
Online Graph Dictionary Learning

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Abstract

Dictionary learning is a key tool for representation learning, that explains the data as linear combination of few basic elements. Yet, this analysis is not amenable in the context of graph learning, as graphs usually belong to different metric spaces. We fill this gap by proposing a new online Graph Dictionary Learning approach, which uses the Gromov Wasserstein divergence for the data fitting term. In our work, graphs are encoded through their nodes' pairwise relations and modeled as convex combination of graph atoms, *i.e.* dictionary elements, estimated thanks to an online stochastic algorithm, which operates on a dataset of unregistered graphs with potentially different number of nodes. Our approach naturally extends to labeled graphs, and is completed by a novel upper bound that can be used as a fast approximation of Gromov Wasserstein in the embedding space. We provide numerical evidences showing the interest of our approach for unsupervised embedding of graph datasets and for online graph subspace estimation and tracking.

challenging, as their nature is by essence non-vectorial, and requires dedicated modelling of their representing structures. Given sufficient data and labels, end-to-end approaches with neural networks have shown great promises in the last years (Wu et al., 2020). In this work, we focus on the unsupervised representation learning problem, where the entirety of the data might not be known beforehand, and is rather produced continuously by different sensors, and available through streams. In this setting, tackling the non-stationarity of the underlying generating process is challenging (Ditzler et al., 2015). Good examples can be found, for instance, in the context of dynamic functional connectivity (Heitmann & Breakspear, 2018) or network science (Masuda & Lambiotte, 2020). As opposed to recent approaches focusing on dynamically varying graphs in online or continuous learning (Yang et al., 2018; Vlaski et al., 2018; Wang et al., 2020), we rather suppose in this work that *distinct* graphs are made progressively available (Zambon et al., 2017; Grattarola et al., 2019). This setting is particularly challenging as the structure, the attributes or the number of nodes of each graph observed at a time step can differ from the previous ones. We propose to tackle this problem by learning a linear representation of graphs with online dictionary learning.

1. Introduction

The question of how to build machine learning algorithms able to go beyond vectorial data and to learn from structured data such as graphs has been of great interest in the last decades. Notable applications can be found in molecule compounds (Kriege et al., 2018), brain connectivity (Ktena et al., 2017), social networks (Yanardag & Vishwanathan, 2015), time series (Cuturi & Blondel, 2018), trees (Day, 1985) or images (Harchaoui & Bach, 2007; Bronstein et al., 2017). Designing good representations for these data is

Dictionary Learning (DL) Dictionary Learning (Mairal et al., 2009; Schmitz et al., 2018) is a field of unsupervised learning that aims at estimating a linear representation of the data, *i.e.* to learn a linear subspace defined by the span of a family of vectors, called *atoms*, which constitute a *dictionary*. These atoms are inferred from the input data by minimizing a reconstruction error. These representations have been notably used in statistical frameworks such as data clustering (Ng et al., 2002), recommendation systems (Bobadilla et al., 2013) or dimensionality reduction (Candès et al., 2011). While DL methods mainly focus on vectorial data, it is of prime interest to investigate flexible and interpretable factorization models applicable to *structured data*. We also consider the dynamic or time varying version of the problem, where the data generating process may exhibit non-stationarity over time, yielding a problem of subspace change or tracking (see *e.g.* (Narayanamurthy & Vaswani, 2018)), where one wants to monitor changes in the subspace best describing the data. In this work, we rely on optimal transport as a fidelity term to compare these structured data.

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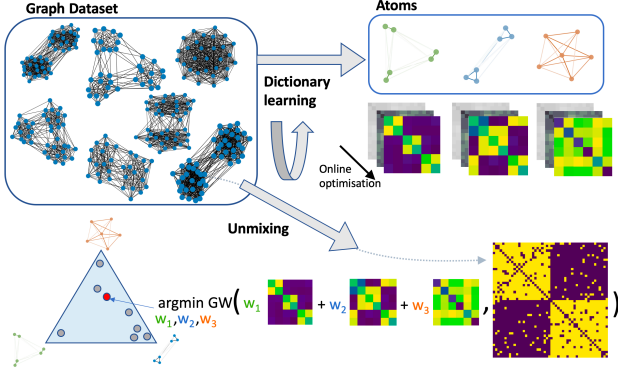


Figure 1. From a dataset of graphs with different number of nodes, our method builds a dictionary of graph atoms with an online procedure. It uses the Gromov-Wasserstein distance as data fitting term between a convex combination of the atoms and a pairwise relations representation for graphs from the dataset.

Optimal Transport for structured data Optimal Transport (OT) theory provides a set of methods for comparing probability distributions, using, *e.g.* the well-known Wasserstein distance (Villani, 2003). It has been notably used by the machine learning community in the context of distributional unsupervised learning (Arjovsky et al., 2017; Schmitz et al., 2018; Peyré & Cuturi, 2019). Broadly speaking the interest of OT lies in its ability to provide correspondences, or relations, between sets of points. Consequently, it has recently garnered attention for learning tasks where the points are described by graphs/structured data (see *e.g.* (Nikolentzos et al., 2017; Maretic et al., 2019; Togninalli et al., 2019; Xu et al., 2019a; Vayer et al., 2019; Barbe et al., 2020)). One of the key ingredient in this case is to rely on the so called Gromov-Wasserstein (GW) distance (Mémoli, 2011; Sturm, 2012) which is an OT problem adapted to the scenario in which the supports of the probability distributions lie in different metric spaces. The GW distance is particularly suited for comparing *relational data* (Peyré et al., 2016; Solomon et al., 2016) and, in a graph context, is able to find the relations between the nodes of two graphs when their respective structure is encoded through the pairwise relationship between the nodes in the graph. GW has been further studied for weighted directed graphs in (Chowdhury & Mémoli, 2019) and has been extended to labeled graphs thanks to the Fused Gromov-Wasserstein (FGW) distance in (Vayer et al., 2018). Note that OT divergences as losses for linear and non-linear DL over vectorial data have already been proposed in (Bonneel et al., 2016; Rolet et al., 2016; Schmitz et al., 2018) but the case of structured data remains quite unaddressed. A non-linear DL approach for graphs based on GW was proposed in (Xu, 2020) but suffers from a lack of interpretability and high computational complexity (see discussions in Section 3). To the best of our knowledge, a linear counterpart does not exist for now.

