

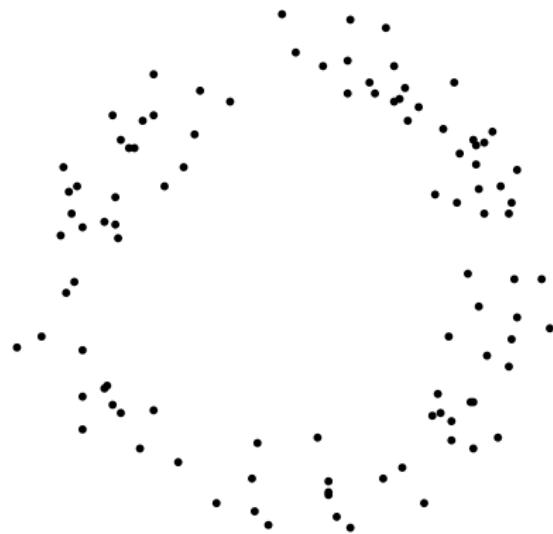
# Persistent connections

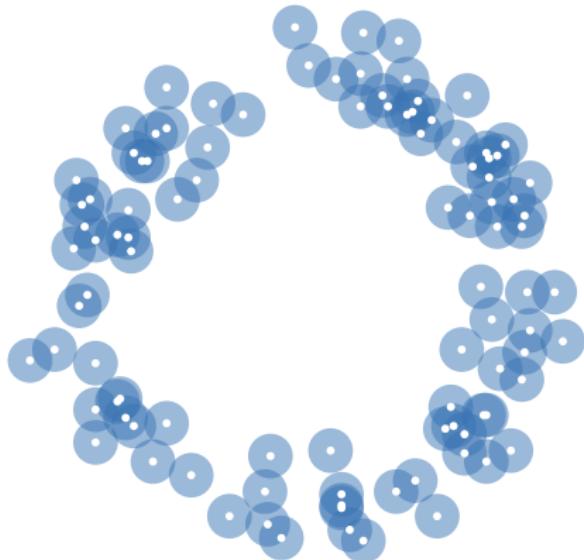
Ulrich Bauer

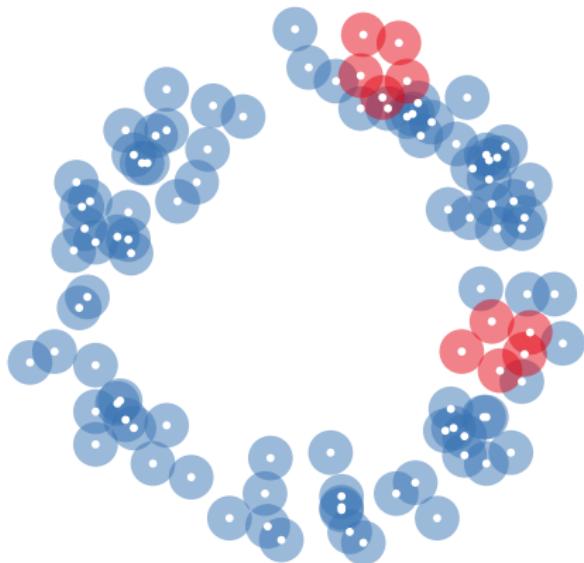
TUM

January 9, 2017

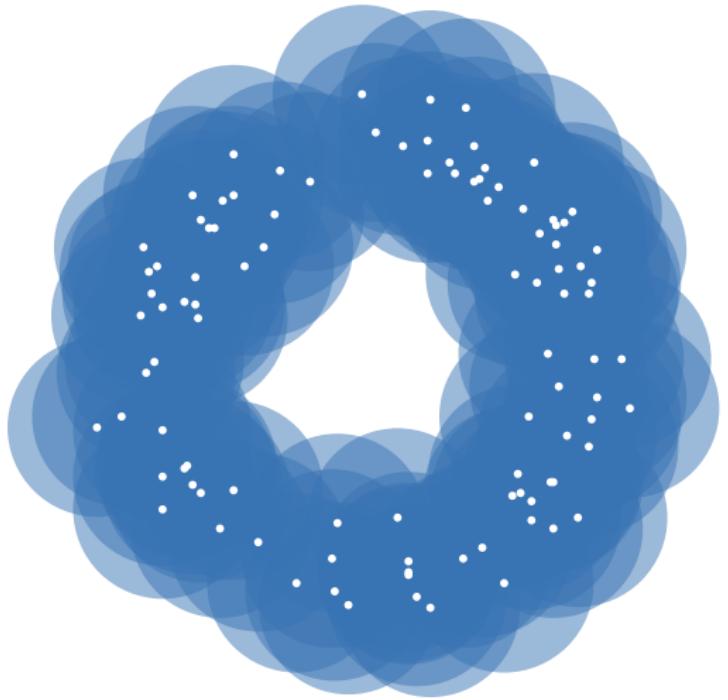
# Persistent homology

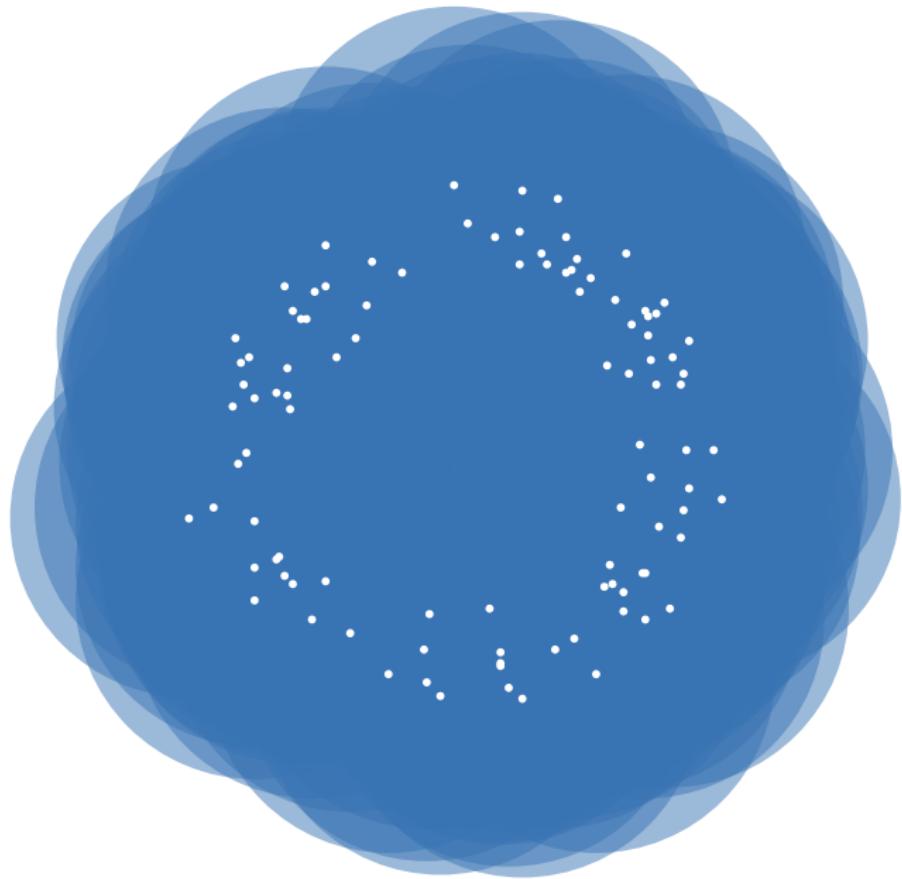


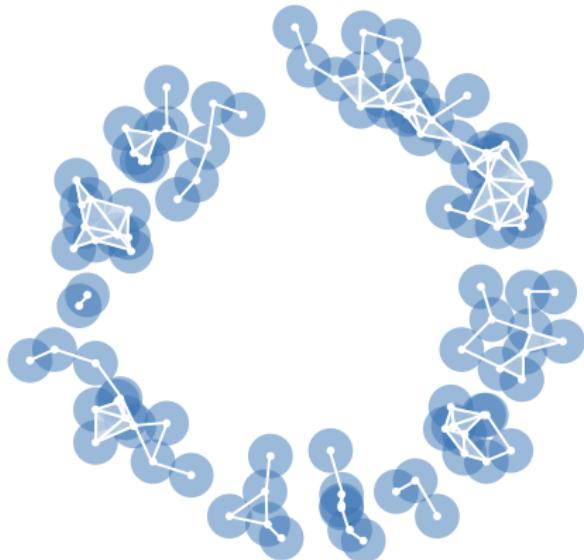


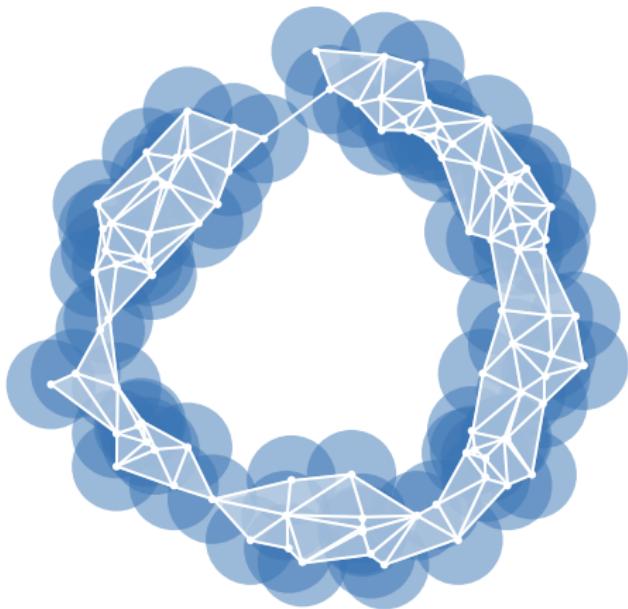


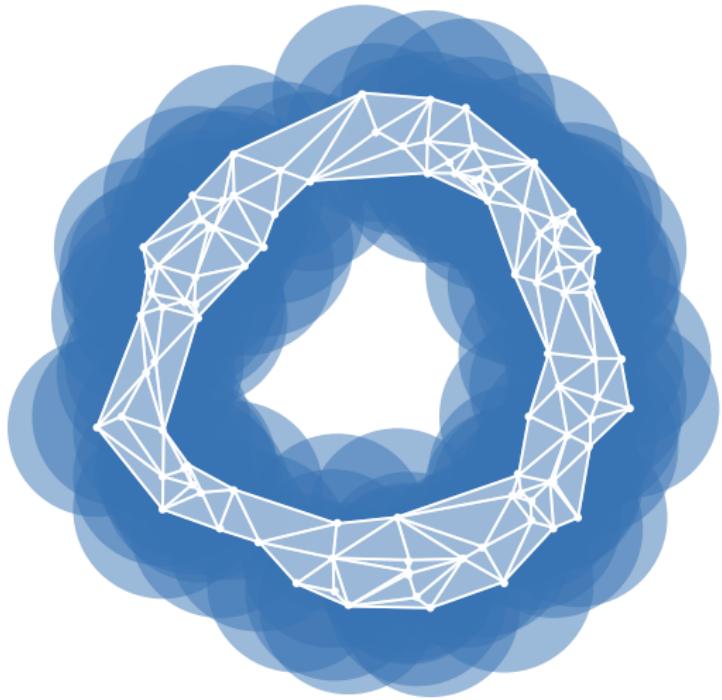


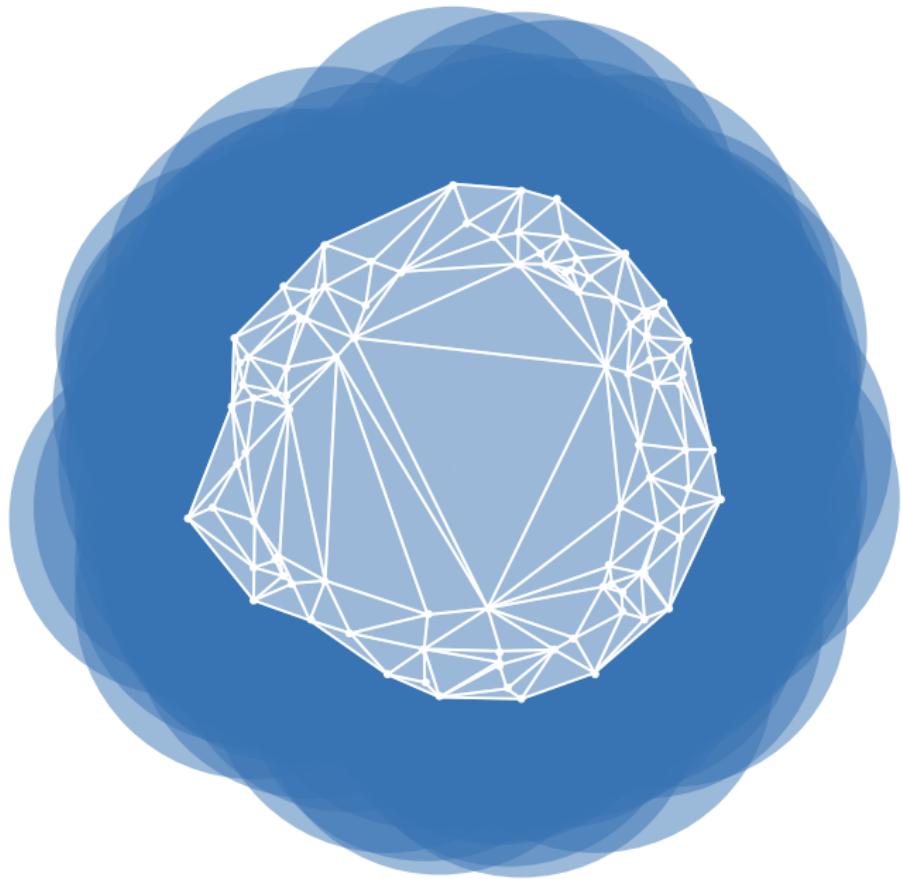




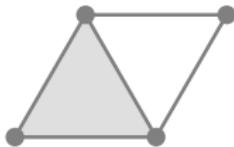






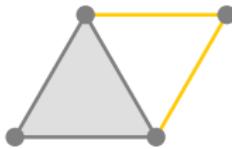


# What is homology?



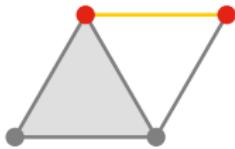
- *Simplicial complex  $K$ :* space built from *simplices* (points, edges, triangles, tetrahedra, ...)

# What is homology?



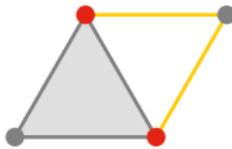
- *Simplicial complex  $K$ :* space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$ :* formal  $\mathbb{Z}_2$  linear combinations of simplices

# What is homology?



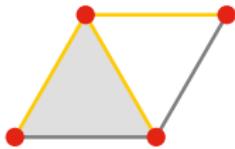
- *Simplicial complex  $K$ :* space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$ :* formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary

# What is homology?



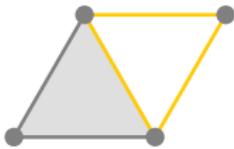
- *Simplicial complex  $K$ :* space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$ :* formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$

# What is homology?



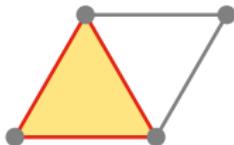
- *Simplicial complex  $K$ :* space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$ :* formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$

# What is homology?



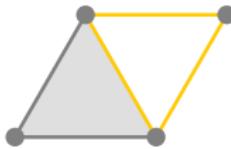
- *Simplicial complex*  $K$ : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains*  $C_*(K)$ : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles*  $Z_*(K) = \ker \partial_*$ : chains with zero boundary

# What is homology?



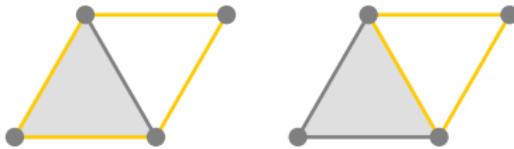
- *Simplicial complex*  $K$ : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains*  $C_*(K)$ : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles*  $Z_*(K) = \ker \partial_*$ : chains with zero boundary
- *Boundaries*  $B_*(K) = \text{im } \partial_*$  are always cycles

