbf{b} \) is shown by \( W(\mathbf{a}, \mathbf{b}) \) and minimizes \( \langle \mathbf{C}, \mathbf{T} \rangle \) where \( \langle \cdot, \cdot \rangle \) indicates the Frobenius inner product and \( \mathbf{C} \) is a symmetric input cost matrix where \( \mathbf{C}(i,j) \) represents the cost of moving \( a[i] \) to \( b[j] \). \( \mathbf{T} \in U(\mathbf{a}, \mathbf{b}) \) where \( \mathbf{T} \) is an optimal transport solution between probability vectors \( \mathbf{a} \) and \( \mathbf{b} \), and \( U(\mathbf{a}, \mathbf{b}) = \{ \mathbf{T} \in \mathbb{R}^{n \times n} | \mathbf{T} \mathbf{1}_m = \mathbf{a}, \mathbf{T}^T \mathbf{1}_n = \mathbf{b} \} \) is a set of all non-negative \( n \times m \) matrices with row and column sums \( \mathbf{a}, \mathbf{b} \) respectively. \( \mathbf{1}_m \) represents \( m \) dimensional vector of ones. The aforementioned problem has complexity \( O(n^3) \) (assuming \( m = n \)) (Peyré, Cuturi et al. 2019). However, computing this distance metric comes with a heavy computational price (Pele and Werman 2009). In order to reduce the complexity of computation, Cuturi et.al. (Cuturi 2013) propose an entropy regularized optimal transport problem between vectors \( \mathbf{a}, \mathbf{b} \):

\[
W_{\rho}(\mathbf{a}, \mathbf{b}) = \minimize_{\mathbf{T} \in U(\mathbf{a}, \mathbf{b})} \langle \mathbf{C}, \mathbf{T} \rangle - \frac{1}{\rho} E(\mathbf{T}),
\]

where \( E(\mathbf{T}) \) is an entropy function and \( \rho \) is a regularization parameter. When \( \rho \geq 0 \), the solution of (2) is called the Sinkhorn divergence (a.k.a. entropy regularized Wasserstein distance) between probability vectors \( \mathbf{a} \) and \( \mathbf{b} \). Eq. (2) is a strictly convex problem with a unique solution and can be computed with vectors \( \mathbf{u} \in \mathbb{R}^n \), \( \mathbf{v} \in \mathbb{R}^n \) such that \( \text{diag}(\mathbf{u}) \mathbf{K} \text{diag}(\mathbf{v}) = U(\mathbf{a}, \mathbf{b}) \). Here, \( \mathbf{K} = \exp(-\rho \mathbf{C}) \in \mathbb{R}^{n \times n} \). Finding the optimal \( \mathbf{u} \) and \( \mathbf{v} \) can be computed via the Sinkhorn’s algorithm (Sinkhorn and Knopp 1967).

Wasserstein Dictionary Learning. There are several techniques for minimizing Wasserstein loss for dictionary learning problem (Sandler and Lindenbaum 2009; Rolet, Cuturi, and Peyré 2016; Qian et al. 2016; Schmitz et al. 2018;
Varol, Nejabakhsh, and McGrory 2019; Xu 2020). Sandler et al. (Sandler and Lindenbaum 2009) introduces the first non-negative matrix factorization problem that minimizes Wasserstein loss by proposing a linear programming problem, however, their method needs heavy computation. Cuturi (Rolet, Cuturi, and Peyré 2016) proposed a Wasserstein dictionary learning problem based on entropy regularization. (Qian et al. 2016) proposes a similar method by exploiting knowledge in both data manifold and features correlation. Xu (Xu 2020) introduces a nonlinear matrix factorization approach for graphs that considers topological structures. Unfortunately, these approaches cannot be directly generalized to tensor inputs due to the challenges mentioned in Section 1.

### 3 SWIFT Framework

We define and solve optimal transport problem for tensor input by proposing Scalable WassersteinFacTorization (SWIFT) for sparse nonnegative tensors. First we define the input and output for SWIFT. Our proposed method requires the N-th order tensor $X \in R^{I_1 \times ... \times I_N}$ and N cost matrices $C_n \in R^{I_1 \times I_n}$ $(n = 1, ..., N)$ capturing the relations between dimensions along each tensor mode as the input to SWIFT. Here, $C_n$ is a cost matrix for mode n and can be computed directly from the tensor input, or derived from external knowledge. It can also be an identity matrix, meaning that the correlation among features are ignored if the cost matrix is not available. SWIFT is based on CP decomposition and computes N non-negative factor matrices $A_n \in R^{I_1 \times I_n}$ $(n = 1, ..., N)$ as the output. These factor matrices can then be used for downstream tasks, such as clustering and classification.

#### 3.1 Wasserstein Distance for Tensors

**Definition 1 Wasserstein Matrix Distance:** Given a cost matrix $C \in R^{M \times M}$, the Wasserstein distance between two matrices $A = [a_1, ..., a_p] \in R^{M \times P}$ and $B = [b_1, ..., b_p] \in R^{M \times P}$ is denoted by $W_M (A, B)$, and given by:

$$W_M (A, B) = \sum_{p=1}^{P} W_V (a_p, b_p)$$

$$= \min \sum_{p=1}^{P} \{ C, T_p \} - \frac{1}{\rho} E(T_p) \tag{3}$$

$$= \min_{T_p \in U(A, B)} \{ C, T \} - \frac{1}{\rho} E(T) \tag{3}$$

Note that sum of the minimization equals minimization of sums since each $T_p$ is independent of others. Here, $\rho$ is a regularization parameter, $C \equiv [C, ..., C]$ and $T = [T_1, ..., T_p, ..., T_P]$ are concatenations of the cost matrices and the transport matrices for the $P$ optimal transport problems, respectively. Note that $U(A, B)$ is the feasible region of the transport matrix $T$ and is given by:

$$U(A, B) = \{ T \in R^{M \times M} \mid T_p 1_M = a_p, T_p^T 1_M = b_p \ \forall p \}$$

$$= \{ T \in R^{M \times M} \mid \Delta (T) = A, \Psi (T) = B \} \tag{4}$$

where $\Delta (T) = [T_1 1_M, ..., T_p 1_M] = T (1_P \otimes 1_M)$.

**Proposition 1** The Wasserstein distance between tensors $X$ and $Y$ denoted by $W_T (X, Y)$ is a valid distance and satisfies the metric axioms as follows:

1. Positivity: $W_T (X, Y) \geq 0$
2. Symmetry: $W_T (X, Y) = W_T (Y, X)$
3. Triangle Inequality: $\forall X, Y, Z \quad W_T (X, Y) \leq W_T (X, Z) + W_T (Z, Y)$

We provide the proof in the appendix.

#### 3.2 Wasserstein Tensor Factorization

Given an input tensor $X$, SWIFT aims to find the low-rank approximation $\hat{X}$ such that their Wasserstein distance in (5) is minimized. Formally, we solve for $\hat{X}$ by minimizing $W_T (\hat{X}, X)$, where $\hat{X} = \{ A_1, ..., A_N \}$ is the CP factorization of $X$. Together with Definitions 1 and 2, we have the following optimization problem:

$$\min_{\{ A_n \geq 0, T_n \}_{n=1}^{N}} \sum_{n=1}^{N} \left( C_n, T_n \right) - \frac{1}{\rho} E(T_n) \tag{6}$$

subject to $\hat{X} = \{ A_1, ..., A_N \}$

$$\forall \bar{X} \in U(\bar{X}(n), X(n)), n = 1, ..., N$$

where the first constraint enforces a low-rank CP approximation, the second one ensures that the transport matrices are inside the feasible region. We also interested in imposing non-negativity constraint on the CP factor matrices for
both well-definedness of the optimal transport problem and interpretability of the factor matrices. Similar to prior works on vector-based, and matrix-based Wasserstein distance minimization problems (Frogner et al. 2015; Qian et al. 2016), in order to handle non-probability inputs, we convert the second hard constraint in (6) to soft regularizations by Lagrangian method using the generalized KL-divergence. Together with the fact that \( \hat{X}_{(n)} = A_n(A_{\circ}^{(-n)})^T \) for CP factorization, we convert (6) into the following objective function:

\[
\begin{align*}
\text{minimize} \quad & \sum_{n=1}^{N} \left( \frac{\langle C_n, T_n \rangle}{n} - \frac{1}{\rho} E(T_n) + \lambda \left( KL(\Delta(T_n)||A_n(A_{\circ}^{(-n)})^T) + KL(\Psi(T_n)||X_{(n)}) \right) \right) \\
\text{Part } P_1 \\
\text{Part } P_2 \\
\text{Part } P_3
\end{align*}
\]

where \( \Delta(T_n) = [T_{n1}, ..., T_{nj}, ..., T_{nI_{(n)-1}}] \), \( \Psi(T_n) = [T_{n1}^T, ..., T_{nj}^T, ..., T_{nI_{(n)-1}}^T] \), and \( \lambda \) is the weighting parameter for generalized KL-divergence regularization.

