

# Novel Homology-Based Centrality Measures for Weighted Graphs

A Presentation for GTDAML 2021

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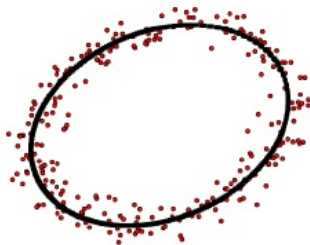
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# Topological Data Analysis

- Properties from the “shape” of data sets. These data sets are usually embedded in metric spaces.

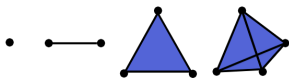
Algebraic Topology and Simplicial Homology

- Focus of our study will be holes.

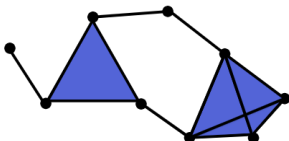


# Vietoris-Rips Simplicial Complex

- Connect, using an edge, any two points  $\epsilon$  units apart.
- A higher-dimensional simplex forms assuming its codimension-1 faces appear.



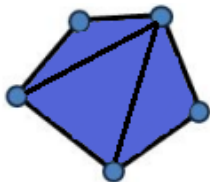
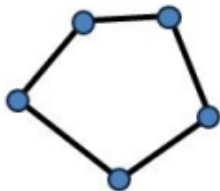
- The resulting structure is a **simplicial complex**, a generalization of graphs.



- Increasing the parameter  $\epsilon$ , we obtain a sequence of simplicial complexes known as a **filtration**.

# Persistence of Homology Classes

- Holes and voids, as equivalence classes, are called **homology classes**.



- For any homology class, we collect the ordered pairs  $(b, d)$  where  $b$  is the **birth** and  $d$  is the **death** of the homology class. The difference  $d - b$  is called the **persistence**.

# Quantifying Importance of a Homology Class

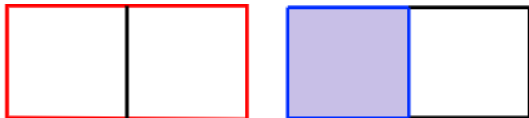
- More persistent homology classes are more important.
- Short-lived homology classes are important in estimation problems such as determining curvatures of surfaces [2].
- (Graph Theory) Loop Centrality is an importance based on the number of walks passing a cycle or loop.

# Merging Homology Classes

- A homology class either trivializes or merges with another homology class.
- The elder rule states that an elder feature in the filtration survives while the other feature dies when two features merge.
- Two  $k$ th homology classes  $[\sigma]$  and  $[\delta]$  **merge** at  $\epsilon$  if

$$\sigma + \delta = \rho$$

for some  $k$ th boundary  $\rho$ .



# Proposed Centrality Measures

- The form of the proposed centrality measures of homology class  $[\sigma]$  is given by

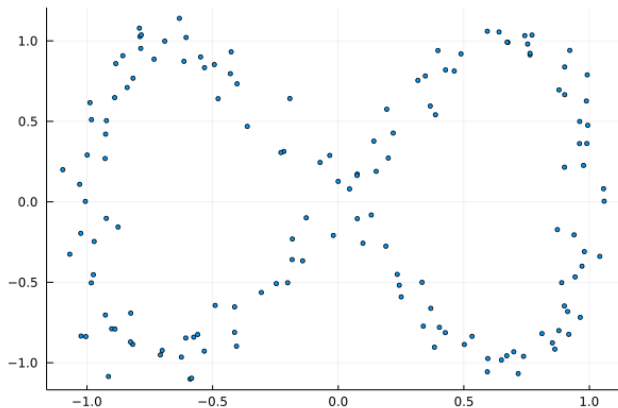
$$J(\sigma, \epsilon) = \begin{cases} 0 & \text{for } \epsilon \leq b(\sigma) \\ [\epsilon - b(\sigma)] + \sum_{\varsigma \in M_1[\sigma, \epsilon]} cP(\varsigma) & \text{for } b(\sigma) < \epsilon \leq d(\sigma) \\ P(\sigma) + \sum_{\varsigma \in M_1[\sigma, \epsilon]} cP(\varsigma) & \text{for } \epsilon > d(\sigma) \end{cases}$$

where  $c \in (0, 1]$ .

- Let  $c = 1$  to denote full weight on the persistence of merging homology classes.
- Consider the merging time and death to give preference to early and late merging.
- Extend those three centrality measures by also considering merging classes of each homology class in the merging class.

# Numerical Simulations on Synthetic Data Sets

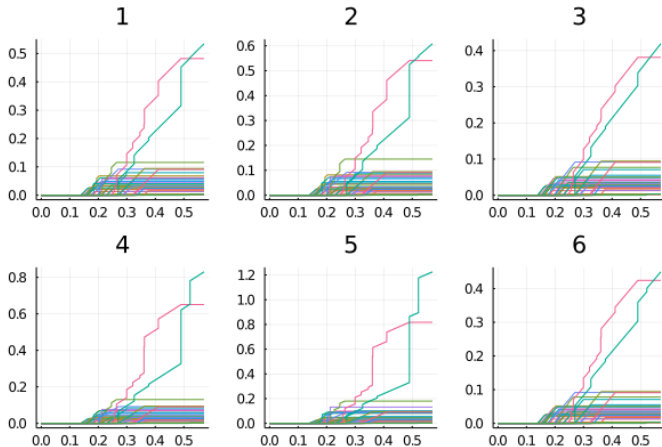
Consider the point cloud depicting a noisy infinity sign (or figure eight).