# What is homology?



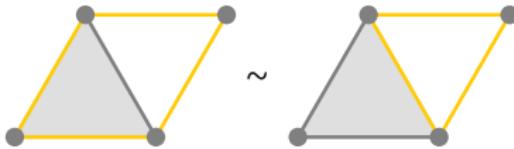
- *Simplicial complex  $K$* : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$* : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles  $Z_*(K) = \ker \partial_*$* : chains with zero boundary
- *Boundaries  $B_*(K) = \text{im } \partial_*$*  are always cycles
- Two cycles are *homologous* if they differ by a boundary

# What is homology?



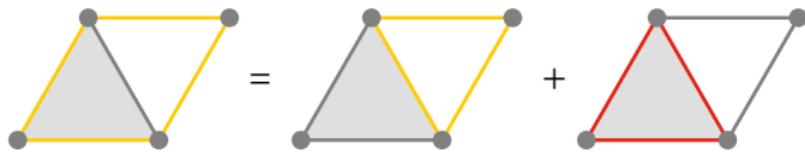
- *Simplicial complex  $K$* : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$* : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles  $Z_*(K) = \ker \partial_*$* : chains with zero boundary
- *Boundaries  $B_*(K) = \text{im } \partial_*$*  are always cycles
- Two cycles are *homologous* if they differ by a boundary

# What is homology?



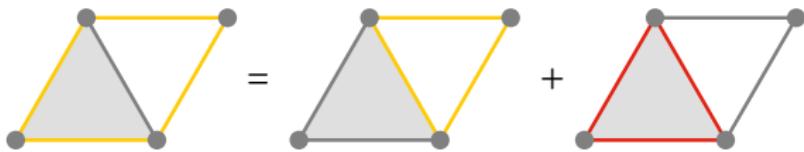
- *Simplicial complex  $K$* : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$* : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles  $Z_*(K) = \ker \partial_*$* : chains with zero boundary
- *Boundaries  $B_*(K) = \text{im } \partial_*$*  are always cycles
- Two cycles are *homologous* if they differ by a boundary

# What is homology?



- *Simplicial complex  $K$* : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$* : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles  $Z_*(K) = \ker \partial_*$* : chains with zero boundary
- *Boundaries  $B_*(K) = \text{im } \partial_*$*  are always cycles
- Two cycles are *homologous* if they differ by a boundary

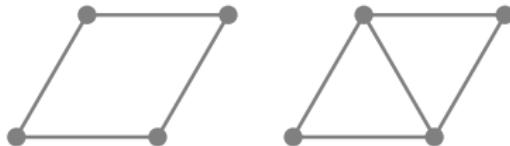
# What is homology?



- *Simplicial complex  $K$* : space built from *simplices* (points, edges, triangles, tetrahedra, ...)
- *Chains  $C_*(K)$* : formal  $\mathbb{Z}_2$  linear combinations of simplices
- Each simplex has a boundary
- Defines linear *boundary map*  $\partial_* : C_*(K) \rightarrow C_*(K)$
- *Cycles  $Z_*(K) = \ker \partial_*$* : chains with zero boundary
- *Boundaries  $B_*(K) = \text{im } \partial_*$*  are always cycles
- Two cycles are *homologous* if they differ by a boundary
- The equivalence classes form the *homology group*  
$$H_*(K) = Z_*(K)/B_*(K)$$

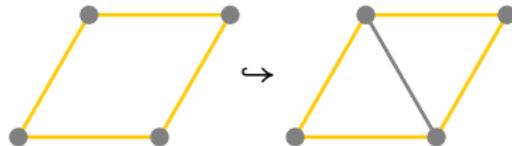
# What is homology?