We use the alternating minimization to solve (7). SWIFT iteratively updates: 1) **Optimal Transports Problems.** For each mode-\( n \) matricization, SWIFT computes at most a set of \( I_{(-n)} \) optimal transport problems. By exploiting the sparsity structure and avoiding explicitly computing transport matrices we can significantly reduce the computation cost. 2) **Factor Matrices.** The CP factor matrices are involved inside the Khatri-Rao product and needs excessive amount of computation. However, by rearranging the terms involved in (7), we can efficiently update each factor matrix. Next, we provide efficient ways to update optimal transport and factor matrices in more details.

3.3 Solution for Optimal Transport Problems

For mode-\( n \), \( \bar{T}_n = [T_{n1}, ..., T_{nj}, ..., T_{nI_{(n)-1}}] \in \mathbb{R}^{I_{n} \times I_{(n)-1}} \) includes a set of \( I_{(-n)} \) different optimal transport problems. The optimal solution of \( j \)-th optimal transport problem for mode \( n \) is \( T_{n} = \text{diag}(u_j)K_{n}\text{diag}(v_j) \), where \( K_n = e^{-\gamma C_n} \in \mathbb{R}^{I_{n} \times I_{n}} \) and \( u_j, v_j \in \mathbb{R}^{I_{(n)-1}} \). This requires computing \( I_{(n)-1} \) transport matrices with size \( I_{n} \times I_{n} \) in (7), which is extremely time-consuming. To reduce the amount of computation, SWIFT proposes the following three strategies:

1) Never explicitly computing transport matrices (\( T_{n} \)): Although we are minimizing (7) with respect to \( T_n \), we never explicitly compute \( T_n \). In stead of directly computing the transport matrices \( T_{n} \), we make use of the constraint \( T_{n}^\odot 1 = \text{diag}(u_j)K_{n}\text{diag}(v_j) \) where \( \odot \) denotes element-wise product. As a result, the following proposition effectively updates objective function 7.

\[
\text{Proposition 2} \quad \Delta(T_n) = [T_{n1}, ..., T_{nj}^\odot 1, ..., T_{nI_{(n)-1}}^\odot 1] = U_n \ast (K_nV_n) \quad \text{minimizes} \quad (7) \text{ where } U_n = (X_n)^\Phi \odot (K_n(X_n^\mathcal{H}) \odot (K_n^\mathcal{H}U_n))^\Phi, \quad V_n = (X_n \odot (K_n^\mathcal{H}U_n))^\Phi, \quad \Phi = \frac{\lambda_p}{\lambda_{p+1}}, \text{ and } \odot \text{ indicates element-wise division. See Section appendix for proof.}
\]

2) Exploiting Sparsity Structure in \( X_{(n)} \in \mathbb{R}^{I_{n} \times I_{(-n)}} \):

We observe that there are many columns with all zero elements in \( X_{(n)} \) due to the sparsity structure in the input data. There is no need to compute transport matrix for those zero columns, therefore, we can easily drop zero value columns in \( X_{(n)} \) and its corresponding columns in \( U_n, V_n, \) and \( \hat{X}_{(n)} \) from our computations. We use \( NNZ_n \) to denote the number of non-zero columns in \( X_{(n)} \). By utilizing this observation, we reduce the number of times to solve the optimal transport problems from \( I_{(n)-1} \) to \( NNZ_n \), where we usually have \( NNZ_n \ll I_{(n)-1} \) for sparse input.

3) Parallelization of the optimal transport computation: The \( NNZ_n \) optimal transport problems for each factor matrix \( X_{(n)} \) can be solved independently. Therefore, parallelization on multiple processes is straightforward for SWIFT.

3.4 Solution for Factor Matrices

All the factor matrices are involved in Part \( P_2 \) of Objective (7) and present in \( N \) different KL-divergence terms. The objective function with respect to factor matrix \( A_n \) for mode \( n \) is:

\[
\begin{align*}
\text{minimize} \quad & \sum_{i=1}^{N} KL \left( \Delta(T_i) || A_i(A_{\circ}^{(-i)})^T \right) \\
\text{Part } P_1 \\
\text{Part } P_2 \\
\text{Part } P_3
\end{align*}
\]

where \( \Delta(T_i) = [T_{i1}, ..., T_{ij}, ..., T_{iI_{(i)-1}}] \), \( A_i \in \mathbb{R}^{I_{(i)} \times R} \), \( A_{\circ}^{(-i)} \in \mathbb{R}^{I_{(-i)} \times R} \). Updating factor matrix \( A_n \) in (8) is expensive due to varying positions of \( A_n \) in the \( N \) KL-divergence terms. Specifically, \( A_n \) is involved in the Khatri-Rao product \( A_{\circ}^{(-i)} \), as defined in (1), for every \( i \neq n \). On the other hand, if \( i = n \), \( A_n \) is not involved in \( A_{\circ}^{(-i)} \).

Efficient rearranging operations. In order to solve (8) efficiently, we introduce operator \( \Pi \), which performs a sequence of reshape, permute and another reshape operations, such that, when applied to the right-hand side of (8), \( A_n \) is no longer involved inside the Khatri-Rao product for all \( i \neq n \). Formally,

\[
\Pi(A_{\circ}^{(-i)})^T, n) = A_n(A_{\circ}^{(-n)})^T \in \mathbb{R}^{I_{n} \times I_{(-n)}} \forall i \neq n.
\]

To maintain equivalence to (8), we apply the same operation to the left-hand side of (8), which leads us to the following formulation:

\[
\begin{align*}
\text{minimize} \quad & KL \left( \left\{ \Pi(\Delta(T_i), n) \right\} || A_i(A_{\circ}^{(-i)})^T \right) \\
\text{Part } P_1 \\
\text{Part } P_2 \\
\text{Part } P_3
\end{align*}
\]

Where \( \Pi(\Delta(T_i), n) \in \mathbb{R}^{I_{n} \times I_{(-n)}} \) for all \( i \). Due to the fact that KL-divergence is computed point-wisely, the above formula is equivalent to (8), with the major difference that \( A_n \) is at the same position in every KL-divergence term in (10), and is no longer involved inside the Khatri-Rao product terms; therefore, it can be much more efficiently updated via multiplicative update rules (Lee and Seung 2001). More details
regarding operator II, and multiplicative update rules for $A_n$ are provided in the appendix.

In every iteration of SWIFT, we first update $N$ different optimal transport problems and then update $N$ factor matrices. Algorithm 1 summarizes the optimization procedure in SWIFT.

**Proposition 3** SWIFT is based on Block Coordinate Descent (BCD) algorithm and guarantees convergence to a stationary point. See detailed proofs in the appendix.

Details regarding complexity analysis are provided in appendix.

**Algorithm 1: SWIFT**

<table>
<thead>
<tr>
<th>Input: $X \in \mathbb{R}_+^{I_1 \times I_2 \times \cdots \times I_N}$, $C_n$, $NNZ_n$, $n = 1, ..., N$, target rank $R$, $\lambda$, and $\rho$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Output: $A_n \in \mathbb{R}_+^{I_n \times R}$, $n = 1, ..., N$</td>
</tr>
</tbody>
</table>

1. Initialize $A_n$, $n = 1, ..., N$ randomly;
2. while stopping criterion is not met do
   1. for $n = 1, ..., N$ do
      1. // Optimal Transport Update (Section 3.3);
      2. $\Phi = \frac{\lambda_F}{\sqrt{I_n}}$;
      3. $U_n = \text{ones}(I_n, NNZ_n) \odot I_n$;
      4. for $s = 1, ..., \text{Sinkhorn Iteration}$ do
         1. $U_n = (\bar{X}_{(n)})^\# \odot (K_n(X_{(n)} \odot (K_n^T U_n))^\#)^*$;
      end
      6. $V_n = (X_{(n)} \odot (K_n^T U_n))^*$;
      7. $\Delta(T_n) = U_n * (K_n V_n)$;
   end
3. for $n = 1, ..., N$ do
   1. // Factor Matrix Update (Section 3.4);
   2. Update $A_n$ based on (10);
4. end

4 Experimental Results

4.1 Experimental Setup

Datasets and Evaluation Metrics

1) BBC News (Greene and Cunningham 2006) is a publicly available dataset from the BBC News Agency for text classification task. A third-order count tensor is constructed with the size of 400 articles by 100 words by 100 words. $\mathcal{X}(i, j, k)$ is the number of co-occurrences of the $j$-th and the $k$-th words in every sentence of the $i$-th article. We use the pair-wise cosine distance as the word-by-word and article-by-article cost matrices with details provided in the appendix. The downstream task is to predict the category (from business, entertainment, politics, sport or tech) of each article and we use accuracy as the evaluation metric.