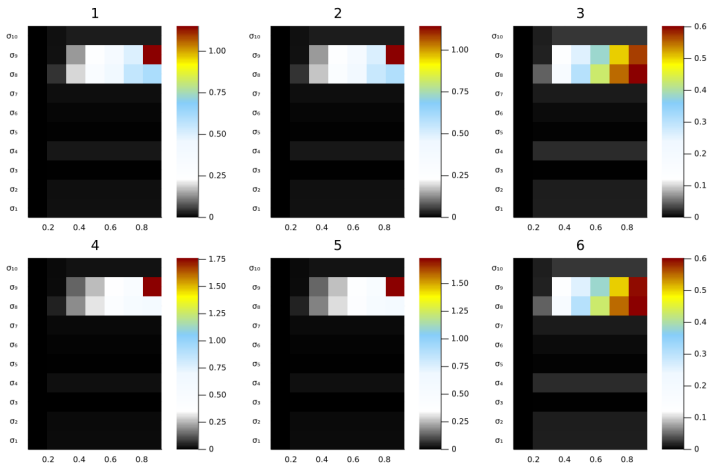


# Representation of Centrality Measures

As expected, we see two relatively important holes. Those two holes are the largest holes on the left and right symmetrical parts of the infinity sign.

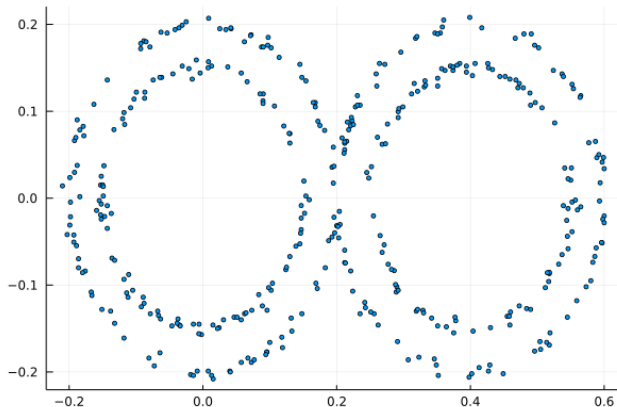


# Alternative Representation



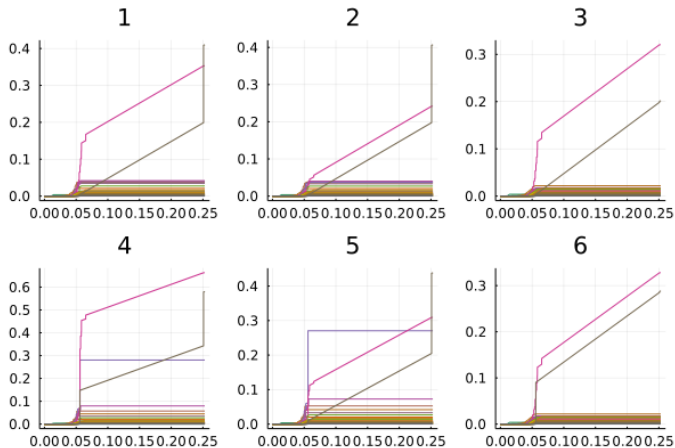
# Another Numerical Simulation

Now, we consider a noisy infinity sign but added with an annular region. Two large holes are visible and expected to be captured by the centrality plot.

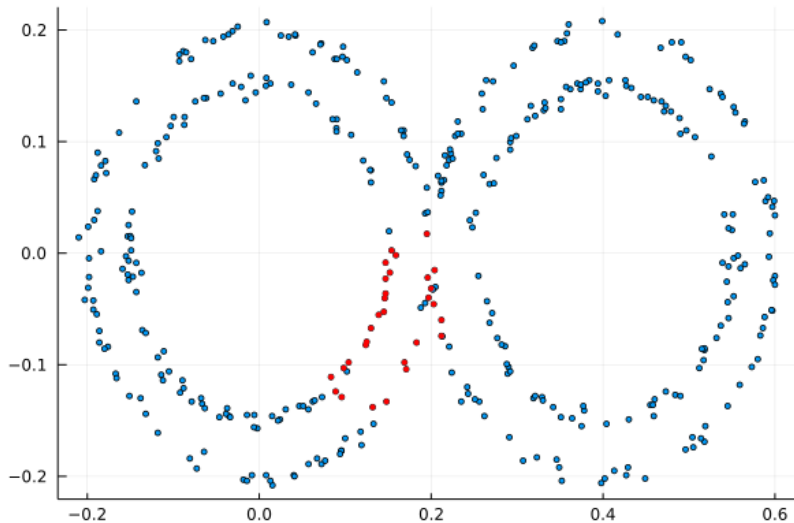


# Another Interpretation

The extended centrality measures capture a third relatively important hole. Centrality 5 gives preference to early merging which means the third important hole merges early with other holes.