Roughly:  $\dim H_*(K)$  is the number of holes



# What is homology?

Roughly:  $\dim H_*(K)$  is the number of holes

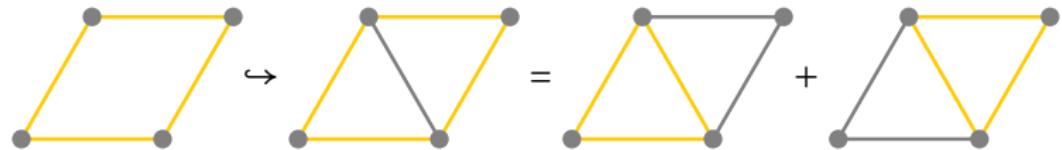


Homology is a *functor*:

- An inclusion map  $L \hookrightarrow K$  induces a linear map  $H_*(L) \rightarrow H_*(K)$

# What is homology?

Roughly:  $\dim H_*(K)$  is the number of holes

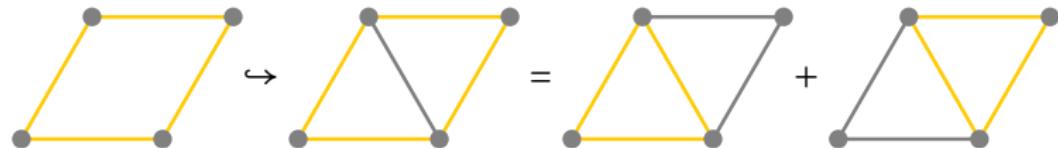


Homology is a *functor*:

- An inclusion map  $L \hookrightarrow K$  induces a linear map  $H_*(L) \rightarrow H_*(K)$
- A “hole” might get mapped to a linear combination of two “holes”

# What is homology?

Roughly:  $\dim H_*(K)$  is the number of holes



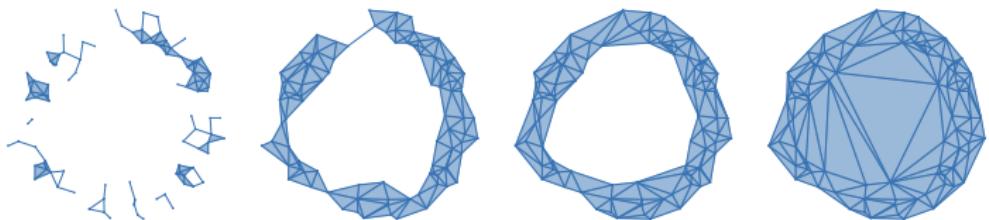
Homology is a *functor*:

- An inclusion map  $L \hookrightarrow K$  induces a linear map  $H_*(L) \rightarrow H_*(K)$
- A “hole” might get mapped to a linear combination of two “holes”

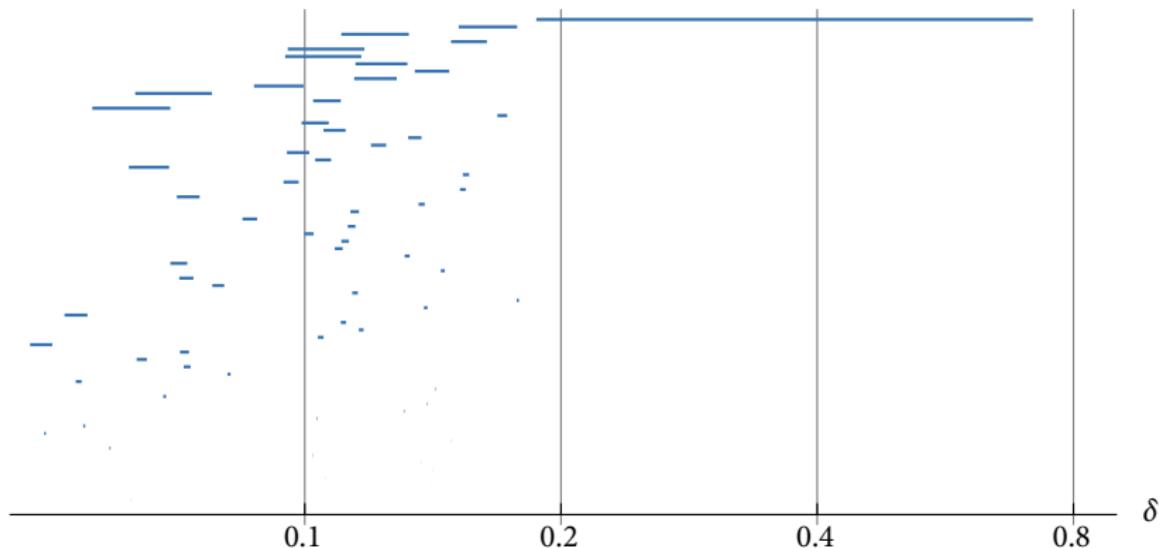
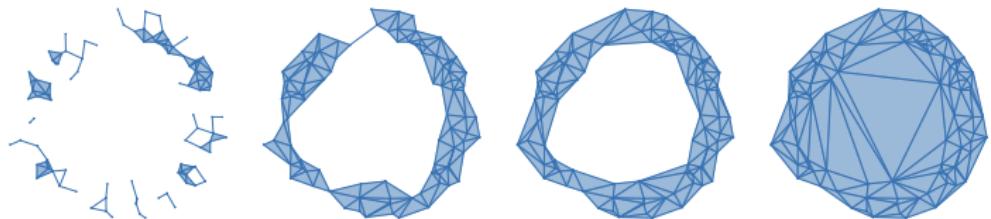
Homology does not have a canonical basis

- Computing (persistent) homology is all about choosing (compatible) bases

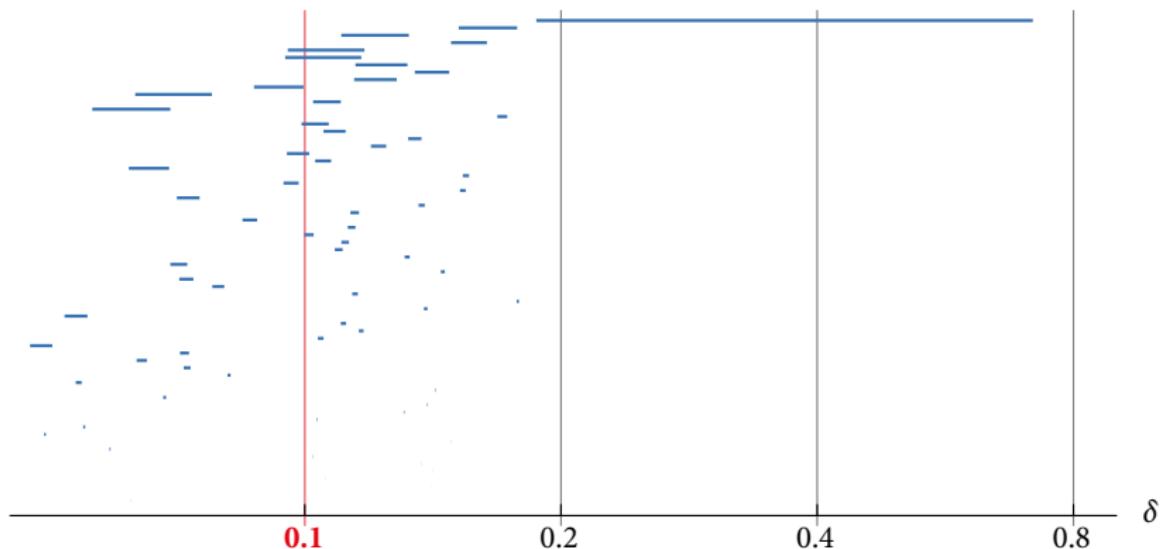
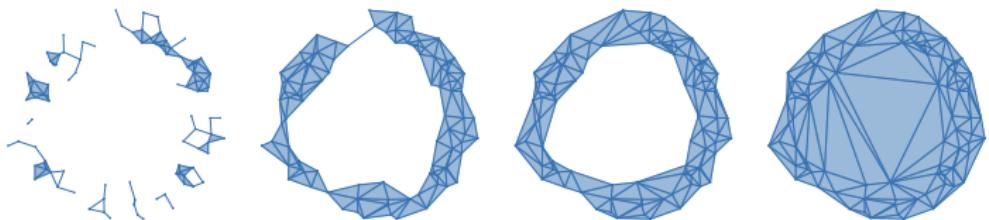
# What is persistent homology?



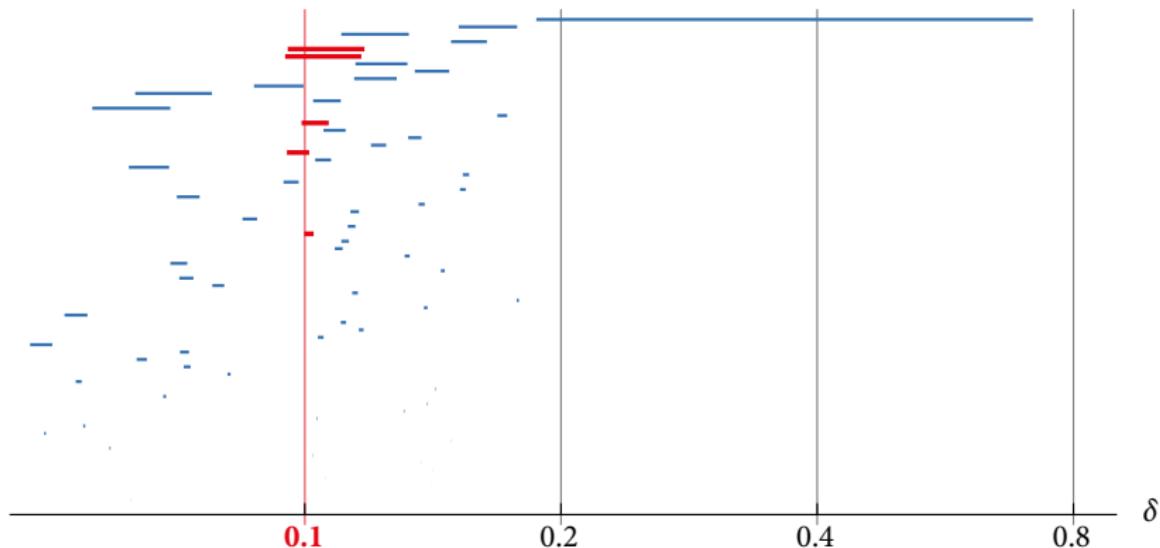
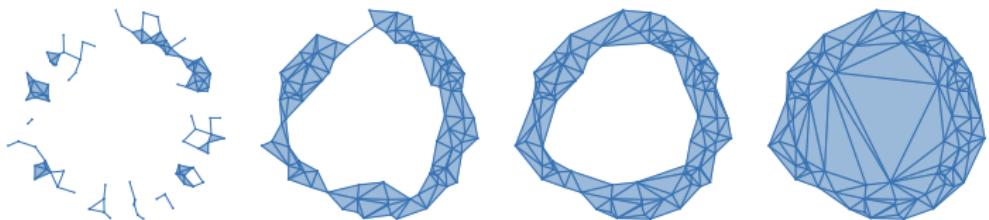
# What is persistent homology?