2) Sutter is a dataset collected from a large real-world health provider network containing the electronic health records (EHRs) of patients. A third-order binary tensor is constructed with the size of 1000 patients by 100 diagnoses by 100 medications. The downstream task is to predict the onset of heart failure (HF) for the patients (200 out of the 1000 patients are diagnosed with HF) and use PR-AUC (Area Under the Precision-Recall Curve) to evaluate the HF prediction task.

We chose these two datasets because of different data types (count in BBC, binary in Sutter). Note that the cost matrices are derived from the original input by cosine similarity without any additional external knowledge. Hence the comparisons are fair since the input are the same.

**Baselines** We compare the performance of SWIFT with different tensor factorization methods with different loss functions and their variants:

- The first loss function minimizes sum square loss and has 4 variants: 1) CP-ALS (Bader and Kolda 2007); 2) CP-NMU (Bader and Kolda 2007); 3) Supervised CP (Kim et al. 2017); and 4) Similarity based CP (Kim et al. 2017). The first one is unconstrained, the second one incorporates non-negativity constraint, the third one utilizes label information, and the last one uses similarity information among features (similar to SWIFT).
- The second loss function is Gamma loss (CP-Continuous (Hong, Kolda, and Duersch 2020)) which is the start of the art method (SOTA) for non-negative continuous tensor.
- The third loss function is Log-loss (CP-Binary (Hong, Kolda, and Duersch 2020)) which is SOTA binary tensor factorization by fitting Bernoulli distribution.
- The fourth loss function is Kullback-Leibler Loss (CP-APR (Chi and Kolda 2012)) which fits Poisson distribution on the input data and is suitable for count data.

4.2 Classification Performance of SWIFT

To evaluate low-rank factor matrices, we utilize the downstream prediction tasks as a proxy to assess the performance of SWIFT and the baselines, similar to the existing works (Ho, Ghosh, and Sun 2014; Yin et al. 2019; Afshar et al. 2020; Yin et al. 2020). We performed 5-fold cross validation and split the data into training, validation, and test sets by a ratio of 3:1:1.

**Outperforming various tensor factorizations:** Table 1 summarizes the classification performance using the factor matrices obtained by SWIFT and the baselines with varying target rank ($R \in \{5, 10, 20, 30, 40\}$). We report the mean and standard deviation of accuracy for BBC News, and that of PR-AUC Score for Sutter dataset over the test set. For BBC News, SWIFT outperforms all baselines for different target ranks with relative improvement ranging from 1.69% to 9.65%. For Sutter, SWIFT significantly outperforms all baselines for all values of $R$ with relative improvement ranging from 5.10% to 11.31%.

**Outperforming various classifiers:** We further compare the performance of SWIFT against widely-adopted classifiers, including Lasso Logistic Regression, Random Forest, Multi-Layer Perceptron, and K-Nearest Neighbor, using raw data. The input for BBC and Sutter to these classifiers are obtained by matricizing the input tensors along the article.
we add the noise by flipping the selected zero value to a count
we inject noise to the raw input tensor to construct the noisy
to measure the performance of SWIFT
SWIFT
the first part reports the average and standard deviation of accuracy on BBC NEWS data.
The second part depicts the average and standard deviation of PR-AUC score on test data for Sutter dataset. Both experiments are based on five-fold cross validation. We used Lasso Logistic Regression as a classifier.

Table 1: The first part reports the average and standard deviation of accuracy on BBC NEWS dataset.
The second part depicts the average and standard deviation of running time of classification.

### Table 1: Average and standard deviation of accuracy on BBC News and Sutter datasets.

<table>
<thead>
<tr>
<th>Dataset</th>
<th>R=5</th>
<th>R=10</th>
<th>R=20</th>
<th>R=30</th>
<th>R=40</th>
</tr>
</thead>
<tbody>
<tr>
<td>CP-ALS</td>
<td>.521 ± .033</td>
<td>.571 ± .072</td>
<td>.675 ± .063</td>
<td>.671 ± .028</td>
<td>.671 ± .040</td>
</tr>
<tr>
<td>CP-NMU</td>
<td>.484 ± .039</td>
<td>.493 ± .048</td>
<td>.581 ± .064</td>
<td>.600 ± .050</td>
<td>.650 ± .031</td>
</tr>
<tr>
<td>Supervised CP</td>
<td>.506 ± .051</td>
<td>.625 ± .073</td>
<td>.631 ± .050</td>
<td>.665 ± .024</td>
<td>.662 ± .012</td>
</tr>
<tr>
<td>Similarity Based CP</td>
<td>.518 ± .032</td>
<td>.648 ± .043</td>
<td>.638 ± .021</td>
<td>.662 ± .034</td>
<td>.673 ± .043</td>
</tr>
<tr>
<td>CP-Continuous</td>
<td>.403 ± .051</td>
<td>.481 ± .056</td>
<td>.528 ± .022</td>
<td>.559 ± .024</td>
<td>.543 ± .043</td>
</tr>
<tr>
<td>CP-Binary</td>
<td>.746 ± .058</td>
<td>.743 ± .027</td>
<td>.737 ± .008</td>
<td>.756 ± .062</td>
<td>.743 ± .044</td>
</tr>
<tr>
<td>CP-APR</td>
<td>.675 ± .059</td>
<td>.768 ± .033</td>
<td>.753 ± .035</td>
<td>.743 ± .033</td>
<td>.746 ± .043</td>
</tr>
<tr>
<td>SWIFT</td>
<td>.759 ± .013</td>
<td>.781 ± .013</td>
<td>.803 ± .010</td>
<td>.815 ± .005</td>
<td>.818 ± .022</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Dataset</th>
<th>R=5</th>
<th>R=10</th>
<th>R=20</th>
<th>R=30</th>
<th>R=40</th>
</tr>
</thead>
<tbody>
<tr>
<td>CP-ALS</td>
<td>.327 ± .072</td>
<td>.333 ± .064</td>
<td>.311 ± .068</td>
<td>.306 ± .065</td>
<td>.332 ± .098</td>
</tr>
<tr>
<td>CP-NMU</td>
<td>.300 ± .054</td>
<td>.294 ± .064</td>
<td>.325 ± .085</td>
<td>.344 ± .068</td>
<td>.302 ± .071</td>
</tr>
<tr>
<td>Supervised CP</td>
<td>.301 ± .044</td>
<td>.305 ± .036</td>
<td>.309 ± .054</td>
<td>.291 ± .037</td>
<td>.293 ± .051</td>
</tr>
<tr>
<td>Similarity Based CP</td>
<td>.304 ± .042</td>
<td>.315 ± .041</td>
<td>.319 ± .063</td>
<td>.296 ± .041</td>
<td>.303 ± .032</td>
</tr>
<tr>
<td>CP-Continuous</td>
<td>.252 ± .059</td>
<td>.237 ± .043</td>
<td>.263 ± .065</td>
<td>.244 ± .053</td>
<td>.256 ± .077</td>
</tr>
<tr>
<td>CP-Binary</td>
<td>.301 ± .061</td>
<td>.325 ± .079</td>
<td>.328 ± .080</td>
<td>.267 ± .074</td>
<td>.296 ± .063</td>
</tr>
<tr>
<td>CP-APR</td>
<td>.305 ± .075</td>
<td>.301 ± .068</td>
<td>.290 ± .052</td>
<td>.313 ± .082</td>
<td>.304 ± .086</td>
</tr>
<tr>
<td>SWIFT</td>
<td>.364 ± .063</td>
<td>.350 ± .031</td>
<td>.350 ± .040</td>
<td>.369 ± .066</td>
<td>.374 ± .044</td>
</tr>
</tbody>
</table>

Table 2: Average and standard deviation of accuracy on BBC NEWS and PR-AUC score on Sutter data sets by performing Lasso LR, RF, MLP, and KNN on raw data sets.

