# An Introduction to Topological Data Analysis

Matthew Zabka

Laber Labs

North Carolina State University

Southwest Minnesota State University

#### Outline

Crash course in algebraic topology

Persistent Homology and examples

JPDwB Tutorial

Persistence Homology and RIPSER

Multiparameter Persistence and RIVET

### **Algebraic Topology**

### What is algebraic topology?

Topology is a branch of mathematics which is good at extracting global qualitative features from complicated geometric structures.

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Topological questions surround different notions of connectedness: connected components, loops, voids, etc.

Two topological spaces are equivalent through the lens of topology if one can be *continuously* deformed to the other.



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### Invariants of topological spaces

- Algebraic Topology assigns invariants to topological spaces. These take the form of groups, rings, fields, vector spaces, etc.
- Our computations will be over the field F<sub>2</sub>, so it suffices to record only the dimensions of vector spaces.
- If two spaces are the same, then the invariants must be the same.

  If the invariants are not the same, then the two spaces are not the same.

# **Topological Data Analysis**

#### Goal

Topological data analysis uses topology to summarize and study the 'shape' of data.

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Topological data analysis uses topology to summarize and study the 'shape' of data.

#### Examples:

- motion tracking problems,
- analysis of brain arteries,
- analysis of social and spatial networks, including neuronal networks, Twitter, co-authorship,
- study of viral evolution,
- measurement of protein compressibility, steganalysis of images,
- analysis of phase transitions,
- financial crash analysis,
- piecewise constant signal analysis,

- study of cosmic web and its filamentary structure.
- identification of breast cancer subtypes,
- study of plant root systems,
- discrimination of EEG signals before and during epileptic seizures,
- sphere packings,
- population activity in the visual cortex,
- fMRI data

#### Overview of TDA

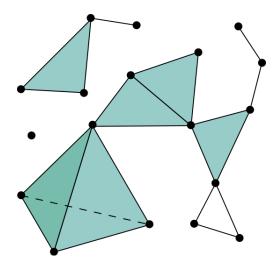
Topological data analysis comes in a variety of flavors.

The two most popular methods in TDA are

- 1. Persistent Homology
  - 2. Mapper

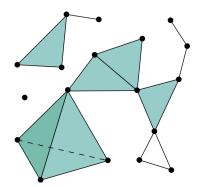
### **Simplicial Complexes**

A *simplicial complex* is a combinatorial object, generalizing the notion of a graph. Each simplicial complex is built out of *simplices* of varying dimensions.



### Betti numbers of simplicial complexes

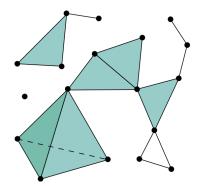
$$eta_0 = \#$$
 of connected components  $eta_1 = \#$  of holes  $eta_2 = \#$  of voids



$$\beta_0 = \beta_1 = \beta_2 = \beta_2$$

### Betti numbers of simplicial complexes

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$$\beta_0 = 3$$

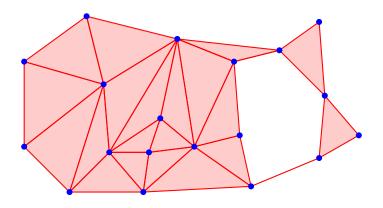
$$\beta_1 = 1$$

$$\beta_2 = 1$$

# Homology of simplicial complexes

### **Definition**

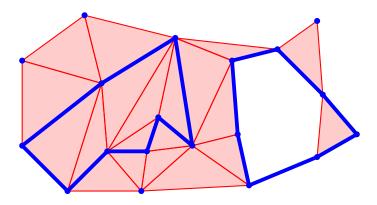
Homology in degree k is given by k-cycles modulo the k-boundaries.



## Homology of simplicial complexes

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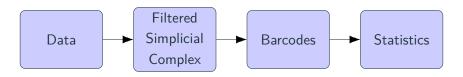
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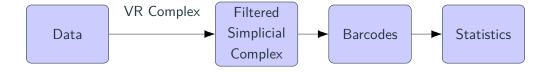
 $\beta_k = \text{rank of homology in degree } k$ 

#### Overview of PH

Persistent homology consists of the following pipeline:



### Overview of PH



# Simplicial Complexes from Point data

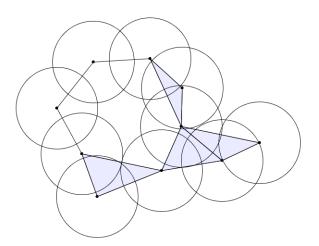
#### **Definition**

A point cloud P is a finite metric space.