Contributions In this paper we use OT distances between structured data to design a linear and online DL for undirected graphs. Our proposal is depicted in Figure 1. It consists in a new factorization model for undirected graphs optionally having node attributes relying on (F)GW distance as data fitting term. We propose an online stochastic algorithm to learn the dictionary which scales to large real-world data (Section 2.3), and uses extensively novel derivations of sub-gradients of the (F)GW distance (Section 2.4). An unmixing procedure projects the graph in an embedding space defined *w.r.t.* the dictionary (Section 2.2). Interestingly enough, we prove that the GW distance in this embedding is upper-bounded by a Mahalanobis distance over the space of *unmixing* weights, providing a reliable and fast approximation of GW (Section 2.1). Moreover, this approximation defines a proper kernel that can be efficiently used for clustering and classification of graphs datasets (sections 4.1-4.2). We empirically demonstrate the relevance of our approach for online subspace estimation and subspace tracking by designing streams of graphs over two datasets (Section 4.3).

Notations The simplex of histograms with N bins is $\Sigma_N := \{\mathbf{h} \in \mathbb{R}_N^+ \mid \sum_i h_i = 1\}$. Let denote $S_N(\mathbb{R})$ the set of symmetric matrices in $\mathbb{R}^{N \times N}$. The Euclidean norm is denoted as $\|\cdot\|_2$ and $\langle \cdot, \cdot \rangle_F$ the Frobenius inner product. We denote the gradient of a function f over x at y in a stochastic context by $\tilde{\nabla}_x f(y)$. The number nodes in a graph is called the *order* of the graph.

2. Online Graph Dictionary Learning

2.1. (Fused) Gromov-Wasserstein for graph similarity

A graph G^X with N^X nodes, can be regarded as a tuple $(C^X, \mathbf{h}^X)$ where $C^X \in \mathbb{R}^{N^X \times N^X}$ is a matrix that encodes a notion of similarity between nodes and $\mathbf{h}^X \in \Sigma_{N^X}$ is a histogram, or equivalently a vector of weights which models the relative importance of the nodes within the graph. Without any prior knowledge uniform weights can be chosen so that $\mathbf{h}^X = \frac{1}{N^X} \mathbf{1}_{N^X}$. The matrix C^X carries the neighbourhood information of the nodes and, depending on the context, it may designate the adjacency matrix, the Laplacian matrix (Maretic et al., 2019) or the matrix of the shortest-path distances between the nodes (Bavaud, 2010). Let us now consider two graphs $G^X = (C^X, \mathbf{h}^X)$ and $G^Y = (C^Y, \mathbf{h}^Y)$, of potentially different orders (*i.e* $N^X \neq N^Y$). The GW_2 distance between G^X and G^Y is defined as the result of the following optimization problem:

$$\min_{T \in \mathcal{U}(\mathbf{h}^X, \mathbf{h}^Y)} \sum_{ijkl} (C_{ij}^X - C_{kl}^Y)^2 T_{ik} T_{jl} \quad (1)$$

where $\mathcal{U}(\mathbf{h}^X, \mathbf{h}^Y) := \{T \in \mathbb{R}_+^{N^X \times N^Y} \mid T \mathbf{1}_{N^Y} = \mathbf{h}^X, T^T \mathbf{1}_{N^X} = \mathbf{h}^Y\}$ is the set of couplings between $\mathbf{h}^X, \mathbf{h}^Y$. The optimal coupling T of the GW problem acts

as a probabilistic matching of nodes which tends to associate pairs of nodes that have similar pairwise relations in C^X and C^Y , respectively. In the following we denote by $GW_2(C^X, C^Y, h^X, h^Y)$ the optimal value of equation 1 or by $GW_2(C^X, C^Y)$ when the weights are uniform.

The previous framework can be extended to graphs with node attributes (typically $\mathbb{R}^d$ vectors). In this case we use the Fused Gromov-Wasserstein distance (FGW) (Vayer et al., 2018; 2019) instead of GW. More precisely, a labeled graph G^X with N^X nodes can be described this time as a tuple $G^X = (C^X, A^X, h^X)$ where $A^X \in \mathbb{R}^{N^X \times d}$ is the matrix of all features. Given two labeled graphs G^X and G^Y , FGW aims at finding an optimal coupling by minimizing an OT cost which is a trade-off of a Wasserstein cost between the features and a GW cost between the similarity matrices. For the sake of clarity, we detail our approach in the GW context and refer the reader to the supplementary material for its extension to FGW.