<table>
<thead>
<tr>
<th>Dataset</th>
<th>Accuracy on BBC</th>
<th>PR-AUC on Sutter</th>
</tr>
</thead>
<tbody>
<tr>
<td>Lasso LR</td>
<td>.728 ± .013</td>
<td>.308 ± .033</td>
</tr>
<tr>
<td>RF</td>
<td>.6281 ± .049</td>
<td>.318 ± .083</td>
</tr>
<tr>
<td>MLP</td>
<td>.690 ± .052</td>
<td>.305 ± .054</td>
</tr>
<tr>
<td>KNN</td>
<td>.5956 ± .067</td>
<td>.259 ± .067</td>
</tr>
<tr>
<td>SWIFT (R=5)</td>
<td>.759 ± .013</td>
<td>.364 ± .063</td>
</tr>
<tr>
<td>SWIFT (R=40)</td>
<td>.818 ± .020</td>
<td>.374 ± .044</td>
</tr>
</tbody>
</table>

4.3 Classification Performance on Noisy Data

To measure the performance of SWIFT against noisy input, we inject noise to the raw input tensor to construct the noisy input tensor. For the binary tensor input, we add Bernoulli noise. Given the noise level (p), we randomly choose zero elements, such that the total number of selected zero elements equals to the number of non-zero elements. Then, we flip the selected zero elements to one with probability p. We follow the same procedure for the count tensor input, except that we add the noise by flipping the selected zero value to a count value with probability p, and the added noise value is selected uniformly at random between 1 and maximum value in the tensor input.

Performance on BBC data with Noise: Figure 1 presents the average and standard deviation of categorizing the arti-

<table>
<thead>
<tr>
<th>Dataset</th>
<th>Accuracy on BBC</th>
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</tr>
</tbody>
</table>

Figure 1: The average and standard deviation of accuracy of different baselines as a function of the noise level on BBC NEWS. SWIFT outperforms other baselines by improving accuracy up to 15%.

4.4 Scalability of SWIFT

In this section, we assess the scalability of SWIFT in comparison to the other 7 CP algorithms introduced earlier. Figure 2 depicts the average and standard deviation of running time of one iteration in seconds. In this experiment, we set R = 40. In order to have fair comparisons, we switched off the par-

Noise Level | Performance on BBC data with Noise: Figure 1 presents the average and standard deviation of categorizing the arti-

mode and the patient mode, respectively. Table 2 summarizes the results, and it clearly shows that SWIFT using Lasso LR classifier even with R=5 outperforms all the other classifiers compared.

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Performance on BBC data with Noise: Figure 1 presents the average and standard deviation of categorizing the arti-

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Table 3: Three phenotypes examples learned on Sutter dataset with SWIFT. The weight value for each phenotype represents the lasso logistic regression coefficient for the heart failure prediction task. “Dx” represents for diagnoses and “Rx” indicates for medications. All phenotypes are considered clinically meaningful by a clinical expert.

<table>
<thead>
<tr>
<th>Phenotype Description</th>
<th>Weight</th>
</tr>
</thead>
<tbody>
<tr>
<td>Atrial Fibrillation</td>
<td>21.93</td>
</tr>
<tr>
<td>Dx-Essential hypertension</td>
<td>98.</td>
</tr>
<tr>
<td>Dx-Cardiac dysrhythmias</td>
<td>106.</td>
</tr>
<tr>
<td>Rx-Calcium Channel Blockers</td>
<td></td>
</tr>
<tr>
<td>Rx-Alpha-Beta Blockers</td>
<td></td>
</tr>
<tr>
<td>Rx-Angiotensin II Receptor Antagonists</td>
<td></td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Phenotype Description</th>
<th>Weight</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cardiometabolic Disease</td>
<td>19.58</td>
</tr>
<tr>
<td>Dx-Diabetes mellitus without complication</td>
<td>49.</td>
</tr>
<tr>
<td>Dx-Essential hypertension</td>
<td>98.</td>
</tr>
<tr>
<td>Dx-Disorders of lipid metabolism</td>
<td>53.</td>
</tr>
<tr>
<td>Rx-Diagnostic Tests</td>
<td></td>
</tr>
<tr>
<td>Rx-Biguanides</td>
<td></td>
</tr>
<tr>
<td>Rx-Diabetic Supplies</td>
<td></td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Phenotype Description</th>
<th>Weight</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mental Disorder</td>
<td>-16.22</td>
</tr>
<tr>
<td>Dx-Anxiety disorders</td>
<td>651.</td>
</tr>
<tr>
<td>Dx-Menopausal disorders</td>
<td>173.</td>
</tr>
<tr>
<td>Dx-Depressive disorders</td>
<td>6572</td>
</tr>
<tr>
<td>Rx-Benzodiazepines</td>
<td></td>
</tr>
<tr>
<td>Rx-Selective Serotonin Reuptake Inhibitors (SSRIs)</td>
<td></td>
</tr>
<tr>
<td>Rx-Serotonin Modulators</td>
<td></td>
</tr>
</tbody>
</table>

Figure 2: The running time of one iteration in seconds on BBC News and Sutter with $R = 40$.

Figure 2: The running time of one iteration in seconds on BBC News and Sutter with $R = 40$.

4.5 Interpretability of SWIFT

To demonstrate the interpretability of results produced by SWIFT we perform computational phenotyping using Sutter dataset.

Computational phenotyping is a fundamental task in healthcare. It refers to extracting meaningful and interpretable medical concepts (patient clusters) from noisy electronic health records (EHRs) (Richesson et al. 2016). Tensor factorization techniques are powerful tools for extracting meaningful phenotyping (Ho, Ghosh, and Sun 2014; Yin et al. 2018, 2020b; Zhao et al. 2019; Afshar et al. 2020; Yin et al. 2020a).

Here we extract heart failure (HF) phenotypes using Sutter dataset. We use the same tensor as we described in the previous section and run SWIFT by selecting $R = 40$ since it achieves the highest PR-AUC in comparison to other values of $R$. $A_2(:, r), A_3(:, r)$ represent the membership value of diagnosis and medication features in $r$-th phenotype.

Results: Table 3 lists three examples of the discovered phenotypes. The weight next to phenotype index indicates the lasso logistic regression coefficient for heart failure prediction (i.e., 21.93, 19.58 and -16.22 in Table 3). “Dx” indicates diagnoses and “Rx” represents medications. All phenotypes are clinically meaningful, endorsed and annotated by a medical expert.

The first phenotype is about Atrial Fibrillation which captures patients with hypertension and cardiac dysrhythmias that are associated with high blood pressure medications. Cardiometabolic disease is another phenotype that captures diabetes patients with hypertension. These two phenotypes have high positive weights (21.93 and 19.58 respectively) for predicting HF diagnosis. The third phenotype is depressive and menopausal disorders, Serotonin and Benzodiazepines appeared in this phenotype are medications that are commonly prescribed to patients with these conditions in clinical practice. This phenotype has a negative association with HF (weight = -16.22). The remaining phenotypes positively associated with HF are listed in the appendix.

5 Conclusion

In this paper, we define the Wasserstein distance between two tensors and propose SWIFT, an effective tensor factorization model based on the defined tensor Wasserstein distance. To efficiently learn the Wasserstein tensor factorization, we introduce introduce several techniques, including exploitation of the sparsity structure of the input tensor, efficient rearrangement by utilizing tensor properties, and parallelization of the optimal transport computation. Experimental results depict that SWIFT consistently outperforms baselines in downstream classification tasks for both binary and count tensor inputs. In the presence of noise, SWIFT also outperforms baselines by a large margin.
Acknowledgments
This work is in part supported by National Science Foundation award SCH-2014438, IIS-1418511, CCF-1533768, IIS-1838042, the National Institute of Health award NIH R01 1R01NS107291-01 and R56HL138415.

References


A Symbols and Notations

Table 4 presents the symbols and notations used in this paper.