### Simplicial Complexes from Point data

#### **Definition**

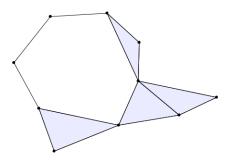
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# Vietoris-Rips parameter

### Question

How do we choose the correct radius for the Vietoris-Rips construction?

Often, there is no one "right" choice.

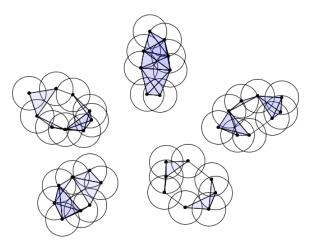
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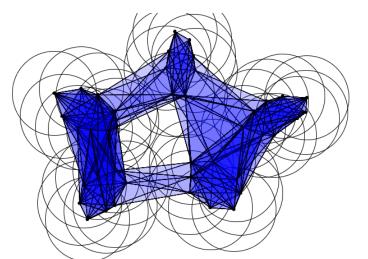


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# **Processing demo**

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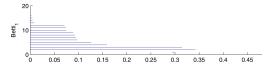
### Overview of PH

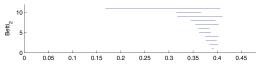


#### **Barcodes**

- The barcode provides a summary of how the homology changes as the radius varies in the Vietoris-Rips construction.
- We look for topological features which 'persist' over many values of radii.

### Barcodes typically look like:





#### **InteractiveJPDwB**

- Recall: Given a set of points X and a parameter r, we can create a formal simplicial complex.
- Let the *n*-simplex  $\{x_0, x_1, \dots, x_n\}$  exist if and only if  $d(x_i, x_j) < r$  for all  $0 \le i, j \le n$ .
- We can visualize this process using InteractiveJPDwB.

#### **InteractiveJPDwB**

- InteractiveJPDwB lets one visualize the generated simplicial complex for data in  $\mathbb{R}^2$  and different values of r.
- In other words, it demonstrates 0-th and 1-st degree persistence homology via barcodes.
- Thanks to Michael Catanzaro, you can easily use this program on your own computer.
- You can download this program from:

https://github.com/MatthewZabka/LaberLabs18

### Anwer the following!

#### https://github.com/MatthewZabka/LaberLabs18

- What is the minimum number of points required so that  $\beta_1 = 1$  for some value of r?
- What is the minimum number of points required so that, for some r, we have  $\beta_0 = 3$  and  $\beta_1 = 2$ ?
- What is the largest degree of homology that is geometrically feasible in  $\mathbb{R}^2$ ?
- What is the minimum number of points required to have  $\beta_2=1$ ? In what dimension must the points lie?
- What is the minimum number of points required to have  $\beta_n = 1$ ? In what dimension must the points lie?

# Discussion

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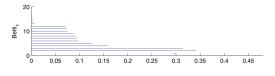
### **Overview of Persistence Homology**

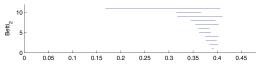


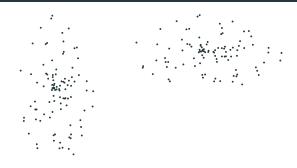
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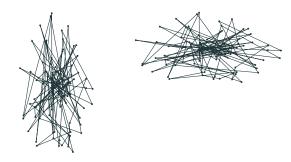


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- If a topological property (like a Betti number) persist over a large range of r, we can conclude something about the structure of the data.



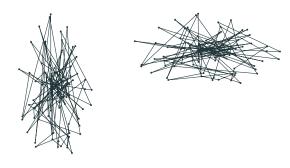
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- If a topological property (like a Betti number) persist over a large range of r, we can conclude something about the structure of the data.
- In this case, we should expect to see  $\beta_0 = 2$  over a large range of r.

# Persistent Homology in Dimension 0 (Clustering)



ullet A graph similar to this should persist over a large range of r.

# Persistent Homology in Dimension 0 (Clustering)



- ullet A graph similar to this should persist over a large range of r.
- How could we see the clusters if these data did not lie in  $\mathbb{R}^2$ ?

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- Developed by Ulrich Bauer, RIPSER is a very fast C++ program for computing Vietoris-Rips persistence barcodes.
- Let's try this on our data with two clusters.



