

Persistent Homology Approximations on Network Distances

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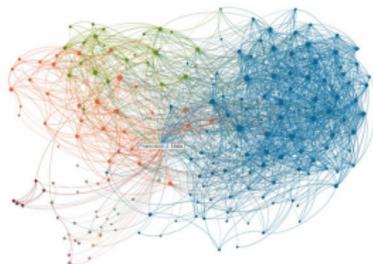
December 16, 2015

GlobalSIP'2015

Internet



Online social media

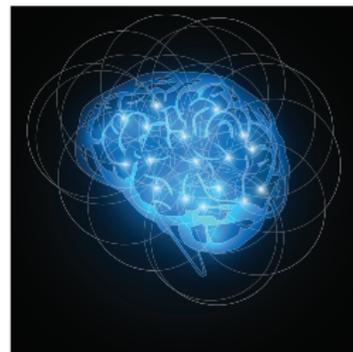


Clean energy & grid

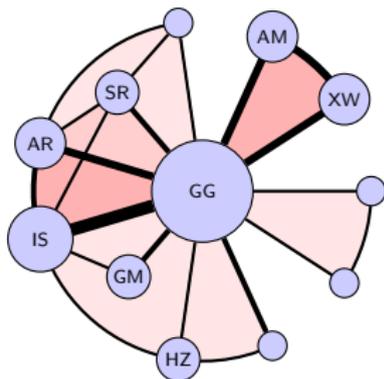


- ▶ Networked data structures encode **relationships** between elements
- ▶ How to evaluate **dissimilarities** between networks remain unclear

- ▶ Neurodegenerate
 - ▶ Association with brain network
 - ▶ Feature heuristics so far
 - ▶ Not specific to a region of the brain
 - ▶ But more about **global properties**
 - ▶ Need to be compared as **unlabeled** entities

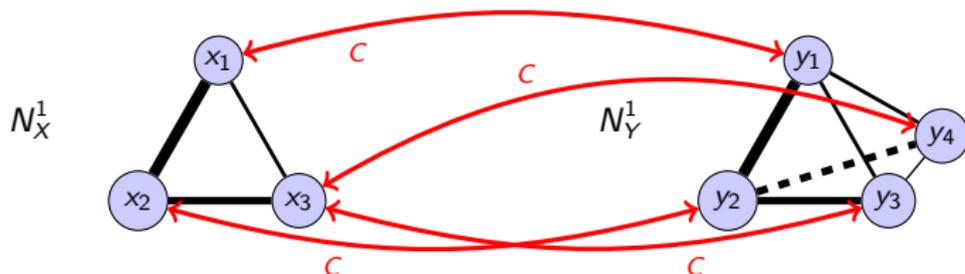


- ▶ Define and estimate **distances** between unlabeled networks
 - ▶ More generalizable
 - ▶ Universal and avoid conflicting statement
- ▶ High order networks
 - ▶ Relationships between three or four nodes, or even singleton



- ▶ Network $N_X^K = (X, r_X^0, r_X^1, \dots, r_X^K)$
 - $\Rightarrow r_X^k$ is a mapping $\overbrace{X \times X \cdots \times X}^{k+1} \rightarrow \mathbb{R}_+$
 - $\Rightarrow r_X^2(x_1, x_2, x_2) = r_X^1(x_1, x_2)$
- ▶ Proximity network P_X^K if $r_X^k(x_{0:k}) \leq r_X^{k-1}(x_{0:k-1})$
 - \Rightarrow Order **decreasing**
- ▶ Dissimilarity network D_X^K if $r_X^k(x_{0:k}) \geq r_X^{k-1}(x_{0:k-1})$
 - \Rightarrow Order **increasing**

- ▶ A **correspondence** C between X and Y is $C \subseteq X \times Y$ s.t.
- ▶ $\forall x \in X$, there exists $y \in Y$ such that $(x, y) \in C$
- ▶ $\forall y \in Y$, there exists $x \in X$ such that $(x, y) \in C$
- ▶ Generalizes permutations
- ▶ $\mathcal{C}(X, Y)$ the set of all correspondences