2.2. Linear embedding and GW unmixing

Linear modeling of graphs We propose to model a graph as a weighted sum of pairwise relation matrices. More precisely, given a graph $G = (C, h)$ and a dictionary $\{\bar{C}_s\}_{s \in [S]}$ we want to find a linear representation $\sum_{s \in [S]} w_s \bar{C}_s$ of the graph G , as faithful as possible. The dictionary is made of pairwise relation matrices of graphs with order N . Thus, each $\bar{C}_s \in S_N(\mathbb{R})$ is called an *atom*, and $w = (w_s)_{s \in [S]} \in \Sigma_S$ is referred as *embedding* and denotes the coordinate of the graph G in the dictionary as illustrated in Fig.1. We rely on the GW distance to assess the quality of our linear approximation and propose to minimize it to estimate its optimal embedding. In addition to being interpretable thanks to its linearity, we also propose to promote sparsity in the weights w similarly to sparse coding (Chen et al., 2001). Finally note that, when the pairwise matrices C are adjacency matrices and the dictionary atoms have components in $[0, 1]$, the model $\sum_{s \in [S]} w_s \bar{C}_s$ provides a matrix whose components can be interpreted as probabilities of connection between the nodes.

Gromov-Wasserstein unmixing We first study the unmixing problem that consists in projecting a graph on the linear representation discussed above, *i.e.* estimate the optimal embedding w of a graph G . The unmixing problem can be expressed as the minimization of the GW distance between the similarity matrix associated to the graph and its linear representation in the dictionary:

$$\min_{w \in \Sigma_S} GW_2^2 \left(C, \sum_{s \in [S]} w_s \bar{C}_s \right) - \lambda \|w\|_2^2 \quad (2)$$

where $\lambda \in \mathbb{R}^+$ induces a **negative** quadratic regularization promoting sparsity on the simplex as discussed in Li et al.

Algorithm 1 BCD for unmixing problem 2

- 1: Initialize $w = \frac{1}{S} \mathbf{1}_S$
 - 2: **repeat**
 - 3: Compute OT matrix T of $GW_2^2(C, \sum_s w_s \bar{C}_s)$, with CG algorithm (Vayer et al., 2018, Alg.1 & 2).
 - 4: Compute the optimal w solving equation 2 for a fixed T with CG algorithm.
 - 5: **until** convergence
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(2016). In order to solve the non-convex problem in equation 2, we propose to use a Block Coordinate Descent (BCD) algorithm (Tseng, 2001).

The BCD (Alg.1) works by alternatively updating the OT matrix of the GW distance and the embeddings w . When w is fixed the problem is a classical GW which is a non-convex quadratic program. We solve it using a Conditional Gradient (CG) algorithm (Jaggi, 2013) based on (Vayer et al., 2019). Note that the use of the exact GW instead of a regularized proxy allowed us to keep a sparse OT matrix as well as to preserve “high frequency” components of the graph, as opposed to regularized versions of GW (Peyré et al., 2016; Solomon et al., 2016; Xu et al., 2019b) that promotes dense OT matrices and leads to smoothed/averaged pairwise matrices. For a fixed OT matrix T , the problem of finding w is a non-convex quadratic program and can also be tackled with a CG algorithm. Note that for non-convex problems the CG algorithm is known to converge to a local stationary point (Lacoste-Julien, 2016). In practice, we observed a typical convergence of the CGs in a few tens of iterations. The BCD itself converges in less than 10 iterations.

Fast upper bound for GW Interestingly, when two graphs belong to the linear subspace defined by our dictionary, there exists a proxy of the GW distance using a dedicated Mahalanobis distance as described in the next proposition:

Proposition 1 For two embedded graphs with embeddings $w^{(1)}$ and $w^{(2)}$, assuming they share the same weights h , the following inequality holds

$$GW_2 \left(\sum_{s \in [S]} w_s^{(1)} \bar{C}_s, \sum_{s \in [S]} w_s^{(2)} \bar{C}_s \right) \leq \|w^{(1)} - w^{(2)}\|_M \quad (3)$$

where $M_{pq} = \langle D_h \bar{C}_p, \bar{C}_q D_h \rangle_F$ and $D_h = \text{diag}(h)$. M is a positive semi-definite matrix hence engenders a Mahalanobis distance between embeddings.

As detailed in the supplementary material, this upper bound is obtained by considering the GW cost between the linear models calculated using the admissible coupling D_h . The

latter coupling assumes that both graph representations are aligned and therefore is a priori suboptimal. As such, this bound is not tight in general. However, when the embeddings are close, the optimal coupling matrix should be close to D_h so that Proposition 1 provides a reasonable proxy to the GW distance into our embedding space. In practice, this upper bound can be used to compute efficiently pairwise kernel matrices or to do retrieval of closest samples (see numerical experiments).

2.3. Dictionary learning and online algorithm

Assume now that the dictionary $\{\bar{C}_s\}_{s \in [S]}$ is not known and has to be estimated from the data. We define a dataset of K graphs $\{G^{(k)} : (C^{(k)}, h^{(k)})\}_{k \in [K]}$. Recall that each graph $G^{(k)}$ of order $N^{(k)}$ is summarized by its pairwise relation matrix $C^{(k)} \in S_{N^{(k)}}(\mathbb{R})$ and weights $h^{(k)} \in \Sigma_{N^{(k)}}$ over nodes, as described in Section 2.1. The DL problem, that aims at estimating the optimal dictionary for a given dataset can be expressed as:

