Soutenance de thèse.

Quelques contributions à l'analyse statistique de données à structure de graphe.

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Theme of this thesis : graphs.



Co-authorship Network [PBV07]



Protein-protein interaction network





[BO04] : Network biology: understanding the cell's functional organization., Barabasi *et al.*, 2008 [HCG+08] : Mapping the structural core of human cerebral cortex, Hagmann *et al.*, 2008 [PBV07] : Quantifying social group evolution, Palla *et al.*, 2007

Objectives

Goals :

• comparison of graph samples

$$(G_1,\ldots,G_N) \stackrel{i.i.d.}{\sim} P \quad ; \qquad (G'_1,\ldots,G'_M) \stackrel{i.i.d.}{\sim} Q$$

• theoretical results (asymptotic in sample size)

Requirements :

- take into account topological information
- graphs can be weighted
- graph sizes (same/different)
- node correspondance (known/unknown)

Outline

1. Heat diffusion distance processes [L21]

- Distances
- Processes
- 2. Functional central limit theorem and beyond [L21]
 - Donsker theorem and Gaussian approximation
 - Two-sample test
- 3. Detecting distribution shift
 - Experiments with MNIST
 - Experiments with Ripsnet

[L21] Heat diffusion distance processes: a statistically founded method to analyze graph data sets, L. , 2021, (Journal of Applied and Computational Topology).

Mathematical representations.



$$(1) (2) (3) (4) (5) (6)$$

$$(1) (0 1 0 0 0 0) (1 0 1 0 1 0 0 0) (1 0 1 0 1 1 1) (1 0 0 0 1 0) (1 0 1 0) (1 0 0) (1 0 0 1 0) (1$$

Mathematical representations.



$$(1) (2) (3) (4) (5) (6)$$

$$(1) \begin{pmatrix} 0 & 2 & 0 & 0 & 0 & 0 \\ 2 & 0 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 1 & 2 & 2 \\ 0 & 0 & 1 & 0 & 3 & 0 \\ (5) \begin{pmatrix} 0 & 0 & 2 & 3 & 0 & 3 \\ 0 & 0 & 2 & 0 & 3 & 0 \end{pmatrix} = W$$

$$(6) Weight Matrix$$

Mathematical representations.



$$(1) (2) (3) (4) (5) (6)$$

$$(1) \begin{pmatrix} 0 & 2 & 0 & 0 & 0 & 0 \\ 2 & 0 & 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 1 & 2 & 2 \\ 0 & 0 & 1 & 0 & 3 & 0 \\ 0 & 0 & 2 & 3 & 0 & 3 \\ (6) \begin{pmatrix} 0 & 0 & 2 & 3 & 0 & 3 \\ 0 & 0 & 2 & 0 & 3 & 0 \end{pmatrix} = W$$

Weight Matrix

$$\begin{bmatrix} 0 & 0 \\ 0 & 0 \\ -2 & -2 \\ -3 & 0 \end{bmatrix} \qquad u^T L u = \sum_{i \neq j} w_{i,j} (u_i - u_j)^2$$

$$L := D - W = \begin{pmatrix} 2 & -2 & 0 & 0 & 0 & 0 \\ -2 & 3 & -1 & 0 & 0 & 0 \\ 0 & -1 & 6 & -1 & -2 & -2 \\ 0 & 0 & -1 & 4 & -3 & 0 \\ 0 & 0 & -2 & -3 & 8 & -3 \\ 0 & 0 & -2 & 0 & -3 & 5 \end{pmatrix}$$

Laplacian Matrix

Heat equation :

 $u_0 \in \mathbb{R}^n$: initial heat distribution.

$$\frac{d}{dt}u_t = -Lu_t, \quad t \ge 0$$



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Comparing graphs

Assumption :

same sizes n & known node correspondance.