<table>
<thead>
<tr>
<th>Symbol</th>
<th>Definition</th>
</tr>
</thead>
<tbody>
<tr>
<td>*</td>
<td>Element-wise multiplication</td>
</tr>
<tr>
<td>⊗</td>
<td>Kronecker product</td>
</tr>
<tr>
<td>⊙</td>
<td>Khatri–Rao product</td>
</tr>
<tr>
<td>⊘</td>
<td>Element-wise division</td>
</tr>
<tr>
<td>X, X, x</td>
<td>Tensor, Matrix, vector, scalar</td>
</tr>
<tr>
<td>X_{(n)}</td>
<td>Mode-n Matricization</td>
</tr>
<tr>
<td>A(i,:)</td>
<td>the i-th row of A</td>
</tr>
<tr>
<td>A(:,r) or a_r</td>
<td>the r-th column of A</td>
</tr>
<tr>
<td>A_n</td>
<td>n-th factor matrix</td>
</tr>
<tr>
<td>A_{(-n)}</td>
<td>Khatri-Rao product of all factor matrices except the n-th one</td>
</tr>
<tr>
<td>&lt; x, y &gt; = x^T y</td>
<td>Inner product between x, y</td>
</tr>
<tr>
<td>vec(A)</td>
<td>converting matrix A into a vector</td>
</tr>
</tbody>
</table>

B Proofs

B.1 Proof of Proposition 1

For simplicity, we ignore the entropy regularization term ($\frac{1}{\rho} E(T)$) similar to (Cuturi 2013). In order to prove that Wasserstein distance for tensors is a valid metric, we assume that the cost matrix of mode-$n$ ($C_{(n)} \in \mathbb{R}_{+}^{I_n \times I_n}$):

- Is symmetric ($C_{(n)}(i,j) = C_{(n)}(j,i); \forall i, j$).
- All of its off diagonal elements are positive and non-zero. ($C_{(n)}(i, j) = 0, \forall i = j$)
• $C_{(n)}(i, j) \leq C_{(n)}(i, k) + C_{(n)}(k, j) \quad \forall i, j, k$

The positivity and symmetry of a distance follow then from

$$W_T(\mathcal{X}, \mathcal{Y}) = \sum_{n=1}^{N} W_M\left(\mathbf{x}_{(n)}, \mathbf{y}_{(n)}\right)$$  \hspace{1cm} (11a)

$$= \sum_{n=1}^{N} \sum_{i_n=1}^{I_{(-n)}} W_V(\mathbf{x}_{n_{i_n}}, \mathbf{y}_{n_{i_n}})$$  \hspace{1cm} (11b)

$$= \sum_{n=1}^{N} I_{(-n)} (C_{(n)} \cdot \mathbf{T}_{n})$$  \hspace{1cm} (11c)

$$= \sum_{n=1}^{N} \sum_{i_n=1}^{I_{(-n)}} W_V(\mathbf{y}_{n_{i_n}}, \mathbf{x}_{n_{i_n}})$$  \hspace{1cm} (11d)

$$= \sum_{n=1}^{N} W_M\left(\mathbf{y}_{(n)}, \mathbf{x}_{(n)}\right)$$  \hspace{1cm} (11e)

$$= W_T(\mathcal{Y}, \mathcal{X})$$  \hspace{1cm} (11f)

Here, $\mathbf{x}_{n_{i_n}} \in \mathbb{R}^{I_{n}}$, $\mathbf{y}_{n_{i_n}} \in \mathbb{R}^{I_{n}}$ are probability vectors and $\|\mathbf{x}_{n_{i_n}}\|_1 = \|\mathbf{y}_{n_{i_n}}\|_1 = 1$. (11a) and (11b) are expanded based on Definitions 2, 1. (11c) is written based on the definition of Wasserstein distance for two vectors. (11d) is written based on the fact that $C_{(n)}$ is a symmetric matrix $\forall n$. Similarly, (11e) and (11f) are written based on Definitions 1, 2. These equations show that Wasserstein distance for two tensors is symmetric. The proof of positivity is straightforward, since all the off diagonal elements of $C_{(n)}$ are positive, therefore $(C_{(n)} \cdot \mathbf{T}_{n}) > 0 \quad \forall i_{n}, n$ which suggests $W_T(\mathcal{X}, \mathcal{Y}) > 0$.

To prove the triangle inequality of Wasserstein distances for tensors we know that

$$W_T(\mathcal{X}, \mathcal{Y}) = \sum_{n=1}^{N} \sum_{i_n=1}^{I_{(-n)}} W_V(\mathbf{x}_{n_{i_n}}, \mathbf{y}_{n_{i_n}})$$  \hspace{1cm} (12)

$$W_T(\mathcal{X}, \mathcal{Z}) = \sum_{n=1}^{N} \sum_{i_n=1}^{I_{(-n)}} W_V(\mathbf{x}_{n_{i_n}}, \mathbf{z}_{n_{i_n}})$$  \hspace{1cm} (13)

$$W_T(\mathcal{Z}, \mathcal{Y}) = \sum_{n=1}^{N} \sum_{i_n=1}^{I_{(-n)}} W_V(\mathbf{z}_{n_{i_n}}, \mathbf{y}_{n_{i_n}})$$  \hspace{1cm} (14)

We already know that triangle inequality of Wasserstein distances for every three probability vectors holds (Cuturi 2013; Peyré, Cuturi et al. 2019) which means:

$$\forall \mathbf{x}_{n_{i_n}}, \mathbf{y}_{n_{i_n}}, \mathbf{z}_{n_{i_n}} \quad W_V(\mathbf{x}_{n_{i_n}}, \mathbf{y}_{n_{i_n}}) \leq W_V(\mathbf{x}_{n_{i_n}}, \mathbf{z}_{n_{i_n}}) + W_V(\mathbf{z}_{n_{i_n}}, \mathbf{y}_{n_{i_n}})$$  \hspace{1cm} (15)

By knowing this fact we can extend it to every triple of tensors $(\mathcal{X}, \mathcal{Y}, \mathcal{Z})$ where their unfolding includes probability vectors:

$$W_T(\mathcal{X}, \mathcal{Y}) \leq W_T(\mathcal{X}, \mathcal{Z}) + W_T(\mathcal{Z}, \mathcal{Y})$$  \hspace{1cm} (16)

B.2 Proof of Proposition 2

Here, we follow the same strategy as (Frogner et al. 2015). $\mathbf{T}_{n_{i_k}}$ is the optimal transport matrix between $\tilde{\mathbf{x}}_{(n)}(:, j), \mathbf{x}_{(n)}(:, j)$ in mode-$n$. In order to compute the $\mathbf{T}_{n_{i_k}}$, we need to take the derivative with respect to $T_{n_{i_k}}$.

$$\frac{\partial W_T(\mathcal{X}, \tilde{\mathcal{X}})}{\partial T_{n_{i_k}}(i, k)} = \frac{\partial}{\partial T_{n_{i_k}}(i, k)} \left( \frac{\partial (\mathbf{C}_{n}, \mathbf{T}_{n})}{\partial T_{n_{i_k}}(i, k)} - \rho E(\mathbf{T}_{n}) + \lambda \left( KL(\Delta(\mathbf{T}_{n})) || \tilde{\mathbf{x}}_{(n)} ) + KL(\Psi(\mathbf{T}_{n}) || \mathbf{x}_{(n)} ) \right) \right) = 0$$
\[ \Rightarrow C_n(i, k) + \frac{1}{\rho} \log(T_{nj}(i, k)) + 1 + \lambda \log \left( T_{nj} \circ X_{nj}(i, j) \right) + \lambda \log \left( T_{nj}^T \circ X_{nj}(i, j) \right) = 0 \]

\[ \Rightarrow \log(T_{nj}(i, k)) + \lambda \rho \left( \log(T_{nj} \circ X_{nj}(i, j)) + \log \left( T_{nj}^T \circ X_{nj}(i, j) \right) \right) = -\rho C_n(i, k) - 1 \]

\[ \Rightarrow T_{nj}(i, k) \ast \left( \frac{T_{nj} \circ X_{nj}(i, j)}{\lambda \rho} \right)^{\lambda \rho} \left( T_{nj}^T \circ X_{nj}(i, j) \right)^{\lambda \rho} \exp(-\rho C_n(i, k) - 1) \]

\[ \Rightarrow T_{nj}(i, k) = \left( X_{nj}(i, j) \circ T_{nj} \right)^{\lambda \rho} \left( T_{nj}^T \circ X_{nj}(i, j) \right)^{\lambda \rho} \exp(-\rho C_n(i, k) - 1) \]

Therefore, \( T_{nj} = \text{diag}(u_j)K_n\text{diag}(v_j) \) where \( u_j = (X_{nj}(i, j) \circ T_{nj} \lambda \rho) \in R^I \) and \( v_j = (X_{nj}(i, j) \circ T_{nj}^T \lambda \rho) \in R^I \) and \( K_n = e^{(-\rho C_n - 1)} \) where \( \circ \) represents element-wise division.

\[ T_{nj}1 = \text{diag}(u_j)K_nv_j = u_j \ast (K_nv_j) = \left( X_{nj}(i, j) \circ T_{nj} \right)^{\lambda \rho} \ast (K_nv_j) \]

\[ \Rightarrow (T_{nj}1)^{(\lambda \rho+1)} = \left( X_{nj}(i, j) \right)^{\lambda \rho} \ast (K_nv_j) \]

\[ \Rightarrow \text{diag}(u_j)(K_nv_j)^{(\lambda \rho+1)} = \text{diag}(X_{nj}(i, j))^{\lambda \rho}(K_nv_j) \]

\[ u_j^{(\lambda \rho+1)} = \left( X_{nj}(i, j) \right)^{\lambda \rho} \ast (K_nv_j) \]

\[ u_j = \left( X_{nj}(i, j) \right)^{\lambda \rho} \ast (K_nv_j) \]

\[ \Rightarrow U_n = [u_1, u_2, ..., u_{I(-n)}] = \left( X_{nj}(i, j) \right)^{\lambda \rho} \ast (K_nv) \ast (K_nV) \ast (K_nU_n)^{\lambda \rho} \]

By applying the similar procedure \( V_n = (X_{nj}(i, j) \circ \left( K_n^T U_n \right) \circ \left( K_n^T U_n \right)^{\lambda \rho}) \). Therefore, \( U_n = \left( X_{nj}(i, j) \right)^{\lambda \rho} \ast (K_nv) \ast (K_nV) \ast (K_nU_n)^{\lambda \rho} \).