$$\min_{\substack{\{w^{(k)}\}_{k \in [K]} \\ \{\bar{C}_s\}_{s \in [S]}}} \sum_{k=1}^K GW_2^2 \left(C^{(k)}, \sum_{s \in [S]} w_s^{(k)} \bar{C}_s \right) - \lambda \|w^{(k)}\|_2^2 \quad (4)$$

where $w^{(k)} \in \Sigma_S$, $\bar{C}_s \in S_N(\mathbb{R})$. Note that the optimization problem above is a classical sparsity promoting dictionary learning on a linear subspace but with the important novelty that the reconstruction error is computed with the GW distance. This allows us to learn a graphs subspace of fixed order N using a dataset of graphs with various orders. The sum over the errors in equation 4 can be seen as an expectation and we propose to devise an online strategy to optimize the problem similarly to the online DL proposed in (Mairal et al., 2009). The main idea is to update the dictionary $\{\bar{C}_s\}_s$ with a stochastic estimation of the gradients on few dataset graphs (minibatch). At each stochastic update the unmixing problems are solved independently for each graph of the minibatch using a fixed dictionary $\{\bar{C}_s\}_s$, using the procedure described in Section 2.2. Then one can compute a gradient of the loss on the minibatch w.r.t $\{\bar{C}_s\}_s$ and proceed to a projected gradient step. The stochastic update of the proposed algorithm is detailed in Alg.2. Note that it can be used on a finite dataset with possibly several epochs on the whole dataset or online in the presence of streaming graphs. We provide an example of such subspace tracking in Section 4.3. We will refer to our approach as GDL in the rest of the paper.

Numerical complexity The numerical complexity of GDL depends on the complexity of each update. The main computational bottleneck is the unmixing procedure that relies on multiple resolution of GW problems. The complexity

Algorithm 2 GDL: stochastic update of atoms $\{\bar{C}_s\}_{s \in [S]}$

- 1: Sample a minibatch of graphs $\mathcal{B} := \{C^{(k)}\}_{k \in \mathcal{B}}$.
- 2: Compute optimal $\{(w^{(k)}, T^{(k)})\}_{k \in [B]}$ by solving B independent unmixing problems with Alg.1.
- 3: Projected gradient step with estimated gradients $\tilde{\nabla}_{\bar{C}_s}$ (equation in supplementary), $\forall s \in [S]$:

$$\bar{C}_s \leftarrow Proj_{S_N(\mathbb{R})}(\bar{C}_s - \eta_C \tilde{\nabla}_{\bar{C}_s}) \quad (5)$$

of solving a GW with the CG algorithm between two graphs of order N and M and computing its gradient is dominated by $\mathcal{O}(N^2M + M^2N)$ operations (Peyré et al., 2016; Vayer et al., 2018). Thus given dictionary atoms of order N , the worst case complexity can be only **quadratic** in the highest graph order in the dataset. For instance, estimating embedding on dataset IMDB-M (see Section 4.2) over 12 atoms takes on average 44 ms per graph (on processor i9-9900K CPU 3.60GHz). We refer the reader to the supplementary for more details. Note that in addition to scale well to large datasets thanks to the stochastic optimization, our method also leads to important speedups when using the representations as input feature for other ML tasks. For instance, we can use the upper bound in equation 3 to compute efficiently kernels between graphs instead of computing all pairwise GW distances.

GDL on labeled graphs We can also define the same DL procedure for labeled graphs using the FGW distance. The unmixing part defined in equation 2 can be adapted by considering a linear embedding of the similarity matrix *and* of the feature matrix parametrized by the *same* w . From an optimization perspective, finding the optimal coupling of FGW can be achieved using a CG procedure so that Alg.2 extends naturally to the FGW case. Note also that the upper bound of Proposition 1 can be generalized to this setting. This discussion is detailed in supplementary material.

2.4. Learning the graph structure and distribution

Recent researches have studied the use of potentially more general distributions h on the nodes of graphs than the naive uniform ones commonly used. (Xu et al., 2019a) empirically explored the use of distributions induced by degrees, such as parameterized power laws, $h_i = \frac{p_i}{\sum_i p_i}$, where $p_i = (deg(x_i) + a)^b$ with $a \in \mathbb{R}_+$ and $b \in [0, 1]$. They demonstrated the interest of this approach but also highlighted how hard it is to calibrate, which advocates for learning these distributions. With this motivation, we extend our GDL model defined in equation 4 and propose to learn atoms of the form $\{\bar{C}_s, \bar{h}_s\}_{s \in [S]}$. In this setting we have two independent dictionaries modeling the relative importance of the nodes with $\bar{h}_s \in \Sigma_N$, and their pairwise

relations through $\bar{C}_s$. This dictionary learning problem reads:

$$\min_{\substack{\{(\mathbf{w}^{(k)}, \mathbf{v}^{(k)})\}_{k \in [K]} \\ \{(\bar{C}_s, \bar{\mathbf{h}}_s)\}_{s \in [S]}}} \sum_{k=1}^K GW_2^2 \left(\mathbf{C}^{(k)}, \tilde{C}(\mathbf{w}^{(k)}), \mathbf{h}^{(k)}, \tilde{\mathbf{h}}(\mathbf{v}^{(k)}) \right) - \lambda \|\mathbf{w}^{(k)}\|_2^2 - \mu \|\mathbf{v}^{(k)}\|_2^2 \quad (6)$$

where $\mathbf{w}^{(k)}, \mathbf{v}^{(k)} \in \Sigma_S$ are the structure and distribution embeddings and the linear models are defined as:

$$\forall k, \tilde{\mathbf{h}}(\mathbf{v}^{(k)}) = \sum_s v_s^{(k)} \bar{\mathbf{h}}_s, \quad \tilde{C}(\mathbf{w}^{(k)}) = \sum_s w_s^{(k)} \bar{C}_s$$