Compare matrix representations :

- Adjacency / weight matrix
- Laplacian matrix
- Heat kernel

Heat Kernel Distance :

$$D_t(G, G') = \|e^{-tL} - e^{-tL'}\|_F \quad [\text{HGJ13}]$$

[HGJ13] : Graph diffusion distance: A difference measure for weighted graphs based on the graph 7 - 1 Laplacian exponential kernel, Hammond *et al*, 2013

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Heat Kernel Distance :



[HGJ13] : Graph diffusion distance: A difference measure for weighted graphs based on the graph 7 - 3 Laplacian exponential kernel, Hammond *et al*, 2013

Using Topological Data Analysis



[CCIL+19]: Perslay : A Neural Network Layer for Persistence Diagrams and New Graph Topological Signatures, Carriere, Chazal, Ike, Lacombe, Royer, Umeda, 2019

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Comparing persistence diagrams



- μ , ν : finite multisets of points in \mathbb{R}^2 . $\Delta = \{(a, a), \forall a \in \mathbb{R}\}$: diagonal
- π : a matching from $\mu \cup \Delta$ to $\nu \cup \Delta$
- $\Pi(\mu, \nu)$: set of all matchings

Bottleneck Distance :
$$d_B(\mu, \nu) := \inf_{\pi \in \Pi(\mu, \nu)} \sup_{x \in \mu \cup \Delta} \|x - \pi(x)\|_{\infty}$$

Another way to compare graphs

Heat Kernel Signature (HKS) : [SOG09] [HRG14]

 $h_t(G): i \to (e^{-tL})_{i,i}$ "Remaining heat at node i"



Heat Persistence Distance (HPD) :

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$$H_t(G, G') = \max_{D_g} d_B \left(Dg \left(G, h_t(G) \right), Dg \left(G', h_t(G') \right) \right)$$

[SOG09]: A concise and provably informative multiscale signature based on heat diffusion, Sun *et al*, 2009 [HRG14] : Stable and informative spectral signatures for graph matching, Hu *et al*, 2014

Recap of the distances.

Assumption : same sizes n & known node correspondance.

Heat Kernel Distance (HKD):

$$D_t(G, G') = \|e^{-tL} - e^{-tL'}\|_F$$

Assumption : No assumption.

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How can we choose t ?

To choose or not to choose?

Functional Point of View								
$D_{\cdot}(G,G'):$	[0,T]	\mapsto	\mathbb{R}	or	$H_{\cdot}(G,G'):$	[0,T]	\mapsto	\mathbb{R}
	t	\mapsto	$D_t(G,G')$			t	\mapsto	$H_t(G,G')$

Empirical Process Point of View					
$\{D_t(G, G'), t \in [0, T]\}$	or	$\{H_t(G, G'), t \in [0, T]\}$			
$\mathcal{F}_{HKD} = \{ D_t(\cdot), t \in [0, T] \}$	or	$\mathcal{F}_{HPD} = \{ H_t(\cdot), t \in [0,T] \}$			

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2. Functional central limit theorem and beyond.

General empirical processes.

- $X_1, \ldots, X_N \sim P$ (i.i.d sample) $X_i \in \mathcal{X}$
- P_N : empirical measure
- $\mathcal{F} = \{f_t, t \in [0,T]\}$: a family of measurable functions.

t fixed,

$$\sqrt{N}(P_N - P)f_t = \sqrt{N} \left(\frac{1}{N} \sum_{i=1}^N f_t(X_i) - \mathbb{E}_P \left[f_t(X) \right] \right)$$

$$\xrightarrow{\mathcal{D}} \mathcal{N}(0, \sigma_t^2)$$

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Definition : \mathcal{F} is said to be *Donsker* if $\{\sqrt{N}(P_N - P)f_t, t \in [0, T]\} \xrightarrow{weak}$ Gaussian Process \mathbb{G} $\forall h : \mathcal{C}([0, T]) \to \mathbb{R}$, continuous and bounded $\lim_{N \to \infty} \mathbb{E} \left[h \left(\sqrt{N}(P_N - P)f_{\cdot} \right) \right] = \mathbb{E} \left[h(\mathbb{G}) \right]$

Definition : $\{G_N f_t, t \in [0, T]\}$ admits a *Gaussian approximation* with rate r_N , if $\forall \lambda > 1, \exists \rho, N_0$, such that $\forall N \ge N_0$ one can construct on the same probability space X_1, \ldots, X_N and a version of the Gaussian limit process $\mathbb{G}^{(N)}$ such that

Main theoretical result

Assumptions :