By knowing \( T_{nj}1 = u_j \ast (K_nv_j) \) we can extend \( \Delta_n = [T_{nj}1, ..., T_{nj}I(-n)1] = U_n \ast (K_nv_n) \).

### B.3 Proof of Proposition 3

Consider SWIFT’s objective function:

\[ \min_{\{A_n \geq 0, T_n\}_{n=1}^N} \sum_{n=1}^N \left( \frac{1}{\rho} E(T_n) + \lambda KL(\Delta(T_n)||A_n(A_{\cap}^{-n})^T) + KL(\Psi(T_n)||X_{nj}(i, j)) \right) \]

The variables in (17) separated into \( N \) different optimal transport problems (\( T_n \)) and \( N \) different factor matrices (\( A_n \)). We use Block Coordinate Descent (BCD) framework. BCD iteratively solves one variable at a time while fixing others. Therefore, in every step the objective function is minimized with respect to each of the variables (Bertsekas 1997 Section 3.7). Each sub-problem in (17) is continuous, differentiable, and strictly convex since minimizing the Frobenius inner product, negative entropy, and KL-Divergence are strictly convex problems (Boyd, Boyd, and Vandenberghe 2004). Therefore, based on (Bertsekas 1997 Section 3.7), (17) converges to a stationary point.

### C Other Technical Details

#### C.1 Details on Updating Factor Matrix \( A_n \)

Algorithm 2 presents the pseudo code for updating factor matrix \( A_n \). As we mentioned earlier, operator \( \Pi(\Delta(T_n), n) \sim R^{I_1 \times I_2} \rightarrow R^{I_1 \times I_2(-n)} \) executes a sequence of reshape, permute, reshape on \( \Delta(T_n) \in R^{I_1 \times I_2(-n)} \) and converts its size to \( n \times I_2(-n) \). More specifically, line 2 in Algorithm 2 reshapes matrix \( \Delta(T_n) \in R^{I_1 \times I_2(-n)} \) to tensor \( D \in R^{I_1 \times I_2(-n)} \times I_1 \times I_2 \). Line 3-9 permute modes \( i, n \) in \( D \) and map it to \( D_{i \rightarrow n} \in R^{I_1 \times I_2 \times I_1 \times I_2 \times I_2(-n)} \). Depending on the position of \( i \), Lines 3-4 show the permutation when \( i < n \). Lines 5-6 show the case when \( i = n \), and lines 7-8 present the permutation when \( i > n \). Here, \( idx(.) \) returns the index of a given mode. For instance, in \( [I_1, ..., I_{i-1}, I_{i+1}, ..., I_n, I_1] \), \( idx(i-1) = i-1, idx(i) = n, idx(i+1) = i \). In line 10, tensor \( D_{i \rightarrow n}(T_n) \) reshapes to a matrix with size \( I_n \times I_2(-n) \). Once we apply \( \Pi \) on both right and left hand side of (9), we get the new form as follows:

\[ \min_{A_n \geq 0} \sum_{i=1}^N KL(\Pi(\Delta(T_n), n) \mid A_n(A_{\cap}^{-n})^T) \]

(18)
Now we can easily update $A_n$ based on a multiplicative update rule (Lee and Seung 2001) with the following form (line 13 in Algorithm 2):

$$
A_n = A_n \ast \left( \left( \sum_{i=1}^{N} \Pi(\Delta(A_i), n) \odot A_n(A_{-n}^{-n})^T \right) A_{-n}^{-(n)} \right) \odot 1A_{-n}^{-(n)} \right) \tag{19}
$$

\begin{algorithm}
\begin{algorithmic}
\State \textbf{Input} : $\Delta(A_i) \in \mathbb{R}_{+}^{I_i \times I(-i)}$ \quad $\forall i = 1, \ldots, N$, mode-$n$, $P = A_n(A_{-n}^{-n})^T$
\State \textbf{Output} : $A_n \in \mathbb{R}^{I_n \times R}$
\For{$i=1,...,N$}
\State // Reshape a matrix to a tensor.
\State $D_i = \text{reshape}(\Delta(A_i), [I_1, \ldots, I(-1), I(i+1), \ldots, I_N, I_i])$
\State // Interchange the $i$-th and $n$-th modes of the tensor.
\If{$i < n$}
\State $D_{i \rightarrow n} = \text{Permute}(D_i, [\text{idx}(1), \ldots, \text{idx}(i-1), \text{idx}(i), \text{idx}(i+1), \ldots, \text{idx}(n-1), \text{idx}(n+1), \ldots, \text{idx}(N), \text{idx}(n)])$
\ElsIf{$i == n$}
\State $D_{i \rightarrow n} = D_i$
\Else\Comment{Reshape a tensor to a matrix.}
\State $\Pi(\Delta(A_i), n) = \text{reshape}(D_{i \rightarrow n}, [I_n, I(-n)])$
\EndIf
\State $\Pi(A(A_{-i}^{-i})^T, n) = P$
\State $A_n = A_n \ast \left( \left( \sum_{i=1}^{N} \Pi(\Delta(A_i), n) \odot P \right) A_{-n}^{-(n)} \right) \odot 1A_{-n}^{-(n)} \right) $
\EndFor
\end{algorithmic}
\end{algorithm}

C.2 Cost Matrix Calculation

The cost matrices are derived from the same input data without any additional external knowledge.

For BBC NEWS data we compute the following cost matrices:

- **Article \times Article**: The articles are converted to a matrix of TF-IDF (term frequency-inverse document frequency) features. Then the cost matrix between articles $i, j$ is computed based on cosine distance ($C(i, j) = 1 - \frac{\langle a_i, a_j \rangle}{||a_i|| \cdot ||a_j||}$) where $a_i$ is the TF-IDF vector of article $i$.

- **Word \times Word**: For each word, we construct a multi-hot encoding vector with the size of documents in the training set. Then we define the cost matrix as the cosine distance between every pair of vectors.

The cost matrices in Sutter data are:

- **Patient \times Patient**: For each patient we create a vector by concatenating diagnosis and medication features and use cosine distance to compute the cost matrix.

- **Diagnosis \times Diagnosis**: We represent each diagnosis as a multi-hot encoding vector with the size of patients in the training set. If a patient has a certain diagnosis then its corresponding value is one, zero otherwise. We use cosine distance to calculate the cost matrix between every pair of diagnosis vectors.

- **Medication \times Medication**: We perform a similar computation as the Diagnosis \times Diagnosis cost matrix.

Note that for Similarity based CP (Kim et al. 2017), we use the same cost matrices.

C.3 Complexity Analysis of SWIFT

Here, we provide the complexity analysis of different parts of SWIFT. In every iteration, we update $N$ different optimal transport problems and $N$ factor matrices based on Algorithm 1.

In order to update $n$-th optimal transport problem ($T_n$), we need to compute the following parts: $K_n$ requires $O(I_n^2)$ flops (i.e. floating-point operation). Computing $A_{-n}^{-(n)}$ needs $O(I(-n)R)$ and $\hat{X}_{(n)} = A_n(A_{-n}^{-n})^T$ requires $O(I_1 \ldots I_N R)$ flops.

The complexity of computing $U_n = (\hat{X}_{(n)})^\Phi \odot (K_n(X_{(n)} \odot (K^T{n} U_{n}))^\Phi$ (line 8 in Algorithm 1) is $O(I_n^2 N N Z_n)$ where $N N Z_n$ indicates the number of non-zero columns in $X_{(n)}$. Line 10 in Algorithm 1 ($V_n = (X_{(n)} \odot (K^T{n} U_{n}))^\Phi$) requires $O(I_n^2 N N Z_n)$ flops and finally computing $\Delta(T_n) = U_n * (K_n V_n)$ involves $O(I_n^2 N N Z_n)$. 