- ▶ Given proximity networks P_X^K and P_Y^K and a correspondence C
- ▶ Define the k -order network difference with respect to C as

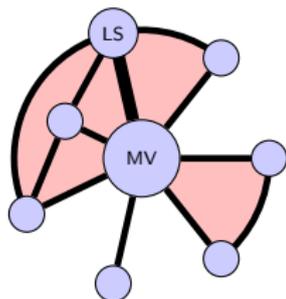
$$\Gamma_{X,Y}^k(C) := \max_{(x_{0:k}, y_{0:k}) \in C} |r_X^k(x_{0:k}) - r_Y^k(y_{0:k})|.$$

- ▶ The k -order proximity network distance is defined as

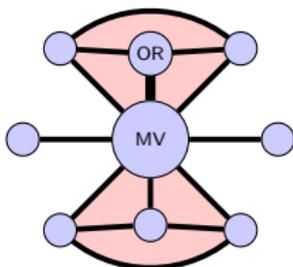
$$d_{\mathcal{P}}^k(P_X^K, P_Y^K) := \min_{C \in \mathcal{C}(X,Y)} \{\Gamma_{X,Y}^k(C)\}.$$

Theorem

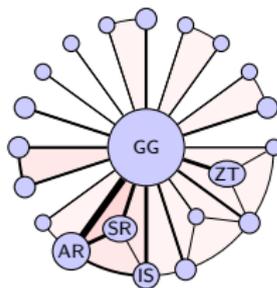
$d_{\mathcal{P}}^k : \mathcal{P}^K \times \mathcal{P}^K \rightarrow \mathbb{R}_+$ is a *metric* in space $\mathcal{P}^K \bmod \cong_k$ for $k \geq 1$ and a *pseudometric* in $\mathcal{P}^K \bmod \cong_0$.