(L) - $\exists k > 0, \forall x \in \mathcal{X}, t \to f_t(x) \text{ is } k\text{-Lipschitz continuous}$ (B) - $\exists M > 0, \forall x \in \mathcal{X}, \forall t \in [0, T], |f_t(x)| \leq M$

Result :

- \mathcal{F} is Donsker
- $\{G_N f_t, t \in [0,T]\}$ admits a Gaussian approximation with

$$r_N = N^{-1/7} (\log N)^{9/14}$$

L21

Refinement :

•
$$\rho = B_1 + B_2 \sqrt{\lambda + 1} + B_3 M (kT + M)^{5/2} (\lambda + 2/7)$$

• $N_0 = \mathcal{O}\left(M^{7+u} + (kT+M)^{2+u}\right)$

Applications : [L21]

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- $\forall (G, G')$ of size n, with weights in $[0, w_{\max}]$, $t \to D_t(G, G')$ is $n^{3/2} w_{\max}$ -lipschitz continuous and bounded by \sqrt{n}
- $\forall (G, G')$ of size at most n, with weights in $[0, w_{\max}]$, $t \to H_t(G, G')$ is $2nw_{\max}$ -lipschitz continuous and bounded by 1.

Two-sample Test (for pairs of graphs)



 $c) \leq \alpha$

Idea : compute
$$T_{N,M} = \frac{\sqrt{NM}}{\sqrt{N+M}} \|P_N D - Q_M D\|_{\infty}$$
.

(- - -

- reject \mathcal{H}_0 , if $T_{N,M} > c$
- retain \mathcal{H}_0 , otherwise

Two-sample Test (for pairs of graphs)

Idea : estimate *c* by resampling.

$$T_{N,M} = \frac{\sqrt{NM}}{\sqrt{N+M}} \|P_N D_{\cdot} - Q_M D_{\cdot}\|_{\infty}$$
$$\hat{T}_{N,M} := \frac{\sqrt{NM}}{\sqrt{N+M}} \|\hat{P}_N D_{\cdot} - \hat{Q}_M D_{\cdot}\|_{\infty}$$
resampled from
$$Z = (X_1, \dots, X_N, Y_1, \dots, Y_M)$$

Two-sample Test (for pairs of graphs)

Idea : estimate c by resampling.



 $ilde{m{c}}$: estimation of the lpha-upper quantile of $\hat{T}_{N,M}|Z|$

 $\lim_{N,M\to\infty} \mathbb{P}_{\mathcal{H}_0} \left(T_{N,M} \ge \tilde{c} \right) \le \alpha$ 17 - 2

if
$$PD_{\cdot} \neq QD_{\cdot}$$
, $\lim_{N,M\to\infty} \mathbb{P}_{\mathcal{H}_1} \left(T_{N,M} \geq \tilde{c} \right) = 1$

Simulations : Stochastic Models









Simulations : Two-sample Tests



Level 95%, bootstrap sample size : 1000, number of tests : 400



HKD

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HPD



: splitting data











By bootstrap : we find $\tilde{\boldsymbol{c}}$ such that if P=Q

$$\lim_{N \to \infty} \mathbb{P}\left(T > \tilde{c}\right) \le \alpha.$$

- If $T > \tilde{c}$, conclude $P \neq Q$.
- If $T \leq \tilde{c}$, conclude P = Q.

Detecting distribution shift.

Distribution shift

A supervized ML algorithm in development :

- Data : train set + test set
- From the train set, learn a function (e.g. classifier)
- From the test set, evaluate the trained function.

Will the trained function perform well once deployed in the "real world"?

Distribution shift

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- Data : train set + test set
- From the train set, learn a function (e.g. classifier)
- From the test set, evaluate the trained function.

Will the trained function perform well once deployed in the "real world"?

- Enhance "real world" performances. (Robustness, data augmentation, ...)
- Detect a potential shift of distribution.

development data
$$\sim P$$

"real world" data $\sim Q$
 $\left. \begin{array}{c} \bullet & \text{If } P = Q, \text{ ok! } \checkmark \\ \bullet & \text{Else, } P \neq Q, \text{ risk of poor behavior!} \end{array} \right.$

Can we detect if the distribution has shifted?