The steps for updating **Factor matrix** \( n \) are as follows: Both reshape and permute operations can be done in \( O(N) \). Factor matrix \( A_n \) is updated based on (19) which can be computed in \( O(RI_1...I_N) \) flops. Finally, the total complexity of SWIFT is \( O(RI_1...I_N) \).

### C.4 Illustration of the second and third strategies

Figure 3 depicts in detail how the second and third strategies introduced in Section (3.3) explore and utilize the sparsity structure in input tensor in more details.

![Diagram showing computation in parallel](image)

**Figure 3:** SWIFT explores sparsity structure in input data \( X(n) \) and drop zero values columns. Also SWIFT parallelize optimal transport problems for each factor matrix \( A_n \). \( NNZ \) in this toy example equals 3.

### D Additional Experiments and Results

#### D.1 Classification Performance of SWIFT.

**Projection on Learned Factor Matrices** Given the learned factor matrices \( (A_{train_1} , \ldots , A_{train_N}) \) from training data, SWIFT is able to project the new unseen data \( X_{new} \in \mathbb{R}^{I_{new_1} \times I_2 \times \ldots \times I_N} \) into existing factors and learn \( A_{new_1} \).

\[
\begin{align*}
\text{minimize} & \quad \sum_{n=1}^{N} \left( \langle C_n, T_n \rangle - \frac{1}{\rho} E(T_n) \right) \\
\text{Subject to} & \quad T_n \in U(X_{new(n)}, \hat{X}_{new(n)}) \quad \forall n = 1, \ldots, N, \\
& \hat{X} = [A_{new_1}, A_{train_2}, \ldots , A_{train_N}],
\end{align*}
\]

(20)

Here, we need to minimize (20) with respect to \( T_n \), for \( n = 1, \ldots , N \) and \( A_{new_1} \), while \( A_{train_2}, \ldots , A_{train_N} \) are fixed. After learning \( A_{new_1} \), we pass it to Lasso Logistic Regression to predict the labels. We use (20) for projecting the validation and test data.

**Training Strategy:** Assume we want to classify the elements in the first factor matrix \( (A_1) \) (e.g. article mode for BBC News and patient mode for Sutter). For all the approaches under the comparison, we use a similar training strategy with the following steps: 1) we split tensor \( \mathcal{X} \) from it’s first mode into training, validation, and test sets by a ratio of 3:1:1 and construct tensors \( (X_{train}), (X_{val}), \) and \( (X_{test}) \). 2) We train the factorization model using the training set \( (X_{train}) \) and compute \( A_{train_1} \), \( n = 1, \ldots, N \). Note that in this step, the label information is not used (except for the baseline Supervised CP which use label information in the factorization step). 3) Then we freeze the factor matrices of all modes except the one with label information (article mode for BBC News and patient mode for Sutter), and project the validation and test sets onto the learned factor matrices to obtain the factor matrix of the mode with labels for the test set based on Equation (20). 4) Finally we use a lasso logistic regression to perform the classification. We used a five-fold cross validation.

**SWIFT Stopping Criteria** In order to provide scalable solutions, SWIFT never directly computes transport matrices, therefore, it is expensive to compute the value of its objective function. Similar to (Frogner et al. 2015; Cuturi 2013; Qian et al. 2016), we set the number of fixed-point iterations to an arbitrary number. Stopping criterion is set to 50 iterations and Sinkhorn iteration to 25.

**Setting hyper-parameters:** For SWIFT, we execute a grid search for \( \lambda \in \{0.1, 1, 10\} \) and \( \rho \in \{10, 20, 50, 100, 1000\} \) and regularization parameter for lasso logistic regression \( \eta \in \{1e^{-2}, 1e^{-1}, 1, 10, 100, 1000, 10000\} \). The hyper-parameters of other baselines including parameter \( \eta \) are carefully tuned for each dataset.
D.2 Performance on Noisy Sutter Data

Classification Performance on Noisy Input: Figure 4 presents the average and standard deviation of PR-AUC Score of heart failure prediction on test data in Sutter data set for five-fold cross validation with respect to different levels of noise \((0.05, 0.1, 0.15, 0.20, 0.25, 0.30)\). SWIFT performs better under various noise levels and relatively improves the PR-AUC score over the best baseline by up to 17%.

D.3 Scalability of SWIFT on BBC NEWS and Sutter Data

In this section, we answer the following question: Q: How scalable is SWIFT more compared to a direct implementation?

**Baseline Construction:** We implement a direct version of Wasserstein tensor factorization for 3-order tensors where for optimal transport computation we do not leverage sparsity structure. For updating factor matrix \(A_n\) in the direct version of Wasserstein tensor factorization we move \(A_n\) into a similar position by vectorizing every \(\Delta(T)\) and \(A_n\), and rearranged the terms using the properties of Khatri-Rao and Kronecker products, so that \(A_n\) has the same position in all \(N\) different KL-divergence terms. We directly utilize Equations (21),(22), (23) to update \(A_1\), \(A_2\), and \(A_3\), respectively. Note that direct Wasserstein TF returns the same solution given the same initialization as SWIFT.

Each of these objective functions are equivalent to (8), (10) but requires more computation.

\[
\begin{align*}
\min_{A_1 \geq 0} & \quad KL \left( \frac{\text{vec}(\Delta(T_1)^T)}{\text{vec}(\Delta(T_2)^T)} \right) \left| \left| \begin{bmatrix}
(A_3 \odot A_2) \odot I_1 \\
(A_2 \odot I_{I_1}) \odot (I_R \odot A_3) \odot I_{I_1} \\
(A_1 \odot I_{I_1}) \odot (I_R \odot A_2) \odot I_{I_1}
\end{bmatrix} \right| \text{vec}(A_1) \right) \\
\min_{A_2 \geq 0} & \quad KL \left( \frac{\text{vec}(\Delta(T_1)^T)}{\text{vec}(\Delta(T_3)^T)} \right) \left| \left| \begin{bmatrix}
(A_3 \odot A_2) \odot I_{I_2} \\
(A_2 \odot I_{I_2}) \odot (I_R \odot A_3) \odot I_{I_2} \\
(A_1 \odot I_{I_2}) \odot (I_R \odot A_2) \odot I_{I_2}
\end{bmatrix} \right| \text{vec}(A_2) \right) \\
\min_{A_3 \geq 0} & \quad KL \left( \frac{\text{vec}(\Delta(T_1)^T)}{\text{vec}(\Delta(T_3)^T)} \right) \left| \left| \begin{bmatrix}
(A_3 \odot A_2) \odot I_{I_3} \\
(A_2 \odot I_{I_3}) \odot (I_R \odot A_3) \odot I_{I_3} \\
(A_1 \odot I_{I_3}) \odot (I_R \odot A_2) \odot I_{I_3}
\end{bmatrix} \right| \text{vec}(A_3) \right)
\end{align*}
\]

We can easily prove that equations (21),(22), (23) are equivalent to (8), (10) by knowing the following properties:

1. \(KL(A||B) = KL(\text{vec}(A)||\text{vec}(B))\)
2. \(KL(A||B) = KL(A^T||B^T)\)
3. \((E^T \odot C)\text{vec}(D) = \text{vec}(CDE)\) (Neudecker 1969)
4. \(\text{vec}(A_3 \odot A_2) = ((I_R \odot A_3) \odot I_{I_2})\text{vec}(A_2)\) (Roemer 2012).
5. \(\text{vec}(A_2 \odot A_1) = (I_{I_2} \odot (A_1(I_R \odot I_{1 \times I_2})))\text{vec}(A_2)\) (Roemer 2012).
6. \( I_{I_1, I_2} = (I_{I_1} \otimes I_{I_2}) = (I_{I_2} \otimes I_{I_1}) = I_{I_2 I_1} \) (Neudecker 1969).

**Results:** In Figure 5, we vary 1) size of a mode, 2) target rank (R), and 3) number of Sinkhorn iterations in optimal transport problems and report the average and standard deviation of running time (in seconds) for one iteration as an average of five different runs for Sutter and BBC NEWS data sets.