Here we exploit fully the GW formalism by estimating simultaneously the graph distribution $\tilde{\mathbf{h}}$ and its geometric structure $\tilde{C}$. Optimization problem 6 can be solved by an adaptation of stochastic Algorithm 2. We estimate the structure/node weights unmixings $(\mathbf{w}^{(k)}, \mathbf{v}^{(k)})$ over a minibatch of graphs with a BCD (see Section 2.3). Then we perform simultaneously a projected gradient step update of $\{\bar{C}_s\}_s$ and $\{\bar{\mathbf{h}}_s\}_s$. More details are given in the supplementary.

The optimization procedure above requires to have access to a gradient for the GW distance *w.r.t.* the weights. To the best of our knowledge no theoretical results exists in the literature for finding such gradients. We provide below a simple way to compute a subgradient for GW weights from subgradients of the well-known Wasserstein distance:

Proposition 2 *Let $(C^1, \mathbf{h}^1)$ and $(C^2, \mathbf{h}^2)$ be two graphs. Let T^* be an optimal coupling of the GW problem between $(C^1, \mathbf{h}^1), (C^2, \mathbf{h}^2)$. We define the following cost matrix $M(T^*) := \left(\sum_{kl} (C_{ik}^1 - C_{jl}^2)^2 T_{kl}^* \right)_{ij}$. Let $\alpha^*(T^*), \beta^*(T^*)$ be the dual variables of the following linear OT problem:*

$$\min_{T \in \mathcal{U}(\mathbf{h}^1, \mathbf{h}^2)} \langle M(T^*), T \rangle_F \quad (7)$$

Then $\alpha^(T^*)$ (resp $\beta^*(T^*)$) is a subgradient of the function $GW_2^2(C^1, C^2, \cdot, \mathbf{h}^2)$ (resp $GW_2^2(C^1, C^2, \mathbf{h}^1, \cdot)$).*

The proposition above shows that the subgradient of GW *w.r.t.* the weights can be found by solving a linear OT problem which corresponds to a Wasserstein distance. The ground cost $M(T^*)$ of this Wasserstein is moreover the gradient (*w.r.t.* the couplings) of the optimal GW loss. Note that in practice the GW problem is solved with a CG algorithm which already requires to solve this linear OT problem at each iteration. In this way, after convergence, the gradient *w.r.t.* the weights can be extracted for free from the last iteration of the CG algorithm. The proof of proposition 2 is given in the supplementary material.

3. Related work and discussion

In this section we discuss the relation of our GDL framework with existing approaches designed to handle graph data. We first focus on existing contributions for graph representation in machine learning applications. Then, we discuss in more details the existing non-linear graph dictionary learning approach of (Xu, 2020).

Graph representation learning Processing of graph data in machine learning applications have traditionally been handled using implicit representations such as with graph kernels (Shervashidze et al., 2009; Vishwanathan et al., 2010). Recent results have shown the interest of using OT based distances to measure graph similarities and to design new kernels (Vayer et al., 2019; Maretic et al., 2019; Chowdhury & Needham, 2020). However, one limit of kernel methods is that the representation of the graph is fixed *a priori* and cannot be adapted to specific datasets. On the other hand, Geometric deep learning approaches (Bronstein et al., 2017) attempt to learn the representation for structured data by means of deep learning (Scarselli et al., 2008; Perozzi et al., 2014; Niepert et al., 2016). Graph Neural Networks (Wu et al., 2020) have shown impressive performance for end-to-end supervised learning problems. Note that both kernel methods and many deep learning based representations for graphs suffer from the fundamental *pre-image* problem, that prevents recovering actual graph objects from the embeddings. Our proposed GDL aims at overcoming such a limit relying on an unmixing procedure that not only provides a simple vectorial representation on the dictionary but also allows a direct reconstruction of interpretable graphs (as illustrated in the experiments). A recent contribution potentially overcoming the pre-image problem is Grattarola et al. (2019). In that paper, a variational autoencoder is indeed trained to embed the observed graphs into a constant curvature Riemannian manifold. The aim of that paper is to represent the graph data into a space where the statistical tests for change detection are easier. We look instead for a latent representation of the graphs that remains as interpretable as possible. As a side note, we point out that our GDL embeddings might be used as input for the statistical tests developed by (Zambon et al., 2017; 2019) to detect stationarity changes in the stochastic process generating the observed graphs (see for instance Figure 6).

Non-linear GW dictionary learning of graphs In a recent work, (Xu, 2020) proposed a non-linear factorization of graphs using a regularized version of GW barycenters (Peyré et al., 2016) and denoted it as Gromov-Wasserstein Factorization (GWF). Authors propose to learn a dictionary $\{\bar{C}_s\}_{s \in [S]}$ by minimizing over $\{\bar{C}_s\}_{s \in [S]}$ and $\{\mathbf{w}^{(k)}\}_{k \in [K]}$ the quantity $\sum_{k=1}^K GW_2^2(\tilde{B}(\mathbf{w}^{(k)}; \{\bar{C}_s\}_s), C^{(k)})$ where $\tilde{B}(\mathbf{w}^{(k)}; \{\bar{C}_s\}_s) \in \arg \min_B \sum_s w_s^{(k)} GW_2^2(B, \bar{C}_s)$ is a

GW barycenter. The main difference between GDL and this work lies in the linear representation of the approximated graph that we adopt whereas (Xu, 2020) relies on the highly non-linear Gromov barycenter. As a consequence, the unmixing requires solving a complex bi-level optimization problem that is computationally expensive. Similarly, reconstructing a graph from this embedding requires again the resolution of a GW barycenter, whereas our linear reconstruction process is immediate. In Section 4, we show that our GDL representation technique compares favorably to GWF, both in terms of numerical complexity and performance.