22 - 2 Even better: can we detect if the algorithm is performing poorly?





 $\sigma: \mathbb{R} \to \mathbb{R}$: activation function (fixed) $w_j \in \mathbb{R}^p$, NN weights (trained) $b_j \in \mathbb{R}$, bias. (trained)





Can we use the underlying graph structure to detect distribution shifts ?

Activation Graphs

Consider :

- a trained neural network,
- one data instance.



- weighted graph
- same vertex set

• characterize the processing of the data

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Consider :

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- one data instance.



• same vertex set

• characterize the processing of the data



The corruptions :



• HD

• HDy (pairing by label)

• HDmt (repeat pairing / multiple testing)

• hammond (use $\max_t D_t(G, G')$)



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• BBSD [LWS18]

26 - 2 [LWS18] Detecting and Correcting for Label Shift with Black Box Predictors, Lipton *et al*, 2018.



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• HD

Compare the means of $D_{\cdot}(G_i^k, G_{\pi(i)}^{k'})$ using $\|\cdot\|_{\infty}$, and the functional CLT.

• HDy (pairing by label)

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Compare the means of $\|D_{\cdot}(G_{i}^{k},G_{\pi(i)}^{k'})\|_{\infty}$ using the standard CLT.

• BBSD [LWS18]

26 - 6 [LWS18] Detecting and Correcting for Label Shift with Black Box Predictors, Lipton *et al*, 2018.

• HD

• HDy (pairing by label)

• HDmt (repeat pairing / multiple testing)



• hammond (use $\max_{t} D_t(G, G')$)



For each output neuron :

For both samples, extract the output neuron values.

Perform a Kolmogorov-Smirnov two-sample test.

Combine tests with Bonferroni procedure (multiple tests).

26 - 7 [LWS18] Detecting and Correcting for Label Shift with Black Box Predictors, Lipton et al, 2018.

Results

<u>Neural Network :</u> Dense layers, 3 hidden layers (16 neurons). Accuracy : \simeq 0.96

Repeat 200x {

1st sample : original 2nd sample : corrupted Compute the test.







Results (full data)



50% increase of the error rate

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Ripsnet [dSHC+22]

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Ripsnet :
$$X = \{x_1, ..., x_k\} \mapsto \phi_2 (\mathbf{op} (\{\phi_1(x_i)\}_{1 \le i \le k})),$$

 $\begin{aligned} \phi_1 : \mathbb{R}^d \to \mathbb{R}^{d'} & \text{dense neural network} \\ \phi_2 : \mathbb{R}^{d'} \to \mathbb{R}^K & \text{dense neural network} \\ \mathbf{op} : & \text{permutation invariant operator ($ *i.e.* $, mean)} \end{aligned}$



[dSHC+22] RipsNet: a general architecture for fast and robust estimation of the persistent homology of point clouds, De Surrel *et al*, 2022.

The data



Results





Publication :

L., 2021, Heat diffusion distance processes: a statistically founded method to analyze graph data sets, arXiv:2109.13213 (Journal of Applied and Computational Topology).

Perspectives :

- Theoretical study of the test power (special cases)
- Interplay between graph size and sample size
- Extensions to classical learning tasks on graphs (clustering, classification, outlier detection, change-point detection for time series)
- Study of neural networks (over-fitting, over-parametrization, ...)

ΤΗΑΝΚ ΥΦΊ ΓΦΚ ΥΦΊΚ ΑΤΤΕΝΤΙΦΝΙ

Simulations : Two-sample Tests

Neyman-Pearson regime :

sample of size NNeyman-Pearson test : $ER(p_1(N))$ vs $ER(p_2(N))$

 $|p_1(N) - p_2(N)| \gg 1/\sqrt{N}$



Influence of the graph sizes

ER-ER vs ER-SBM



Results (CNN)

Neural Network :

Convolution layers + 3 dense layers (32 neurons).

Accuracy : $\simeq 0.989$

Repeat 200x





Ripsnet [dSHC+22]



[dSHC+22] RipsNet: a general architecture for fast and robust estimation of the persistent homology of point clouds, De Surrel *et al*, 2022.