Figures 5a depicts the time as we increase the number of patients (\{500, 750, 1000, 1250\}). SWIFT is up to \(293\times\) faster than the Wasserstein TF model. Execution in the direct Wasserstein TF failed for a tensor of \(1250 \times 100 \times 100\), due to the excessive amount of memory needed. Next, we compare the scalability of both methods by varying the value for \(R\) on Figure 5b. SWIFT achieves up to \(928\times\) faster computation. Figure 5c depicts the optimal transport running time for SWIFT and Wasserstein TF for a tensor with size \(1500 \times 500 \times 500\). Generally, more Sinkhorn iterations lead to a better solution for the optimal transport problems. We increase the value of Sinkhorn iterations and SWIFT achieves up to \(12\times\) speed up over Wasserstein TF suggests that speedup strategies introduced in Section 3.3 are beneficial.

Next, we perform the experiments for BBC NEWS data. Figure 5d demonstrates the running time for one iteration as we increase the number of articles in BBC NEWS data (\{500, 750, 1000, 1250\}). SWIFT is up to \(97\times\) faster than the direct Wasserstein tensor factorization model in Figure 5d. Same as before, execution in the direct Wasserstein TF failed for a tensor of \(1250 \times 100 \times 100\), due to the excessive amount of memory. We measure the scalability of both methods by varying the value for \(R\) (\(R = \{5, 10, 20, 40\}\)) on Figure 5e where SWIFT achieves up to \(718\times\) faster computation. Again for \(R = 40\), direct Wasserstein TF failed to execute due to out of memory. Finally, Figure 5f represents the optimal transport running time for SWIFT and Wasserstein TF for a tensor with size \(1500 \times 500 \times 500\). We increase the value of Sinkhorn iterations (\(\{1, 10, 20, 50\}\)). SWIFT achieves up to \(10\times\) speed up over Wasserstein tensor factorization.

Figure 5 suggests the strategies we introduced in Sections 3.3 and 3.4 are beneficial and significantly reduce the running time of SWIFT in compare to the direct implementation.

---

2We avoid updating factor matrices here and just record the running time of optimal transport problem.
D.4 Effect of Cost Matrix on Classification Performance

In order to understand the power of cost matrix (C) on classification performance, we compare different cost matrices including the one we introduced in section C.2, random and 1-identity (matrix of ones with a zero diagonal) cost matrices for both BBC NEWS and Sutter datasets. When the cost matrix is 1-identity then Wasserstein distance can be seen as total variation distance (Cuturi and Avis 2014).

From Table 5 we summarize our observations as follow: 1) using random cost matrices drop the performance. 2) 1-identity cost matrix performs better than random cost matrices but still not as good as cost matrices introduced in Section (C.2). As you can see SWIFT with 1-identity cost matrices have better performance than other well-known CP baselines (Comparing with Table 1).

Table 5: Average and standard deviation of accuracy on BBC NEWS and PR-AUC score on Sutter data sets by considering different cost matrices for different values of target-rank ($R = [5, 10, 20, 30, 40]$).

<table>
<thead>
<tr>
<th></th>
<th>R=5</th>
<th>R=10</th>
<th>R=20</th>
<th>R=30</th>
<th>R=40</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>BBC NEWS</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SWIFT (Random)</td>
<td>.621 ± .041</td>
<td>.693 ± .053</td>
<td>.684 ± .038</td>
<td>.712 ± .047</td>
<td>.684 ± .022</td>
</tr>
<tr>
<td>SWIFT (1-Identity)</td>
<td>.709 ± .016</td>
<td>.775 ± .019</td>
<td>.793 ± .032</td>
<td>.775 ± .023</td>
<td>.771 ± .013</td>
</tr>
<tr>
<td>SWIFT</td>
<td><strong>.759 ± .013</strong></td>
<td><strong>.781 ± .013</strong></td>
<td><strong>.803 ± .010</strong></td>
<td><strong>.815 ± .005</strong></td>
<td><strong>.818 ± .022</strong></td>
</tr>
<tr>
<td><strong>Sutter</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>SWIFT (Random)</td>
<td>.340 ± .103</td>
<td>.347 ± .091</td>
<td>.341 ± .083</td>
<td>.352 ± .041</td>
<td>.354 ± .074</td>
</tr>
<tr>
<td>SWIFT (1-Identity)</td>
<td>.363 ± .105</td>
<td>.349 ± .079</td>
<td>.344 ± .073</td>
<td>.358 ± .063</td>
<td>.368 ± .074</td>
</tr>
<tr>
<td>SWIFT</td>
<td><strong>.364 ± .063</strong></td>
<td><strong>.350 ± .031</strong></td>
<td><strong>.350 ± .040</strong></td>
<td><strong>.369 ± .066</strong></td>
<td><strong>.374 ± .044</strong></td>
</tr>
</tbody>
</table>

D.5 More Results on HF Phenotyping

We show the remaining phenotypes positively associated with HF in Table 6.
Table 6: The phenotypes positively associated with HF. All phenotypes are annotated and endorsed by a medical expert. The unlisted ones are either negatively or not associated with HF.

<table>
<thead>
<tr>
<th>HF with long-term Diabetes (Weight= 14.62)</th>
<th>HF with Dysrhythmias (Weight= 13.73)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Dx-Diabetes with ketoacidosis or uncontrolled diabetes</td>
<td>Dx-Carotid dysrhythmias [106.]</td>
</tr>
<tr>
<td>Dx-Other mycoses</td>
<td>Dx-Diabetes mellitus without complication [49.]</td>
</tr>
<tr>
<td>Dx-Diabetes mellitus without complication [49.]</td>
<td>Dx-Other viral infections [7.]</td>
</tr>
<tr>
<td>Rx-Corticosteroids - Topical</td>
<td>Rx-Coumarin Anticoagulants</td>
</tr>
<tr>
<td>Rx-Biguanides</td>
<td>Rx-Nonsteroidal anti-inflammatory agents</td>
</tr>
<tr>
<td>Rx-Central Muscle Relaxants</td>
<td>Rx-Calcium Channel Blockers</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>CAD related HF (Weight= 10.04)</th>
<th>Aging and frail related HF (Weight= 9.72)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Dx-Carotid dysrhythmias [106.]</td>
<td>Dx-Urinary tract infections [159.]</td>
</tr>
<tr>
<td>Dx-Acute cerebrovascular disease [109.]</td>
<td>Dx-Genitourinary symptoms and ill-defined conditions [163.]</td>
</tr>
<tr>
<td>Dx-Disorders of lipid metabolism [53.]</td>
<td>Dx-Other bone disease and musculoskeletal deformities [212.]</td>
</tr>
<tr>
<td>Rx-Coumarin Anticoagulants</td>
<td>Rx-Urinary Anti-infectives</td>
</tr>
<tr>
<td>Rx-Oil Soluble Vitamins</td>
<td>Rx-Fluoroquinolones</td>
</tr>
<tr>
<td>Rx-HMG CoA Reductase Inhibitors</td>
<td>Rx-Anti-infective Misc. - Combinations</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>HF with Pulmonary disease (Weight= 9.58)</th>
<th>HF with air pathway blockage (Weight= 8.98)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Dx-Other and unspecified asthma</td>
<td>Dx-Essential hypertension [98.]</td>
</tr>
<tr>
<td>Dx-Chronic airway obstruction; not otherwise specified</td>
<td>Dx-Allergic reactions [253.]</td>
</tr>
<tr>
<td>Dx-Acute bronchitis [125.]</td>
<td>Dx-Esophageal disorders [138.]</td>
</tr>
<tr>
<td>Rx-Sympathomimetics</td>
<td>Rx-Calcium Channel Blockers</td>
</tr>
<tr>
<td>Rx-Fluoroquinolones</td>
<td>Rx-Beta Blockers Cardio-Selective</td>
</tr>
<tr>
<td>Rx-Opioid Combinations</td>
<td>Rx-Proton Pump Inhibitors</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>HF with systematic inflammation (Weight= 6.45)</th>
<th>HF with thyroid dysfunction (Weight= 4.08)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Dx-Essential hypertension [98.]</td>
<td>Dx-Immunizations and screening for infectious disease [10.]</td>
</tr>
<tr>
<td>Dx-Chronic kidney disease [158.]</td>
<td>Dx-Essential hypertension [98.]</td>
</tr>
<tr>
<td>Dx-Disorders of lipid metabolism [53.]</td>
<td>Dx-Other thyroid disorders</td>
</tr>
<tr>
<td>Rx-ACE Inhibitors</td>
<td>Rx-Antihypertensive Combinations</td>
</tr>
<tr>
<td>Rx-HMG CoA Reductase Inhibitors</td>
<td>Rx-Impotence Agents</td>
</tr>
<tr>
<td>Rx-Calcium Channel Blockers</td>
<td>Rx-Angiotensin II Receptor Antagonists</td>
</tr>
</tbody>
</table>