

Exploiting Edge Features in Graph-based Learning with Fused Network **Gromov-Wasserstein Distance**

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Motivation

Problem: Graph-based Learning with Optimal Transport

Existing Works: Graph Classification with FGW [5], Graph Clustering with GW [4], FGW [5] or NGW [2], Graph Dictionary Learning with (F)GW [6], Supervised Graph Prediction with FGW [1]

Goal: We want to unlock OT-based learning for edge featured graphs. We target especially Supervised Graph Prediction task.



Supervised Graph Prediction with FNGW

Given input space \mathcal{X} , output graph space \mathcal{G} , relaxed graph space $\mathcal{G}_m = \{(F, A, E, \mathbf{p}) \mid C \in \mathcal{G}\}$ $[0,1]^{m \times m}, F = (F_i)_{i=1}^m \in \operatorname{Conv}(\mathcal{F})^m, E = (E_{ij}) \in \operatorname{Conv}(\mathcal{T})^{m \times m}, p = m^{-1}\mathbb{1}_m\}$ where $\mathcal{F} \subset \mathbb{R}^S$ and $\mathcal{T} \subset \mathbb{R}^T$ are finite node and edge feature spaces, and training samples $\{(x_i, g_i)\}_{i=1}^n$, Supervised Graph Prediction requires finding an estimator $f: \mathcal{X} \to \mathcal{G}_m$ of the minimizer f^* of the expected risk $\mathcal{R}(f) = \mathbb{E}_{\rho}[\text{FNGW}_{\alpha,\beta}(f(X),G)]$. Based on the work of [3, 1], we propose an estimator of the form

$$\hat{f}(x) = \underset{g \in \mathcal{G}_m}{\operatorname{arg\,min}} \sum_{i=1}^{n} \xi(x)_i \operatorname{FNGW}_{\alpha,\beta}(g,g_i)$$

with $\xi(x) = \mathbf{K}S^{\mathsf{T}}(S\mathbf{K}^2S^{\mathsf{T}} + n\lambda S\mathbf{K}S^{\mathsf{T}})^{\dagger}S\boldsymbol{\kappa}_x$ where $\mathbf{K} \in \mathbb{R}^{n \times n}$ is the input kernel Gram matrix, $\boldsymbol{\kappa}_x = (k(x, x_1), \dots, k(x, x_n))^{\mathsf{T}} \in \mathbb{R}^n$, and $S \in \mathbb{R}^{s \times n}$ with $s \ll n$ is a sketching matrix.

• The FNGW loss admits an Implicit Loss Embedding (ILE) $\rightarrow \hat{f}$ is universally consistent and its learning rate is of order $n^{-1/4}$ with additional assumptions.

Experiment: Fingerprint to Molecule

Node and Edge Featured Graph

A node and edge featured graph of size m is a quadruple of the form (F, A, E, p) where

- $F \in \Psi^m$ is a tuple of points valued in a metric space (Ψ, d_{Ψ})
- $A \in \mathbb{R}^{m \times m}$ is a real-valued matrix
- $E \in \Omega^{m \times m}$ is a tuple of points valued in a metric space (Ω, d_{Ω}) • $p \in \Sigma_m$ is a simplex histogram

We denote \mathcal{G} as a set of such quadruples.



Fused Network Gromov-Wasserstein Distance

Given $g = (F, A, E, \mathbf{p})$ of size $m, \tilde{g} = (\tilde{F}, \tilde{A}, \tilde{E}, \tilde{\mathbf{p}})$ of size \tilde{m} corresponding to two tuples of \mathcal{G} , and trade-off parameters $(\alpha, \beta) \in [0, 1]^2$, the Fused Network Gromov-Wasserstein distance between them for $(p,q) \in [1,\infty]$ is written as :





GED w/o edge feature \downarrow	GED w/ edge feature 、
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NNBary-FGW NNBary-FNGW	5.000 ± 0.140 5.311 ± 0.090	-5.756 ± 0.073
Sketched ILE-FGW Sketched ILE-FNGW	3.037 ± 0.111 1.449 \pm 0.034	-1.534 ± 0.029

Table. Graph edit distances of different methods on the Fin2Mol test set.



$$\operatorname{YNGW}_{\alpha,\beta,q,p}(g,\tilde{g}) = \min_{\pi \in \Pi(\boldsymbol{p},\tilde{\boldsymbol{p}})} \mathcal{E}_{\alpha,\beta,q,p}((F,A,E),(\tilde{F},\tilde{A},\tilde{E}),\pi)$$

with

$$\mathcal{E}_{\alpha,\beta,q,p}((F,A,E),(\tilde{F},\tilde{A},\tilde{E}),\pi) = \left(\sum_{i,j,k,l} \left[\alpha d_{\Omega} \left(E(i,k),\tilde{E}(j,l)\right)^{q} + \beta |A(i,k) - \tilde{A}(j,l)|^{q} + (1-\alpha-\beta) d_{\Psi} \left(F(i),\tilde{F}(j)\right)^{q}\right]^{p} \pi_{k,l}\pi_{i,j}\right)^{\frac{1}{p}}$$

The FNGW distance satisfies the following **metric** properties: positivity, symmetry, equality with a corresponding notion of weak isomorphism, relaxed triangle inequality with a factor of 2^{q-1} .



FNGW Barycenter

Given a set $\{g_k\}_{k=1}^K$ and a set of weights $\{\lambda_k\}_{k=1}^K$ such that $\sum_k \lambda_k = 1$, the FNGW Barycenter for a pre-defined histogram $\boldsymbol{p} \in \Sigma_n$ is defined as follows:

Figure. Qualitative comparison of the predicted QM9 molecules.

Experiment: Metabolite Identification

To solve Metabolite Identification problem, the learning algorithm is expected to predict the metabolite (small molecules) given a tandem mass spectra. For each input spectra, a known set of metabolite candidates is provided.

	Top-1	Top-10	Top-20
WL kernel	9.8%	29.1%	37.4%
Fingerprint with linear kernel	28.6%	54.5%	59.9%
Fingerprint with gaussian kernel	41.0 %	62.0 %	67.8 %
FGW diffuse	28.1%	53.6%	59.9%
FNGW diffuse + Bond stereo	27.7%	55.2%	$\begin{array}{c} 60.9\% \\ 60.0\% \\ 61.9\% \end{array}$
FNGW diffuse + Bond type	34.6%	55.1%	
FNGW diffuse + Mix	36.2%	58.2%	

Table. Top-k accuracies on the metabolite identification test set.

$Bary(\{\lambda_k\}_k, \{g_k\}_k, \boldsymbol{p}) = \operatorname*{arg\,min}_{F \in \mathbb{R}^{n \times S}, A \in \mathbb{R}^{n \times n}, E \in \mathbb{R}^{n \times n \times T}} \sum_k \lambda_k FNGW_{\alpha, \beta}((F, A, E, \boldsymbol{p}), g_k)$

We employ the Block Coordinate Descent (BCD) algorithm to obtain the FNGW barycenter where the tensor E can be updated by $E = \frac{1}{\mathcal{I}_{n \times T} \times_2 p p^{\mathsf{T}}} \sum_k \lambda_k (E_k \times_2 \pi_k) \times_1 \pi_k.$



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