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 $\verb|https://github.com/MatthewZabka/LaberLabs18||$ 



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• Input cloud1.txt into RIPSER:

https://live.ripser.org/

• Suppose we have data that lie on the circle.



**Figure 1:** An unrealistic example cloud2.txt, where data lie perfectly on  $S^1$ .

• Suppose we have data that lie on the circle.



Figure 1: An unrealistic example cloud2.txt, where data lie perfectly on  $S^1$ .

• Input cloud2.txt into RIPSER.

• Suppose we have data that almost lie on the circle.

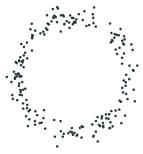


Figure 2: A slightly more realistic example cloud3.txt.

• Suppose we have data that **almost** lie on the circle.



Figure 2: A slightly more realistic example cloud3.txt.

• We want to analyze data we cannot see!

### https://github.com/MatthewZabka/LaberLabs18

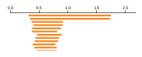
- Now it is your turn! Data are located in the RIPSERdata folder that you have already downloaded!
- Input cloud1.txt int RIPSER. Confirm that two generators of  $H_0$  persist. (i.e.  $\beta_0=2$ )
- Input cloud2.txt into RIPSER. Confirm that one generator for  $H_0$  and one generator for  $H_1$  persist. (i.e.  $\beta_0=1$  and  $\beta_1=1$ )
- Input cloud3.txt into RIPSER. Confirm that one generator for  $H_0$  and one generator for  $H_1$  persist. (i.e.  $\beta_0=1$  and  $\beta_1=1$ )
- Try inputting cloud4.txt into RIPSER up to distance 2. What can you say about the data's shape?
- Try some actual data! cloud5.txt (Test up to distance 150.)
- More actual data! cloud6.txt (Test up to distance 2.)

# Discussion

- What is the shape of cloud4.txt?
- What is the shape of cloud5.txt?
- What is the shape of cloud6.txt?

# **Combining Persistence and Other Methods**

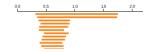
• Suppose we had a barcode in dimension 1 that looked as follows:



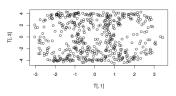
• What are are possibilities for the manifold on which the data lie?

# **Combining Persistence and Other Methods**

• Suppose we had a barcode in dimension 1 that looked as follows:



• Suppose we perform PCA and get the following projection:



- What does PCA suggest about the dimension of the manifold?
- What does this suggest about the space on which the data lie?

• Let's think about how this could go wrong and how we could fix it!

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- Suppose that, instead of data that look like this:

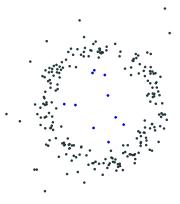


- Let's think about how this could go wrong and how we could fix it!
- We had data that looked like this:

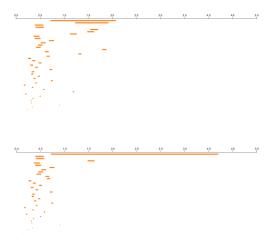


• This is a much more realistic example.

- This is a much more realistic example.
- ullet The blue points noise will make it hard to see the generator of  $H_1$ .



The first barcode includes the entire data set. The second barcode eliminates the blue 'noise'.



• One way to deal with such situations: consider the density of the points.

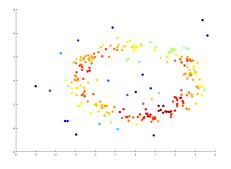


Figure 3: A density heat map of the data

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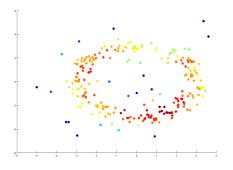


Figure 3: A density heat map of the data

• Letting the density threshold vary results in **multi-parameter persistence**!

- Michael Lesnick and Matthew Wright have written a program for visualizing multiparameter persistence.
- Installation is not so simple.
- Let us try to use RIVET on this example together.



#### References

#### General Overviews:

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- Robert Ghrist. "Homological algebra and data". (2017). URL: https://www.math.upenn.edu/~ghrist/preprints/HAD.pdf.
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#### References

#### Software

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- Michael Lesnick and Matthew Wright. "Interactive visualization of 2-d persistence modules". arXiv preprint arXiv:1512.00180 (2015). http://rivet.online/.
- Luke Wolcott. "InteractiveJPDwB: an interative program for persistence homology". https://github.com/lukewolcott/InteractiveJPDwB. 2016.