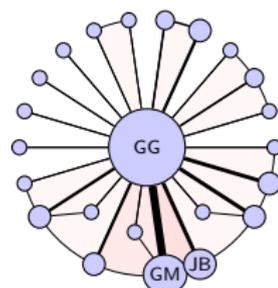
MV, 2004 - 2008



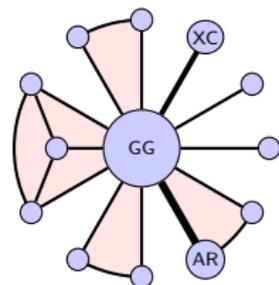
MV, 2009 - 2013



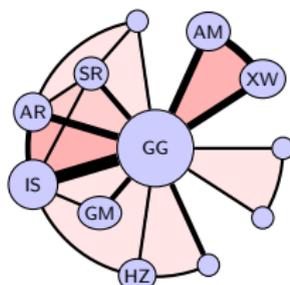
GG, 2004 - 2008



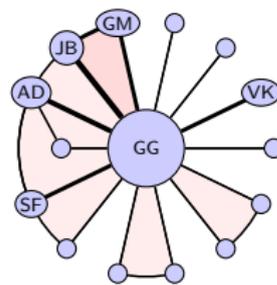
GG, 2009 - 2013



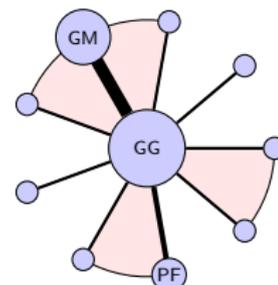
GG, 2006 - 2007



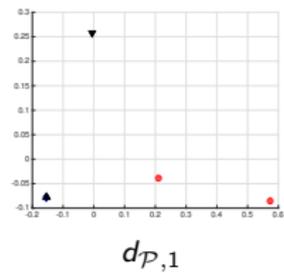
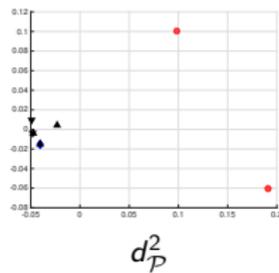
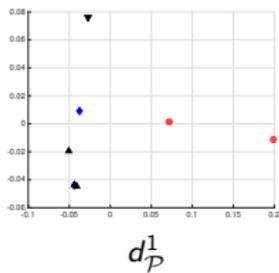
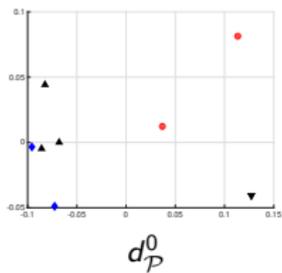
GG, 2008 - 2009



GG, 2010 - 2011

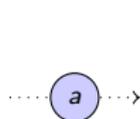


GG, 2012 - 2013



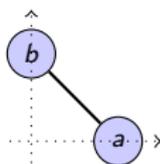
- ▶ Computable for large networks
- ▶ Admittedly, lower bounds may suffer same problems as features
- ▶ Large lower bound entails a large distance
- ▶ **Upper bound** easy to establish
 - ⇒ Used to estimate **distance intervals**

- k -simplex $[x_{0:k}]$ is the convex hull of the set of points $x_{0:k}$



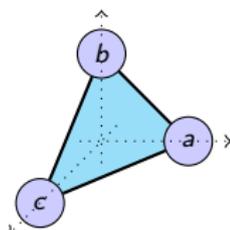
vertex $[a]$

0-simplex



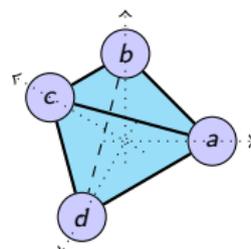
edge $[a, b]$

1-simplex



triangle $[a, b, c]$

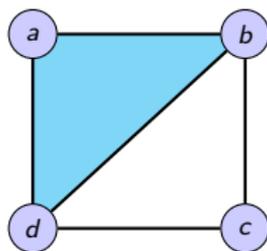
2-simplex



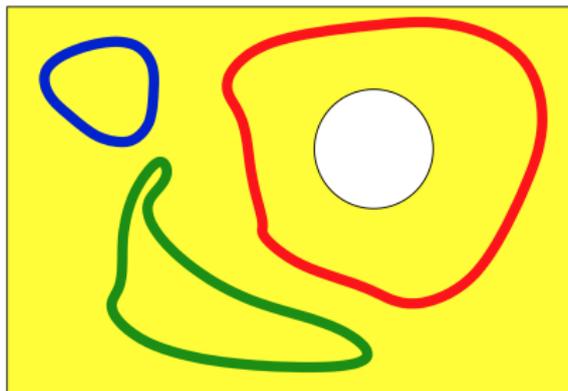
tetrahedron $[a, b, c, d]$

3-simplex

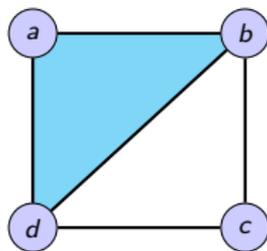
- ▶ **Simplicial complex L** is the collection of simplices glued together



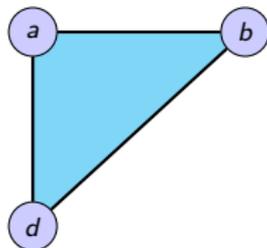
- ▶ We want to describe **holes** that do not have **interiors**
- ▶ Why do we consider them \Rightarrow think of rubber bands
 - \Rightarrow Rubber band enclosing them cannot be diminished



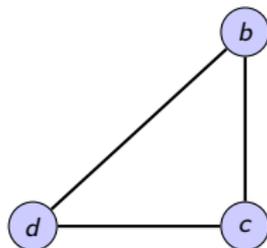
- ▶ We want to describe **holes** that do not have **interiors**
 - ⇒ **Homological features** are defined to formalize this
 - ⇒ **Cycles** without any **interiors**