4. Numerical experiments

This section aims at illustrating the behavior of the approaches introduced so far for both clustering (Sections 4.1-4.2) and online subspace tracking (Section 4.3).

Implementation details The base OT solvers that are used in the algorithms rely on the POT toolbox (Flamary & Courty, 2017). For our experiments, we considered the Adam algorithm (Kingma & Ba, 2014) as an adaptive strategy for the update of the atoms with a fixed dataset, but used SGD with constant step size for the online experiments in Section 4.3. The code is available at <https://github.com/cedricvincentcuaz/GDL>.

4.1. GDL on simulated datasets

The GDL approach discussed in this section refers to equation 4. First we illustrate it on datasets simulated according to the well understood Stochastic Block Model (SBM, Holland et al., 1983; Wang & Wong, 1987) and show that we can recover embeddings and dictionary atoms corresponding to the generative structure.

Datasets description We consider two datasets of graphs, generated according to SBM, with various orders, randomly sampled in $\{10, 15, \dots, 60\}$. The first scenario (D_1) adopts three different generative structures (also referred to as *classes*): dense (no clusters), two clusters and three clusters (see Figures 2). Nodes are assigned to clusters into equal proportions. For each generative structure 100 graphs are sampled. The second scenario (D_2) considers the generative structure with two clusters, but with varying proportions of nodes for each block (see top of Figure 3), 150 graphs are simulated accordingly. In both scenarios we fix $p = 0.1$ as the probability of inter-cluster connectivity and $1 - p$ as the probability of intra-cluster connectivity. We consider adjacency matrices for representing the structures of the graphs in the datasets and uniform weights on the nodes.

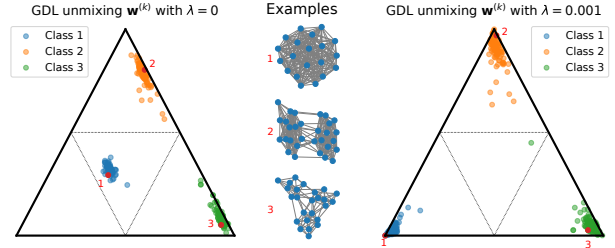


Figure 2. Visualizations of the embeddings of the graphs from D_1 with our GDL on 3 atoms. The positions on the simplex for the different classes are reported with no regularization (left) and sparsity promoting regularization (right). Three simulated graphs from D_1 are shown in the middle and their positions on the simplex reported in red.

Results and interpretation First we learn on dataset D_1 a dictionary of 3 atoms of order 6. The unmixing coefficients for the samples in D_1 are reported in Fig. 2. On the left, we see that the coefficients are not sparse on the simplex but the samples are clearly well clustered and graphs sharing the same class (i.e. color) are well separated. When adding sparsity promoting regularization (right part of the figure) the different classes are clustered on the corners of the simplex, thus suggesting that regularization leads to a more discriminant representation. The estimated atoms for the regularized GDL are reported on the top of Fig. 1 as both matrices $\bar{C}_s$ and their corresponding graphs. As it can be seen, the different SBM structures in D_1 are recovered. Next we estimate on D_2 a dictionary with 2 atoms of order 12. The interpolation between the two estimated atoms for some samples is reported in Fig. 3. As it can be seen, D_2 can be modeled as a one dimensional manifold where the proportion of nodes in each block changes continuously. We stress that the grey links on the bottom of Figure 3 correspond to the entries of the reconstructed adjacency matrices. Those entries are in $[0, 1]$, thus encoding a probability of connection (see Section 2.2). The darker the link, the higher the probability of interaction between the corresponding nodes. The possibility of generating random graphs using these probabilities opens the door to future researches.

We evaluate in Fig. 4 the quality of the Mahalanobis upper bound in equation 3 as a proxy for the GW distance on D_1 . On the left, one can see that the linear model allows us to recover the true GW distances between graphs most of the time. Exceptions occur for samples in the same class (i.e. "near" to each other in terms of GW distance). The right part of the figure shows that the correlation between the Mahalanobis upper bound (cf. Proposition 1) and the GW distance between the embedded graphs is nearly perfect (0.999). This proves that our proposed upper bound provides a nice approximation of the GW distance between the input graphs, with a correlation of 0.96 (middle of the figure), at a much lower computational cost.

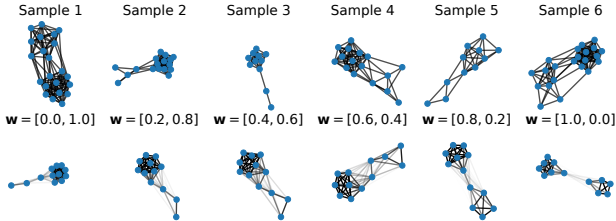


Figure 3. On the top, a random sample of real graphs from D_2 (two blocks). On the bottom, reconstructed graphs as linear combination of two estimated atoms (varying proportions for each atom).