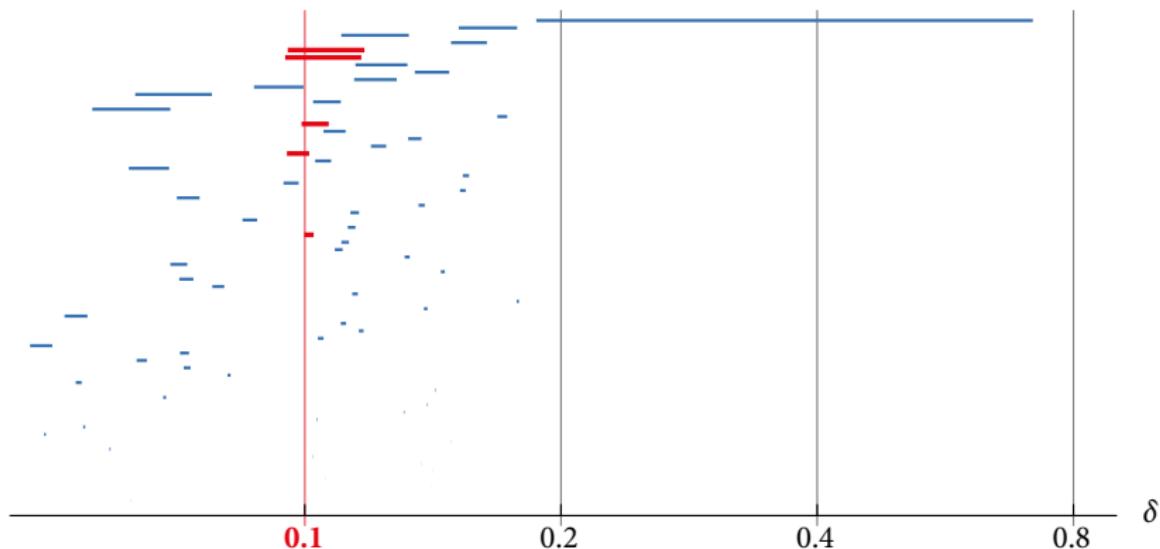
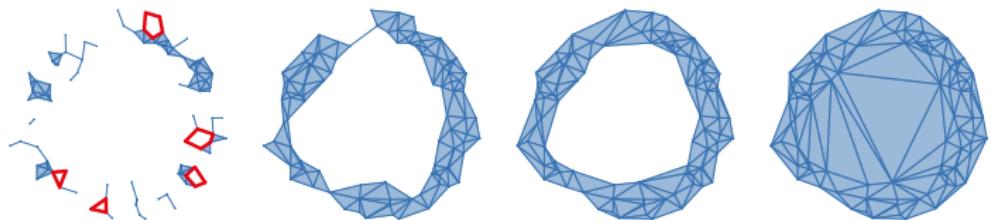
# What is persistent homology?



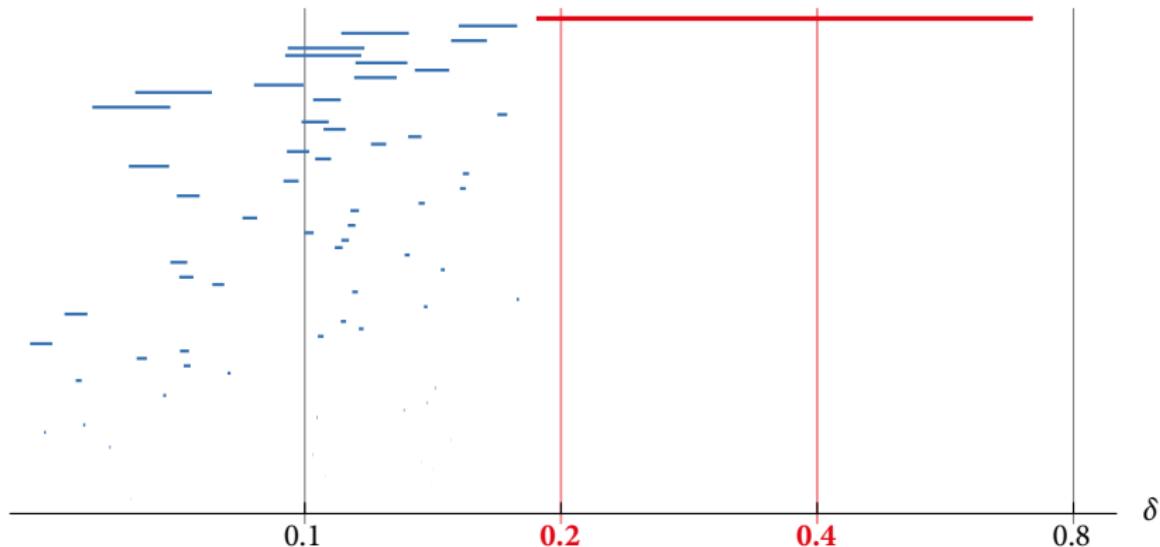
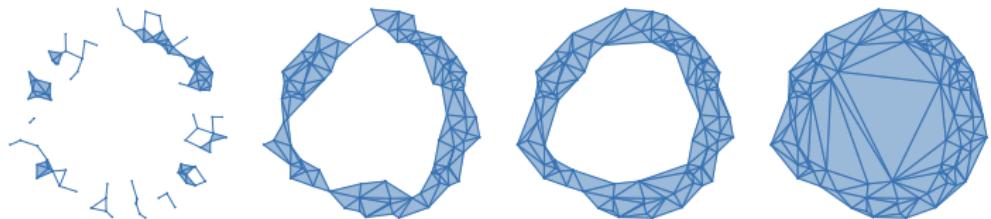
# What is persistent homology?



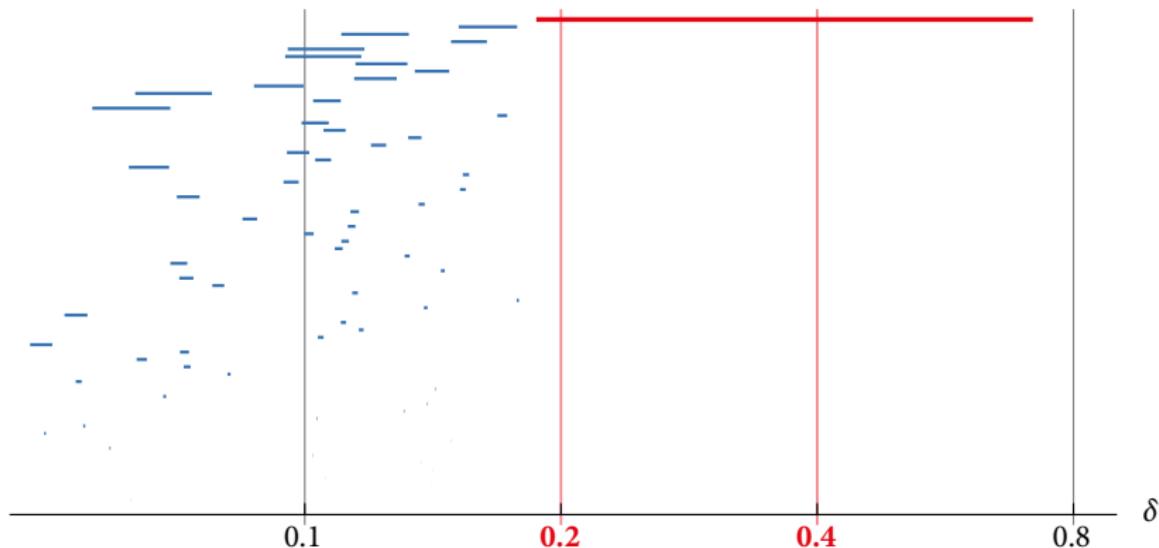
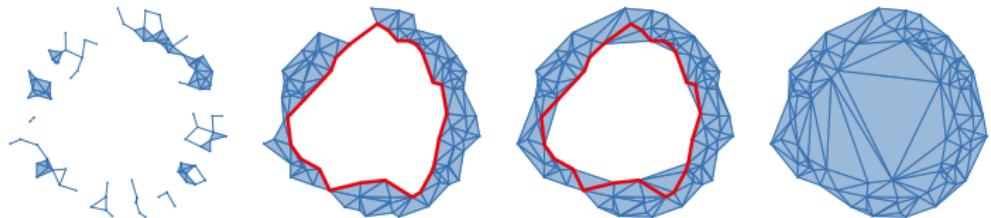
# What is persistent homology?



# What is persistent homology?



# What is persistent homology?



# What is persistent homology?

# What is persistent homology?

Persistent homology is the homology of a filtration.