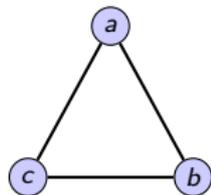
- ▶ **Homological features** describe **hole** with no **interior**
- ▶ $[a, b], [b, d], [d, a]$ forms a **hole** that has **interior**
⇒ Not a **Homological feature**
- ▶ $[d, b], [b, c], [c, d]$ forms a hole with no interior
⇒ Is homological feature



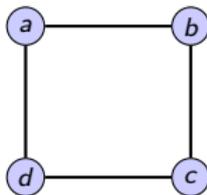
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⇒ Is **homological feature**



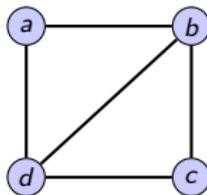
- ▶ **Homological features** describe **hole** with no **interior**



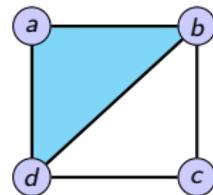
1 feature



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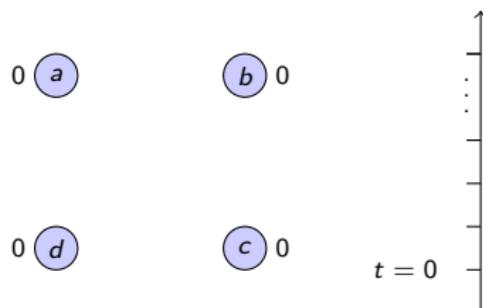


2 features

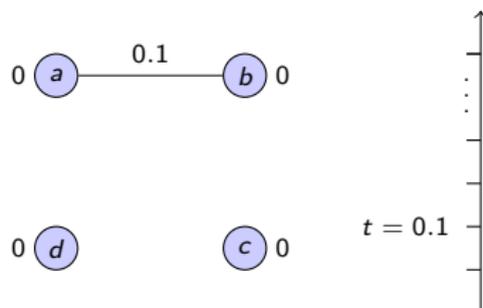


1 feature

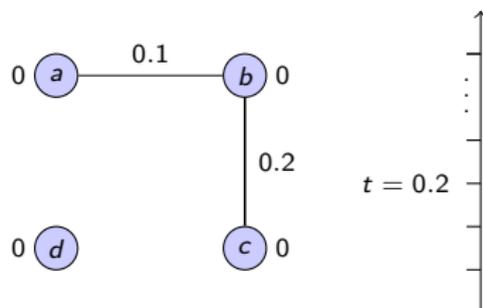
- ▶ Simplicial complexes form **unweighted** networks
 - ▶ No way to incorporate **weights**
 - ▶ To solve this problem, assign each simplex a **value**
 - ▶ The time when this simplex **appears**



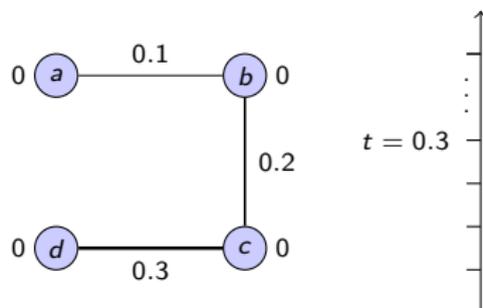
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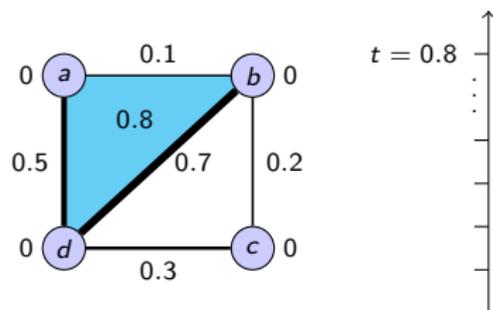
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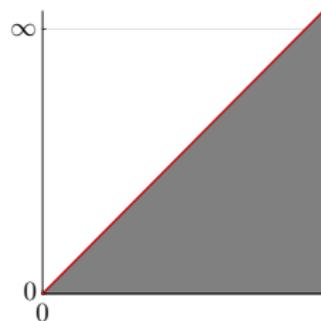
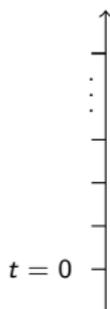
- Quantify when do **holes** and **interiors** appear