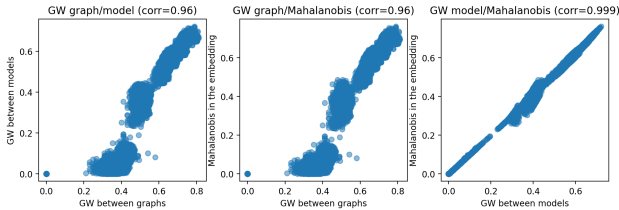


Figure 4. Plot of the pairwise distances in D_1 and their Pearson correlation coefficients. GW distance between graphs versus its counterpart between the embedded graphs (left). GW distance between graphs versus Mahalanobis distance between the embeddings (middle). GW distance between the embedded graphs versus Mahalanobis between the corresponding embeddings (right).

4.2. GDL on real data for clustering and classification

We now show how our *unsupervised* GDL procedure can be used to find meaningful representations for well-known graph classification datasets. The knowledge of the classes will be employed as a ground truth to validate our estimated embeddings in *clustering* tasks. For the sake of completeness, in supplementary material we also report the supervised classification accuracies of some recent supervised graph *classification* methods (e.g. GNN, kernel methods) showing that our DL and embedding is also competitive for classification.

Datasets and methods We considered well-known benchmark datasets divided into three categories: i) IMDB-B and IMDB-M (Yanardag & Vishwanathan, 2015) gather graphs without node attributes derived from social networks; ii) graphs with discrete attributes representing chemical compounds from MUTAG (Debnath et al., 1991) and cuneiform signs from PTC-MR (Krichene et al., 2015); iii) graphs with real vectors as attributes, namely BZR, COX2 (Sutherland et al., 2003) and PROTEINS, ENZYMES (Borgwardt & Kriegel, 2005). We benchmarked our models for clustering tasks with the following state-of-the-art OT models: i) GWF (Xu, 2020), using the proximal point algorithm detailed in that paper and exploring two configurations, i.e. with either fixed atom order (GWF-f) or random atom order (GWF-r, default for the method); ii) GW k-means (GW-k) which is

a k-means using GW distances and GW barycenter (Peyré et al., 2016); iii) Spectral Clustering (SC) of (Shi & Malik, 2000; Stella & Shi, 2003) applied to the pairwise GW distance matrices or the pairwise FGW distance matrices for graphs with attributes. We complete these clustering evaluations with an ablation study of the effect of the negative quadratic regularization proposed with our models. As introduced in equation 4, this regularization is parameterized by λ , so in this specific context we will distinguish GDL ($\lambda = 0$) from GDL_λ ($\lambda > 0$).

Experimental settings For the datasets with attributes involving FGW, we tested 15 values of the trade-off parameter α via a logspace search in $(0, 0.5)$ and symmetrically $(0.5, 1)$ and select the one minimizing our objectives. For our GDL methods as well as for GWF, a first step consists into learning the atoms. A variable number of $S = \beta k$ atoms is tested, where k denotes the number of classes and $\beta \in \{2, 4, 6, 8\}$, with a uniform number of atoms per class. When the order N of each atom is fixed, for GDL and GWF-f, it is set to the median order in the dataset. The atoms are initialized by randomly sampling graphs from the dataset with corresponding order. We tested 4 regularization coefficients for both methods. The embeddings w are then computed and used as input for a k-means algorithm. However, whereas a standard Euclidean distance is used to implement k-means over the GWFs embeddings, we use the Mahalanobis distance from Proposition 1 for the k-means clustering of the GDLs embeddings. Unlike GDL and GWF, GW-k and SC do not require any embedding learning step. Indeed, GW-k directly computes (a GW) k-means over the input graphs and SC is applied to the GW distance matrix obtained from the input graphs. The cluster assignments are assessed by means of Rand Index (RI, Rand, 1971), computed between the true class assignment (known) and the one estimated by the different methods. For each parameter configuration (number of atoms, number of nodes and regularization parameter) we run each experiment five times, independently, with different random initializations. The mean RI was computed over the random initializations and the dictionary configuration leading to the highest RI was finally retained.

Results and interpretation Clustering results can be seen in Table 1. The mean RI and its standard deviation are reported for each dataset and method. Our model outperforms or is at least comparable to the state-of-the-art OT based approaches for most of the datasets. Results show that the negative quadratic regularization proposed with our models brings additional gains in performance. Note that for this benchmark, we considered a fixed batch size for learning our models on labeled graphs, which turned out to be a limitation for the dataset ENZYMES. Indeed, comparable conclusions regarding our models performance have been

Table 1. Clustering: Rand Index computed for benchmarked approaches on real datasets.

MODELS	NO ATTRIBUTE		DISCRETE ATTRIBUTES		REAL ATTRIBUTES			
	IMDB-B	IMDB-M	MUTAG	PTC-MR	BZR	COX2	ENZYMES	PROTEIN
GDL (ours)	51.32(0.30)	55.08(0.28)	70.02(0.29)	51.53(0.36)	62.59(1.68)	58.39(0.52)	66.97(0.93)	60.22(0.30)
GDL _λ (ours)	51.64(0.59)	55.41(0.20)	70.89(0.11)	51.90(0.54)	66.42(1.96)	59.48(0.68)	66.79(1.12)	60.49(0.71)
GWf-r	51.24 (0.02)	55.54(0.03)	68.83(1.47)	51.44(0.52)	52.42(2.48)	56.84(0.41)	72.13(0.19)	59.96(0.09)
GWf-f	50.47(0.34)	54.01(0.37)	58.96(1.91)	50.87(0.79)	51.65(2.96)	52.86(0.53)	71.64(0.31)	58.89(0.39)
GW-k	50.32(0.02)	53.65(0.07)	57.56(1.50)	50.44(0.35)	56.72(0.50)	52.48(0.12)	66.33(1.42)	50.08(0.01)
SC	50.11(0.10)	54.40(9.45)	50.82(2.71)	50.45(0.31)	42.73(7.06)	41.32(6.07)	70.74(10.60)	49.92(1.23)