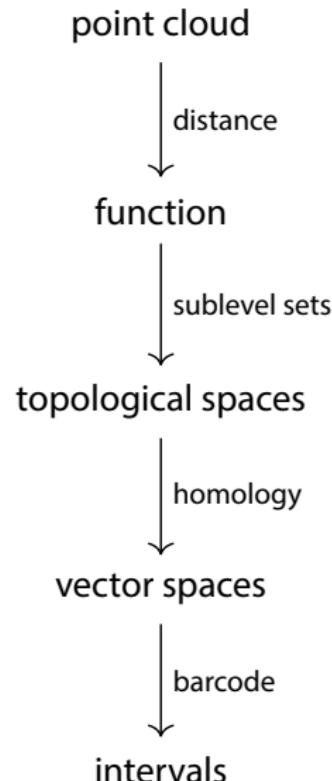
# What is persistent homology?

Persistent homology is the homology of a filtration.

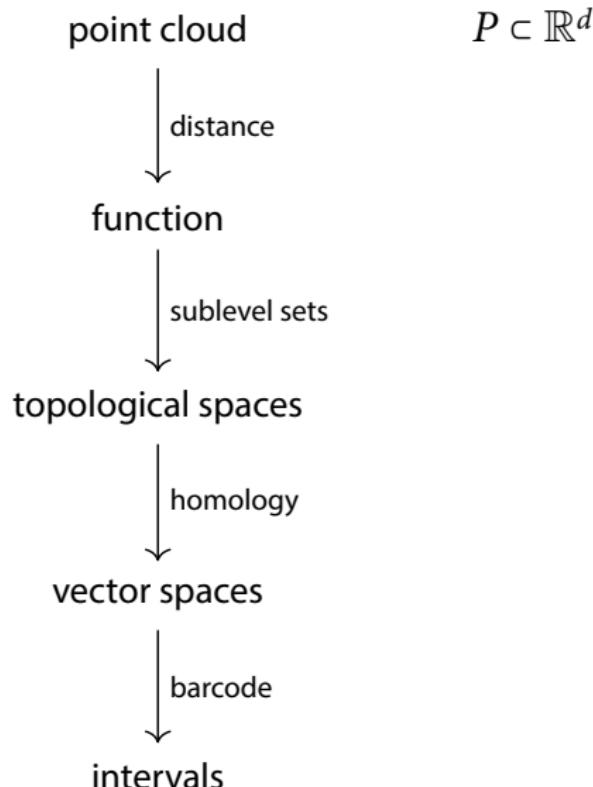
- A filtration is a certain diagram  $K : \mathbf{R} \rightarrow \mathbf{Top}$ , where  $\mathbf{R}$  is the poset category of  $(\mathbb{R}, \leq)$ :
  - A topological space  $K_t$  for each  $t \in \mathbb{R}$
  - A continuous map  $K_s \rightarrow K_t$  for each  $s \leq t \in \mathbb{R}$

# Stability

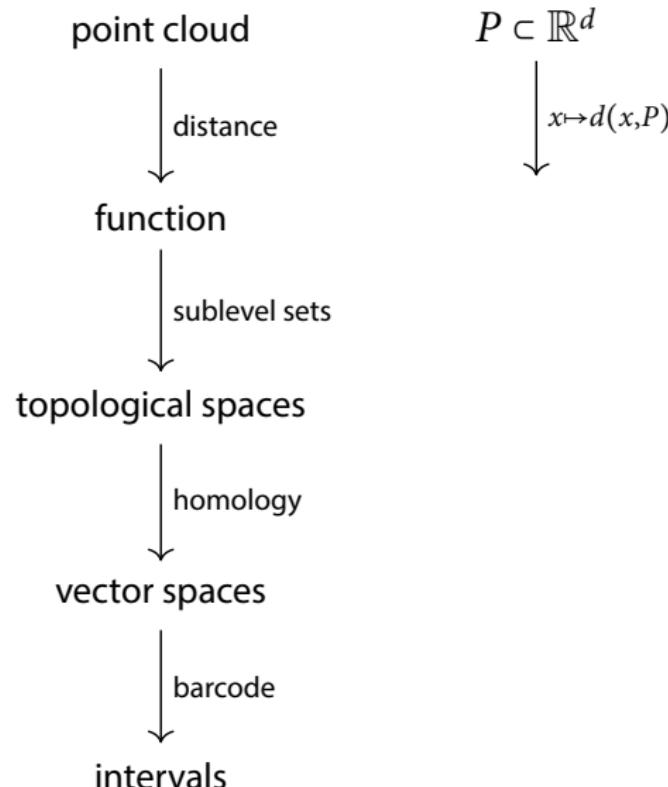
# The pipeline of topological data analysis



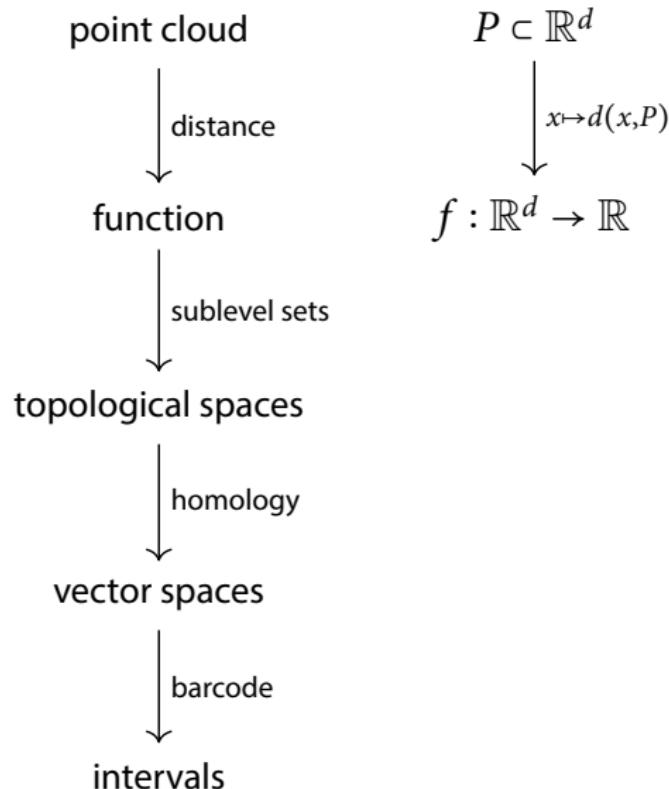
# The pipeline of topological data analysis



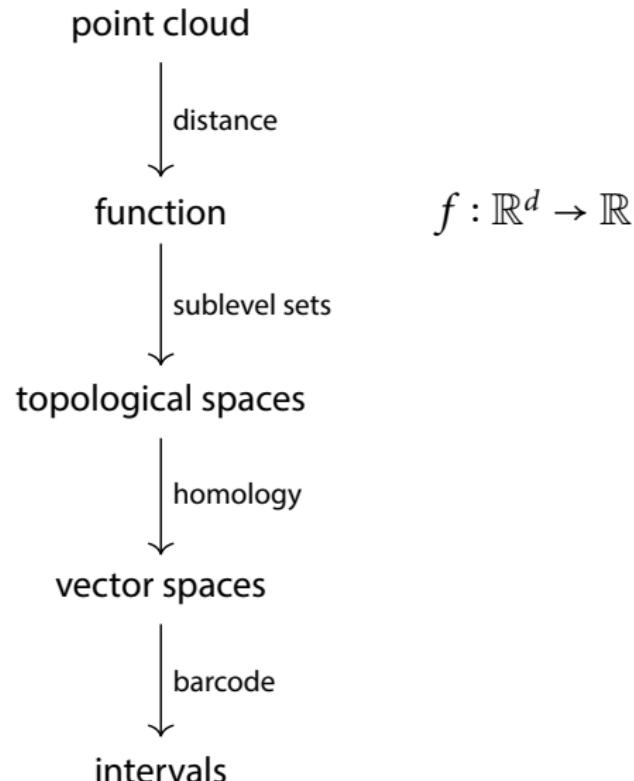
# The pipeline of topological data analysis



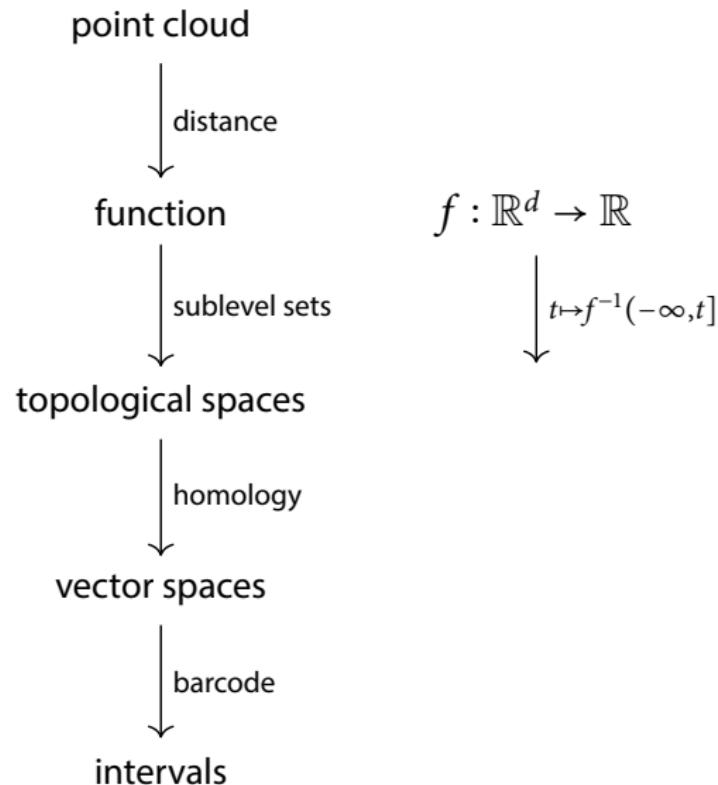
# The pipeline of topological data analysis