0 *a*

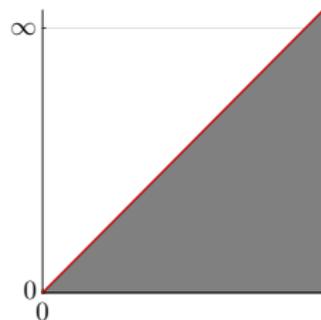
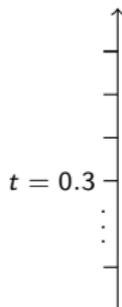
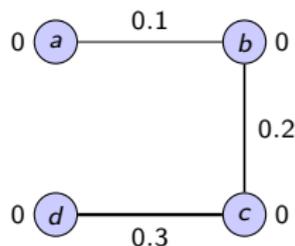
b 0

0 *d*

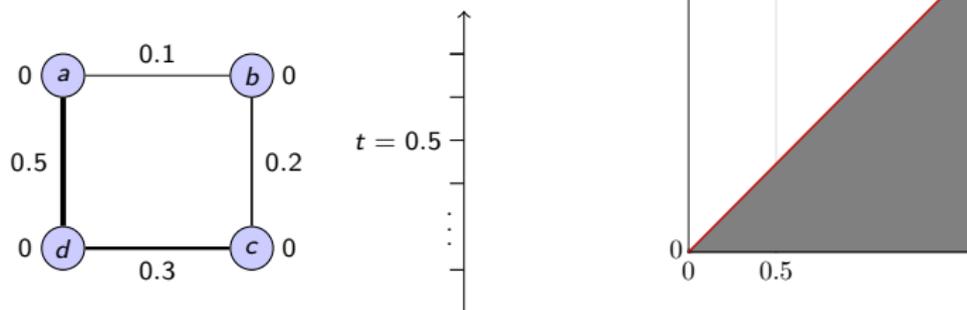
c 0



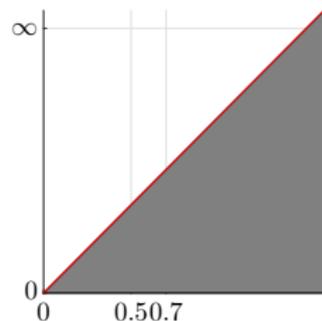
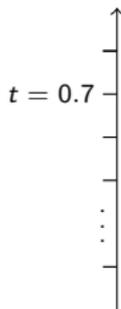
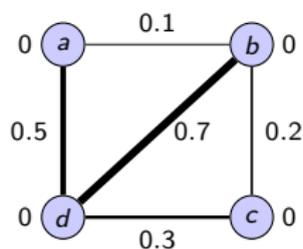
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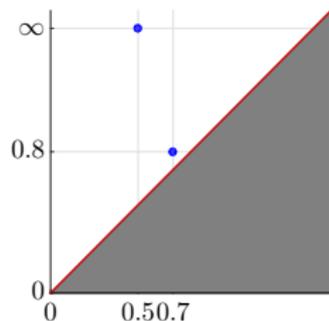
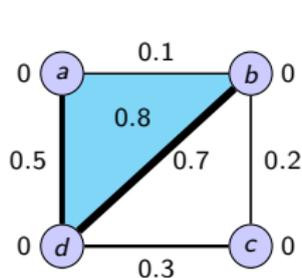
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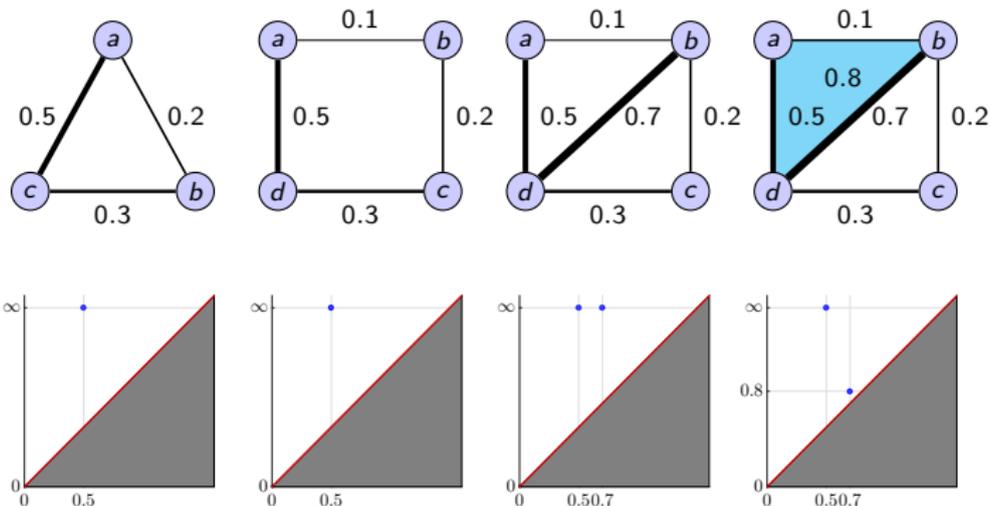
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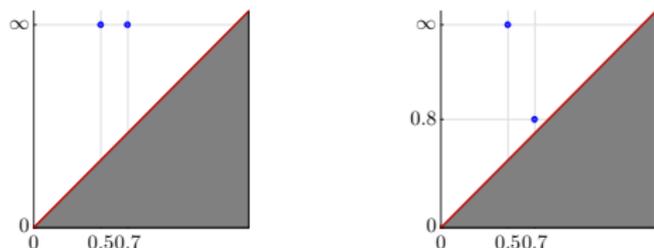
- Quantify when do **holes** and **interiors** appear



- ▶ Bottleneck distance $d_B^\infty(Q, \tilde{Q})$ between two point sets Q and \tilde{Q}