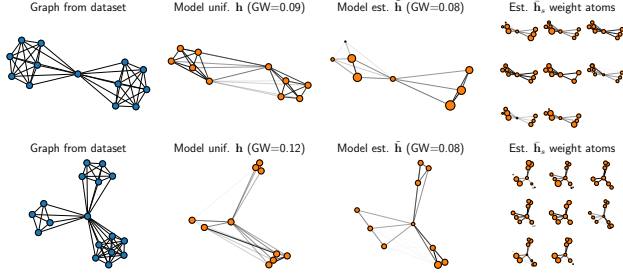


Figure 5. Modeling of two real life graphs from IMDB-M with our GDL approaches with 8 atoms of order 10. (left) original graphs from the dataset, (center left) linear model for GDL with uniform weights as in equation 4, (center right) linear model for GDL with estimated weights as in equation 6 and (right) different $\bar{h}_s$ on the estimated structure.

observed by setting a higher batch size for this latter dataset and are reported in the supplementary material. This might be due to both a high number of heterogeneous classes and a high structural diversity of labeled graphs inside and among classes.

We illustrate in Fig. 5 the interest of the extension of GDL with estimated weights for IMDB-M dataset. We can see in the center-left part of the figure that, without estimating the weights, GDL can experience difficulties producing a model that preserves the global structure of the graph because of the uniform weights on the nodes. In opposition, simultaneously estimating the weights brings a more representative modeling (in the GW sense), as illustrated in the centred-right columns. The weights estimation can re-balance and even discard non relevant nodes, in the vein of attention mechanisms. We report in the supplementary material a companion study for clustering tasks which further supports our extension concerning the learning of node weights.

4.3. Online graph subspace estimation and change detection

Finally we provide experiments for online graph subspace estimation on simulated and real life datasets. We show that our approach can be used for subspace tracking of graphs as well as for change point detection of subspaces.

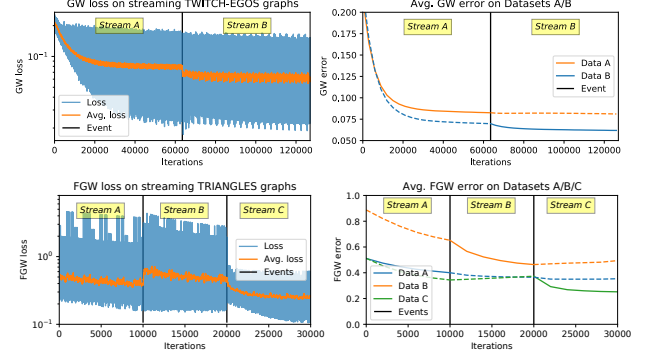


Figure 6. Online GDL on dataset TWITCH-EGOS with 2 atoms of 14 nodes each (top) and on TRIANGLES with 4 atoms of 17 nodes each (bottom).

Datasets and experiments In this section we considered two new large graph classification datasets: TWITCH-EGOS (Rozemberczki et al., 2020) containing social graphs without attributes belonging to 2 classes and TRIANGLES (Knyazev et al., 2019) that is a simulated dataset of labeled graphs with 10 classes. Here we investigate how our approach fits to online data, *i.e.* in the presence of a stream of graphs. The experiments are designed with different time segments where each segment streams graphs belonging to the same classes (or group of classes). The aim is to see if the method learns the current stream and detects or adapts to abrupt changes in the stream. For TWITCH-EGOS, we first streamed all graphs of a class (A), then graphs of the other class (B), both counting more than 60.000 graphs. All these graphs consist in a unique high-frequency (a hub structure) with sparse connections between non-central nodes (sparser for class B). For TRIANGLES, the stream follows the three groups A,B and C, with 10,000 graphs each, where the labels associated with each group are: $A = \{4, 5, 6, 7\}$, $B = \{8, 9, 10\}$ and $C = \{1, 2, 3\}$.

Results and discussion The online (F)GW losses and a running mean of these losses are reported for each dataset on the left part of Fig. 6. On the right part of the Figure, we report the average losses computed on several datasets containing data from each stream at some time instant along the iterations. First, the online learning for both datasets can be seen in the running means with a clear decrease of loss on

each time segment. Also, note that at each event (change of stream) a jump in terms of loss is visible suggesting that the method can be used for change point detection. Finally it is interesting to see on the TRIANGLES dataset that while the loss on Data B is clearly decreased during Stream B it increases again during Stream C, thus showing that our algorithm performs subspace tracking, adapting to the new data and forgetting old subspaces no longer necessary.

5. Conclusion

We present a new *linear* Dictionary Learning approach for graphs with different orders relying on the Gromov Wasserstein (GW) divergence, where graphs are modeled as convex combination of graph atoms. We design an online stochastic algorithm to efficiently learn our dictionary and propose a computationally light proxy to the GW distance in the described graphs subspace. Our experiments on clustering classification and online subspace tracking demonstrate the interest of our unsupervised representation learning approach. We envision several extensions to this work, notably in the context of graph denoising or graph inpainting.

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