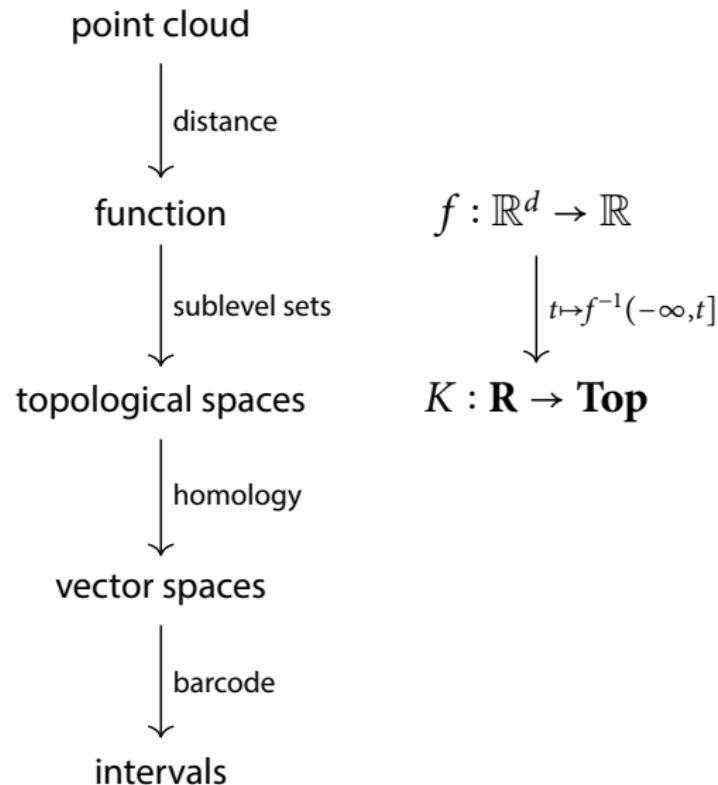
# The pipeline of topological data analysis



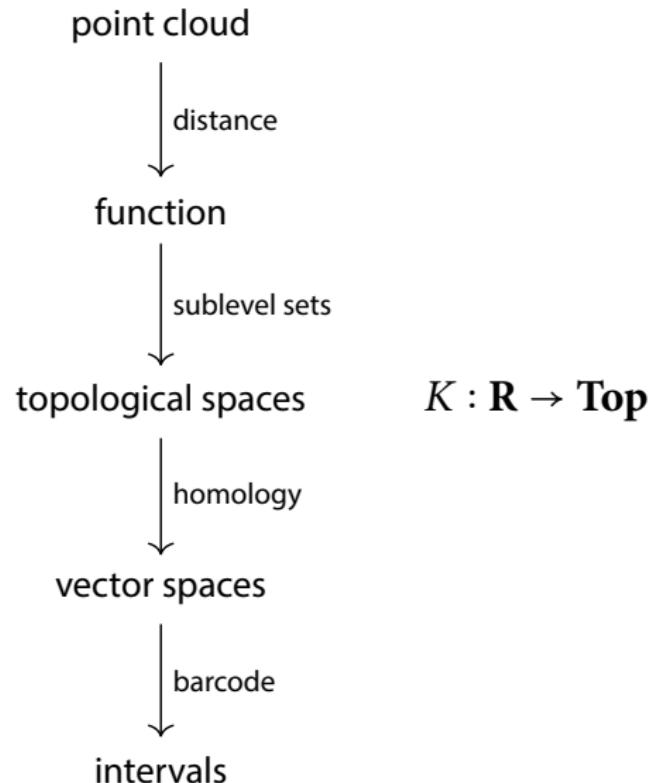
# The pipeline of topological data analysis



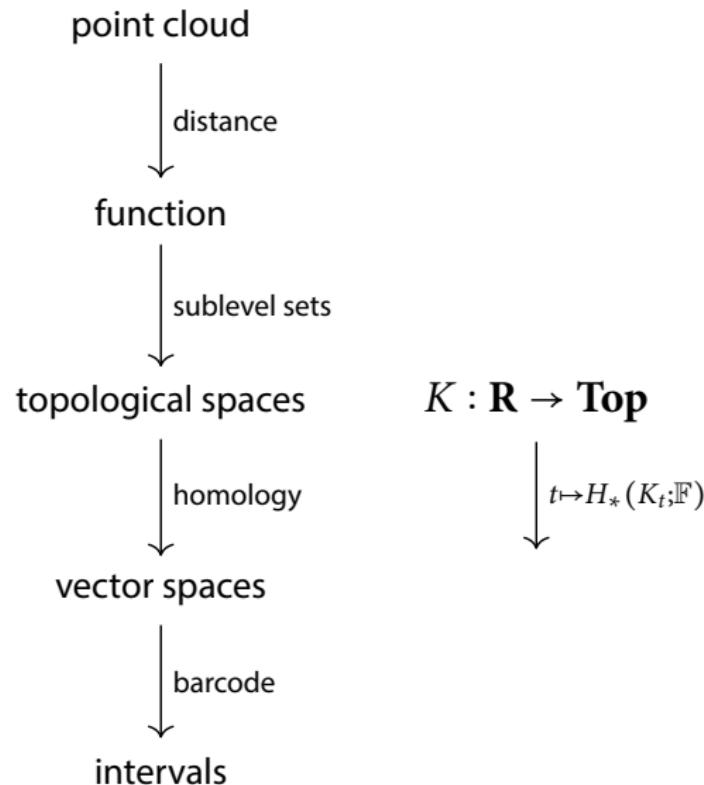
# The pipeline of topological data analysis



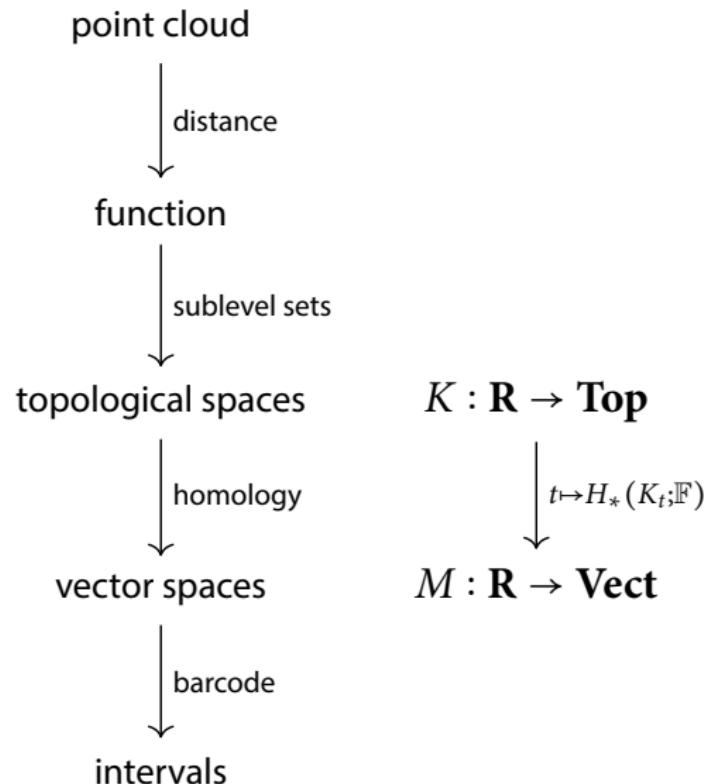
# The pipeline of topological data analysis



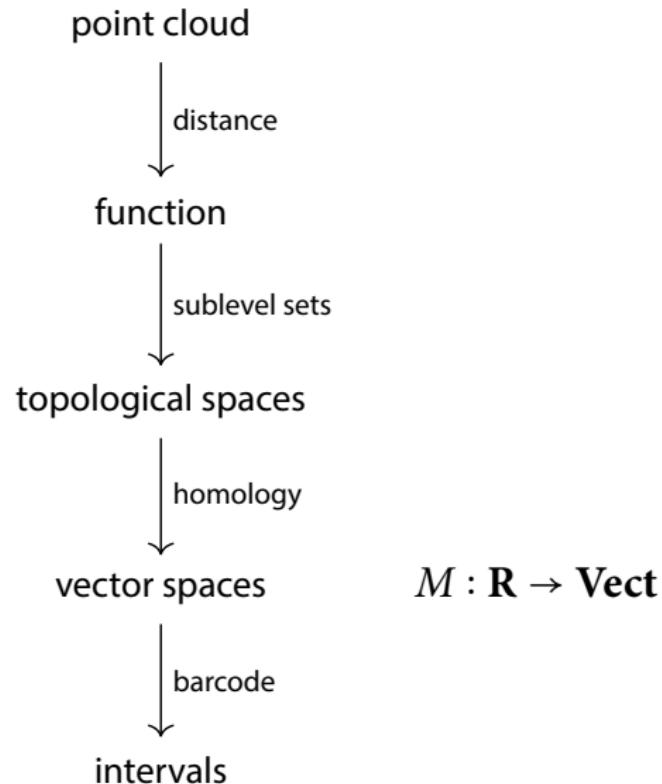
# The pipeline of topological data analysis



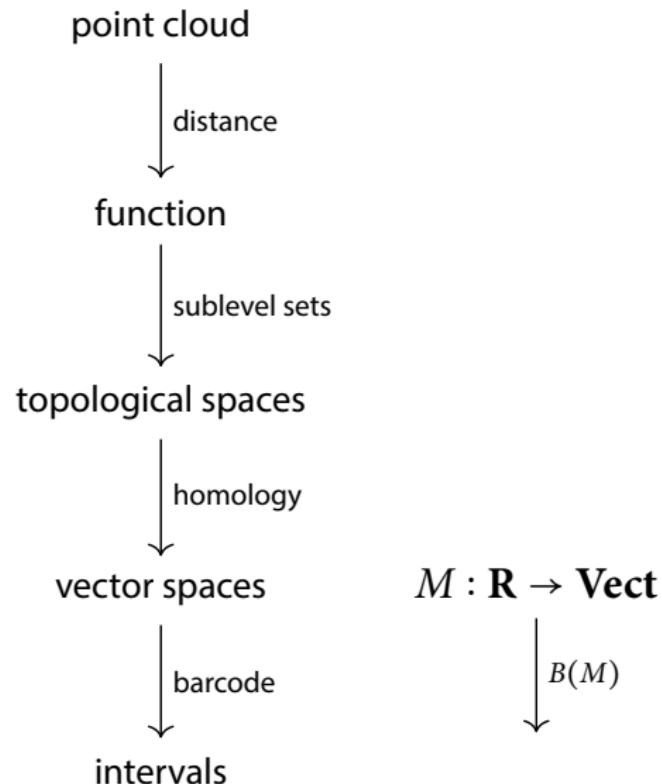
# The pipeline of topological data analysis