$$d_B^\infty(Q, \tilde{Q}) = \min_{\pi} \max_{q \in Q} \|q - \pi(q)\|_\infty,$$

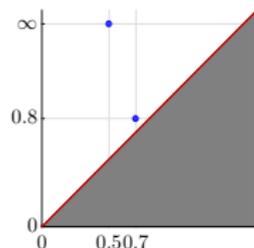
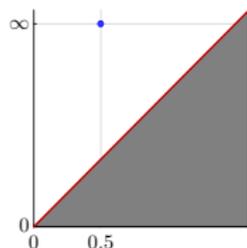
- ▶ where π ranges over all bijections from Q to \tilde{Q}
- ▶ $|Q| = |\tilde{Q}|$ are point sets in two dimensional space



$$d_B^\infty(Q, \tilde{Q}) = \max\{|\infty - 0.8|, |0.7 - 0.7|\} = \infty$$

- ▶ d_B^∞ ill-defined if for diagrams with different cardinalities
- ▶ Homological features trivialized at the same time they appear
- ▶ Add diagonal points to the persistence diagram with fewer nodes
- ▶ Linear Bottleneck Assignment Problem : $\min_{\pi} \max_i c(q_i, \tilde{q}_{\pi(i)})$

$$c(q, \tilde{q}) = \min \left\{ \|q - \tilde{q}\|_\infty, \frac{1}{2} \max \{|q_x - \tilde{q}_x|, |q_y - \tilde{q}_y|\} \right\}.$$