# Third Relatively Important Homology Class



# Differences to Common Topological Summaries

- A barcode or a persistence diagram does not capture the third hole as relatively important.
- Less persistent holes may be important different to popular notion of importance for homology classes.

# Properties of the Centrality Measures

- The centrality measure is a monotonic function.
- For each homology class  $[\sigma]$  in the filtered simplicial complex, we define

$$\|J_\sigma(\epsilon)\|_p = \left\{ \int_{[0, d(\sigma)]} |J_\sigma(x)|^p dx \right\}^p$$

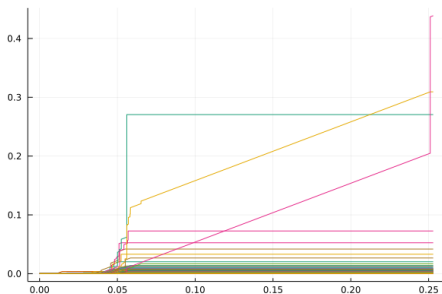
for  $1 \leq p < \infty$ . Furthermore, we form the set

$$\{\|J_\sigma(\epsilon)\|_p : \sigma \text{ is a homology class}\}.$$

We want to define a distance between two centrality plots.

# Properties (Continuation)

The landscape distance [1] compares the  $k$ th order landscapes. The centrality plot cannot have such reasonable ordering. Hence, we consider all possible matchings between the centrality curves similar to the idea of the bottleneck distance.





# Properties (Continuation)

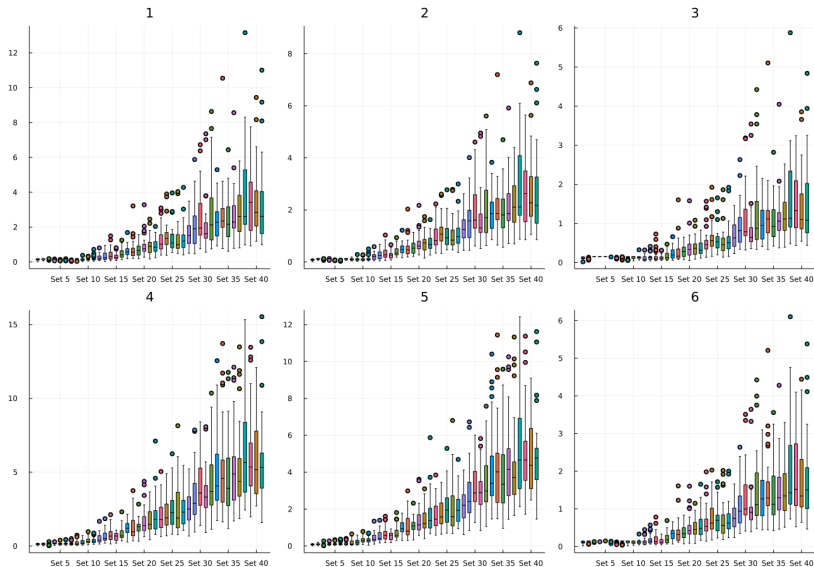
Assuming the birth times are equal, the implementation [4] of the bottleneck distance between two persistence diagrams depending only on the death times,  $\delta_x$  and  $\delta_{\phi(x)}$ , is given by  $\inf_{\phi} \sup_{x \in X} \|x - \phi(x)\|_{\infty}$  where





$$\|x - \phi(x)\|_{\infty} = \begin{cases} \frac{1}{2} \max\{\delta_x, \delta_{\phi(x)}\} & \text{if } \phi(x) \in \Delta \\ |\delta_x - \delta_{\phi(x)}| & \text{otherwise} \end{cases} .$$

The image  $\phi(x)$  of  $x$  under a bijection  $\phi$  is a matched point for  $x$  and  $\Delta$  is the set of diagonal points. We replace the death times to  $|J_{\sigma}(\epsilon)|_{\rho}$  which is a number that only depends on  $d(\sigma)$ .

- (Current Work) Stability of centrality plots under the defined distance.

# Empirical Evidence



-  P. BUBENIK AND P. DŁOTKO, *A persistence landscapes toolbox for topological statistics*, Journal of Symbolic Computation, 78 (2017), pp. 91 – 114.
-  P. BUBENIK, M. HULL, D. PATEL, AND B. WHITTLE, *Persistence homology detects curvature*, Inverse Problems, 36 (2020).
-  G. E. CARLSSON, *Topology and data*, Bulletin (New Series) of the American Mathematical Society, 46 (2009), pp. 255–308.
-  P. S. IGNACIO, J.-A. BULAUAN, AND D. UMINSKY, *Lumáwig: An efficient algorithm for dimension zero bottleneck distance computation in topological data analysis*, Algorithms, 13 (2020).

Thank you for listening! Any  
Questions?