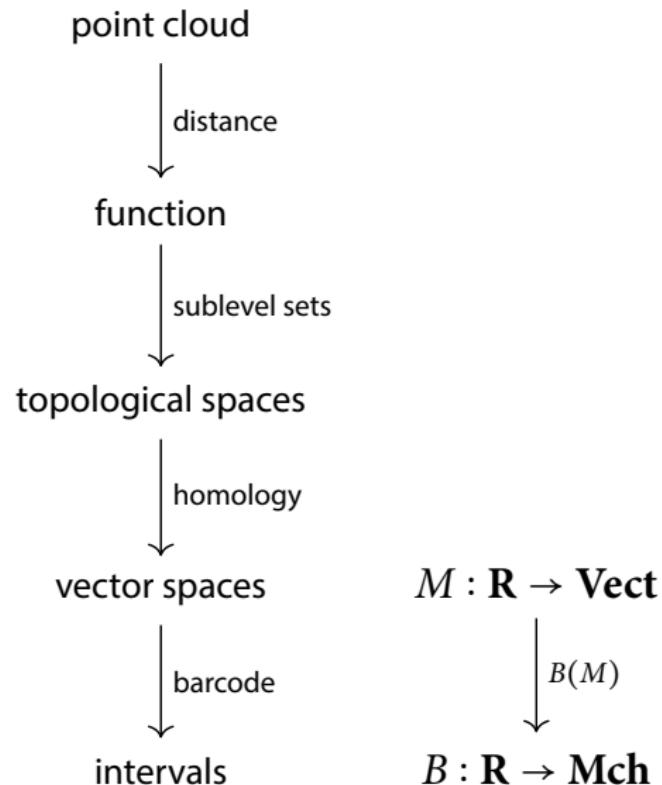
# The pipeline of topological data analysis



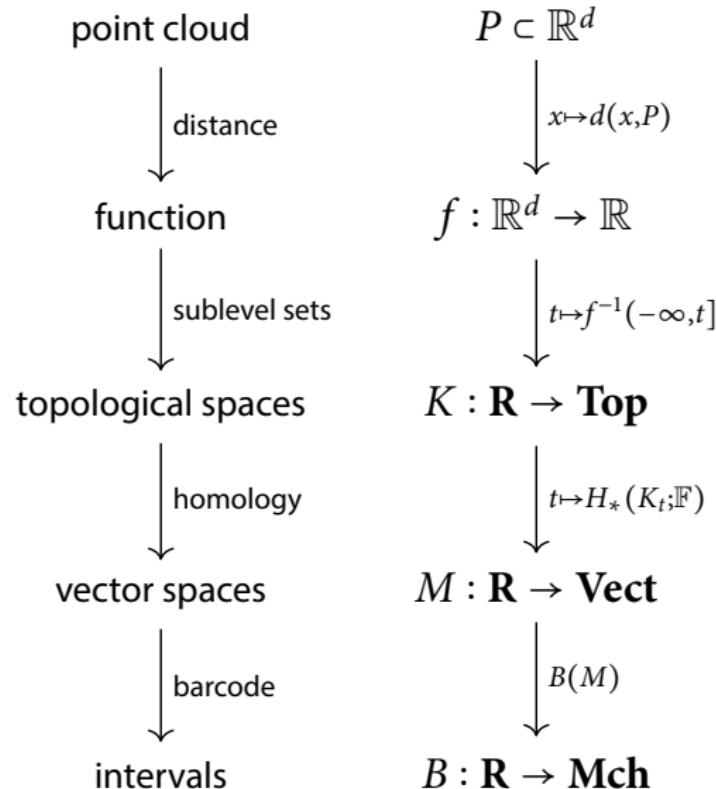
# The pipeline of topological data analysis



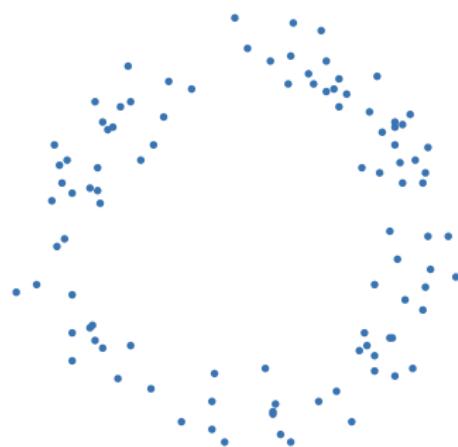
# The pipeline of topological data analysis



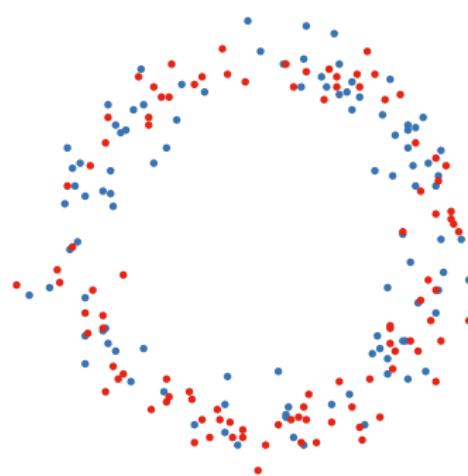
# The pipeline of topological data analysis