$$d_B^\infty(Q, \tilde{Q}) = \frac{1}{2} |0.8 - 0.7| = 0.05$$

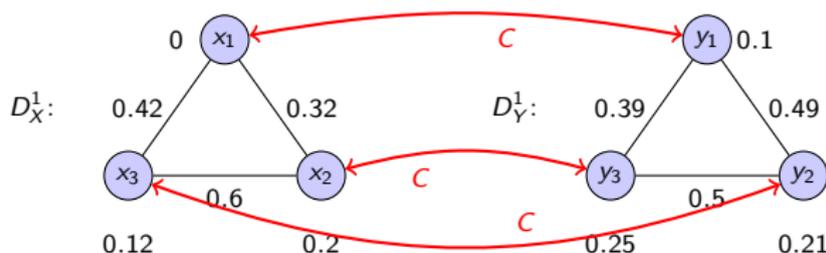
Theorem

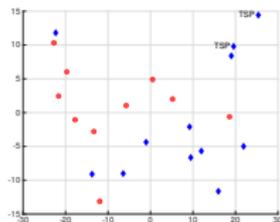
d_B^∞ between the k -persistence diagrams of the filtrations $\mathcal{L}(D_X^K)$ and $\mathcal{L}(D_Y^K)$ is at most $d_{\mathcal{D},\infty}(D_X^K, D_Y^K)$ for any $0 \leq k \leq K$, i.e.

$$d_B^\infty(\mathcal{P}_k \mathcal{L}(D_X^K), \mathcal{P}_k \mathcal{L}(D_Y^K)) \leq d_{\mathcal{D},\infty}(D_X^K, D_Y^K).$$

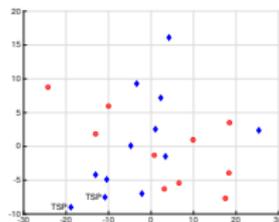
Theorem

Any k -order relationships between full rank tuples of D_X^k appear either in the death time of the $(k - 1)$ -th dimensional homological features or the birth time of the k -th dimensional homological features.

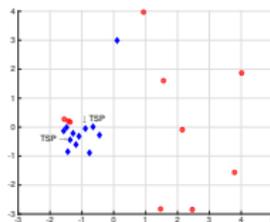




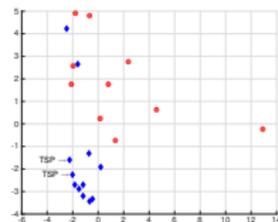
$d_B^\infty(\mathcal{P}_0\mathcal{L})$ removed



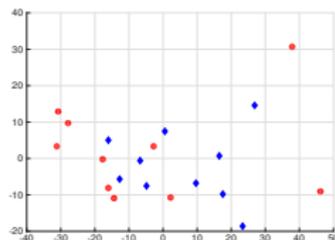
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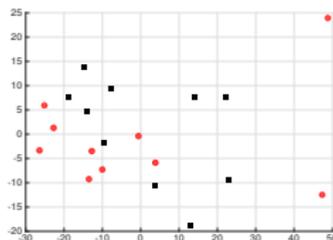
$d_B^\infty(\mathcal{P}_1\mathcal{L})$ replaced



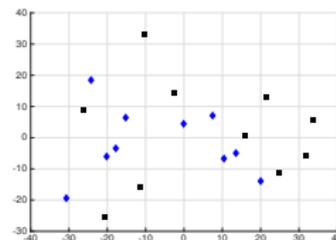
$d_B^\infty(\mathcal{P}_2\mathcal{L})$ replaced



TAC - TSP



TAC - TWC



TSP - TWC

- ▶ Applications
 - ▶ Brain networks at finer scale
 - ▶ Pattern recognition from time series of observations
- ▶ Theory
 - ▶ Clustering based on distance intervals
 - ▶ Graph structure inference