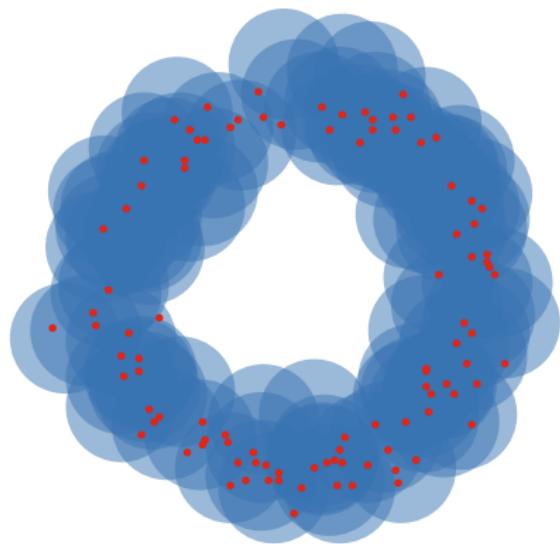
# Geometric interleavings



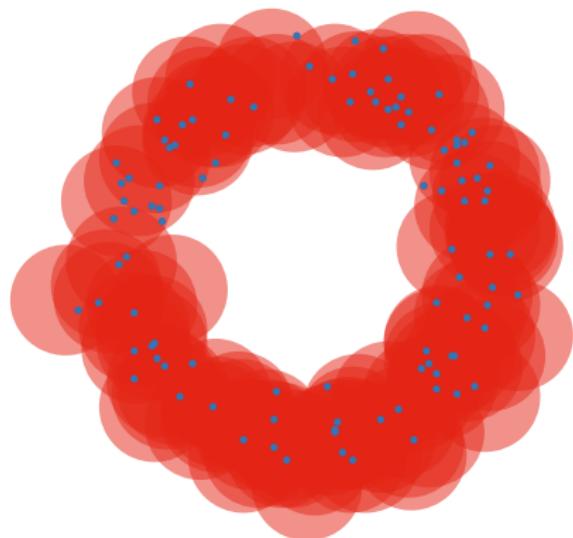
# Geometric interleavings



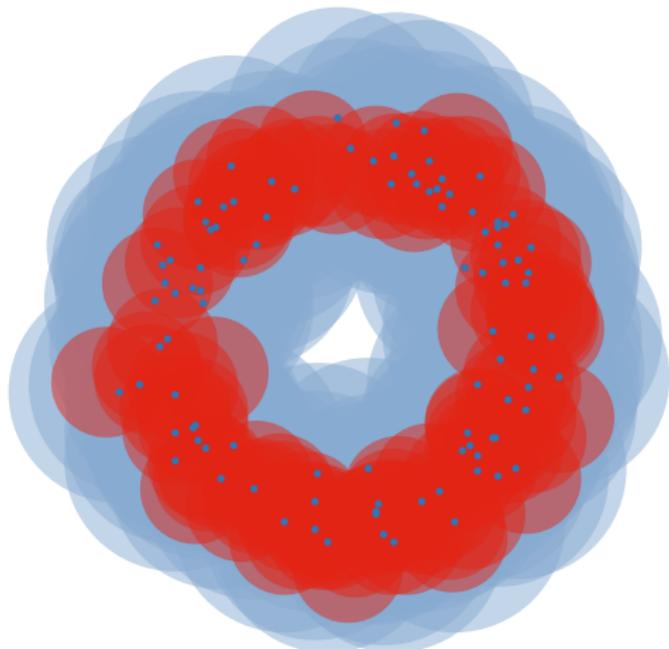
# Geometric interleavings



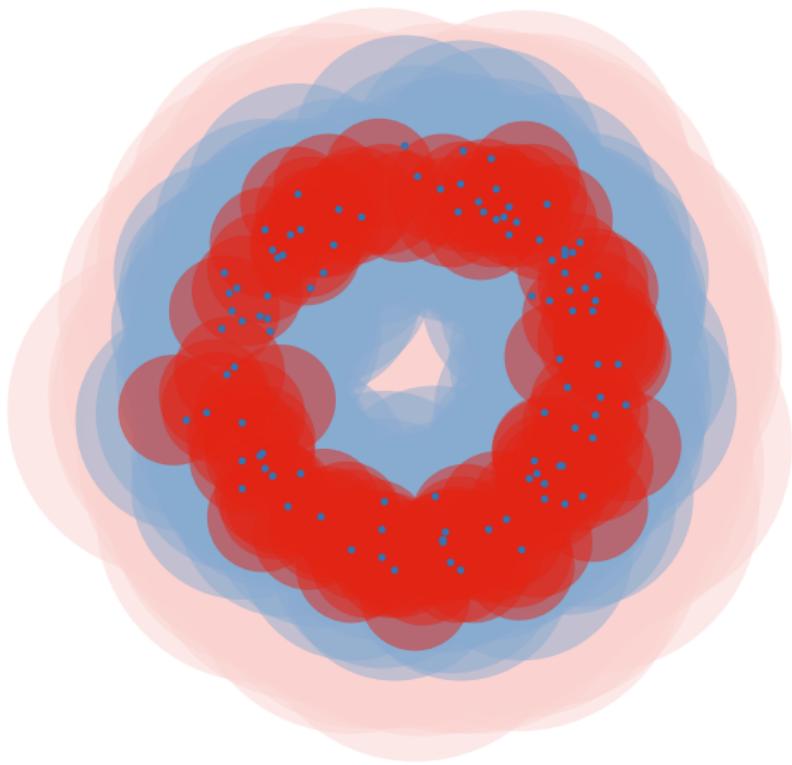
# Geometric interleavings



# Geometric interleavings



# Geometric interleavings



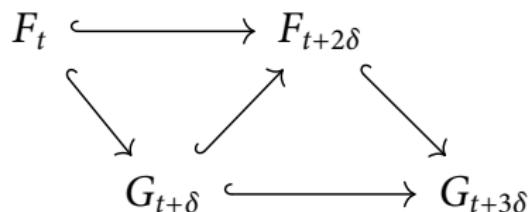
# Interleavings of sublevel sets

Let

- $F_t = f^{-1}(-\infty, t]$ ,
- $G_t = g^{-1}(-\infty, t]$ .

If  $\|f - g\|_\infty \leq \delta$  then  $F_t \subseteq G_{t+\delta}$  and  $G_t \subseteq F_{t+\delta}$ .

So the sublevel sets are  $\delta$ -interleaved:



# Interleavings

## Definition

Two diagrams (functors)  $M, N$  indexed over the reals (as a poset category) are  $\delta$ -interleaved if there are natural transformations

$$\mu_t : M_t \rightarrow N_{t+\delta}, \quad \nu_t : N_t \rightarrow M_{t+\delta} \quad \forall t \in \mathbb{R}$$

such that this diagram commutes for all  $t$ :

$$\begin{array}{ccccc} M_t & \xrightarrow{\hspace{2cm}} & M_{t+2\delta} & & \\ \searrow \mu_t & & \nearrow \nu_{t+\delta} & & \searrow \mu_{t+2\delta} \\ N_{t+\delta} & \xrightarrow{\hspace{2cm}} & N_{t+3\delta} & & \end{array}$$

- Interleavings are approximate isomorphisms, up to an index shift of  $\delta$
- Interleaving distance: infimum  $\delta$  admitting an interleaving

# Stability via functoriality?

$$\begin{array}{ccc} F_t & \xleftarrow{\quad} & F_{t+2\delta} \\ \curvearrowright & & \curvearrowright \\ G_{t+\delta} & \xleftarrow{\quad} & G_{t+3\delta} \end{array}$$

# Stability via functoriality?

$$\begin{array}{ccc} H_*(F_t) & \longrightarrow & H_*(F_{t+2\delta}) \\ \searrow & & \nearrow \\ H_*(G_{t+\delta}) & \longrightarrow & H_*(G_{t+3\delta}) \end{array}$$

# Stability via functoriality?

$$\begin{array}{ccc} H_*(F_t) & \longrightarrow & H_*(F_{t+2\delta}) \\ & \searrow & \nearrow \\ & H_*(G_{t+\delta}) & \longrightarrow H_*(G_{t+3\delta}) \end{array}$$



# Stability via functoriality?

$$\begin{array}{ccc} B(H_*(F_t)) & \rightarrow & B(H_*(F_{t+2\delta})) \\ \searrow & & \nearrow \\ & B(H_*(G_{t+\delta})) & \rightarrow B(H_*(G_{t+3\delta})) \end{array}$$



# Stability via functoriality?

$$\begin{array}{ccc} B(H_*(F_t)) & \rightarrow & B(H_*(F_{t+2\delta})) \\ & \searrow & \nearrow \\ & B(H_*(G_{t+\delta})) & \rightarrow B(H_*(G_{t+3\delta})) \end{array}$$



# Stability via functoriality?

$$\begin{array}{ccc} B(H_*(F_t)) & \rightarrow & B(H_*(F_{t+2\delta})) \\ \searrow & & \nearrow \\ & B(H_*(G_{t+\delta})) & \rightarrow B(H_*(G_{t+3\delta})) \end{array}$$



## Proposition (B, Lesnick 2015)

*There exists no functor  $\mathbf{Vect} \rightarrow \mathbf{Mch}$  sending each vector space of dimension  $d$  to a set of cardinality  $d$ .*

# Interleavings and (co)kernels

Consider a  $\delta$ -interleaving in **Vect** or **Mch**:

$$\begin{array}{ccccc} M_t & \xrightarrow{\quad} & M_{t+2\delta} & & \\ \searrow \mu_t & & \nearrow v_{t+\delta} & & \searrow \mu_{t+2\delta} \\ N_{t+\delta} & \xrightarrow{\quad} & N_{t+3\delta} & & \end{array}$$

## Proposition

The diagrams  $\ker \mu$  and  $\text{coker } \mu$  are  $2\delta$ -trivial.

- A diagram  $K$  is  $\epsilon$ -trivial if the maps  $K_t \rightarrow K_{t+\epsilon}$  are zero.
- For example, if a barcode  $B : \mathbf{R} \rightarrow \mathbf{Mch}$  is  $\epsilon$ -trivial, then the intervals of  $B$  have length at most  $\epsilon$ .

# The induced matching theorem

Theorem (B, Lesnick 2015, 2016)

Let  $\mu$  be a natural transformation between  $M, N : \mathbf{R} \rightarrow \mathbf{Vect}$ . Then

- $\mu$  induces a natural transformation  $X(\mu)$  between the barcodes  $B(M), B(N) : \mathbf{R} \rightarrow \mathbf{Mch}$ .
- If  $\ker \mu$  is  $\epsilon$ -trivial then so is  $\ker X(\mu)$ , and likewise for coker.

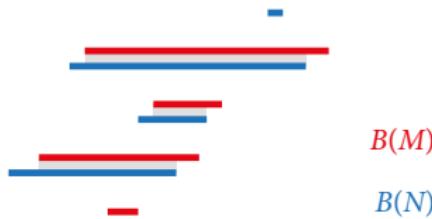


# The induced matching theorem

Theorem (B, Lesnick 2015, 2016)

Let  $\mu$  be a natural transformation between  $M, N : \mathbf{R} \rightarrow \mathbf{Vect}$ . Then

- $\mu$  induces a natural transformation  $X(\mu)$  between the barcodes  $B(M), B(N) : \mathbf{R} \rightarrow \mathbf{Mch}$ .
- If  $\ker \mu$  is  $\epsilon$ -trivial then so is  $\ker X(\mu)$ , and likewise for  $\text{coker}$ .



This implies the algebraic stability for persistence barcodes:

Theorem (Chazal et al. 2009; B, Lesnick 2015, 2016)

Two diagrams  $M, N : \mathbf{R} \rightarrow \mathbf{Vect}$  are  $\delta$ -interleaved if and only if their barcodes  $B(M), B(N)$  are.

# History

# When was persistent homology invented?

- [Edelsbrunner/Letscher/Zomorodian 2000]

# When was persistent homology invented?

- [Edelsbrunner/Letscher/Zomorodian 2000]
- [Robbins 1999]

# When was persistent homology invented?

- [Edelsbrunner/Letscher/Zomorodian 2000]
- [Robbins 1999]
- [Frosini 1990]

# When was persistent homology invented?

- [Edelsbrunner/Letscher/Zomorodian 2000]
- [Robbins 1999]
- [Frosini 1990]
- [Leray 1946]?

When was persistent homology invented first?

# When was persistent homology invented first?

ANNALS OF MATHEMATICS  
Vol. 41, No. 2, April, 1940

## RANK AND SPAN IN FUNCTIONAL TOPOLOGY

BY MARSTON MORSE

(Received August 9, 1939)

### 1. Introduction.

The analysis of functions  $F$  on metric spaces  $M$  of the type which appear in variational theories is made difficult by the fact that the critical limits, such as absolute minima, relative minima, minimax values etc., are in general infinite in number. These limits are associated with relative  $k$ -cycles of various dimensions and are classified as 0-limits, 1-limits etc. The number of  $k$ -limits suitably counted is called the  $k^{\text{th}}$  type number  $m_k$  of  $F$ . The theory seeks to establish relations between the numbers  $m_k$  and the connectivities  $p_k$  of  $M$ . The numbers  $p_k$  are finite in the most important applications. It is otherwise with the numbers  $m_k$ .

The theory has been able to proceed provided one of the following hypotheses is satisfied. The critical limits cluster at most at  $+\infty$ ; the critical points are isolated;<sup>1</sup> the problem is defined by analytic functions; the critical limits taken in their natural order are well-ordered. These conditions are not generally fulfilled. The generality of the theory rested upon the fact that the cases treated approximate in a certain sense the most general problems which it is

# When was persistent homology invented first?

Web Images More... Sign in

Google Scholar 9 results (0.02 sec) My Citations ▾

All citations Rank and span in functional topology

Articles  Search within citing articles

Case law

My library

Exact homomorphism sequences in homology theory ed.ac.uk [PDF]

JL Kelley, E Pitcher - Annals of Mathematics, 1947 - JSTOR

The developments of this paper stem from the attempts of one of the authors to deduce relations between homology groups of a complex and homology groups of a complex which is its image under a simplicial map. Certain relations were deduced (see [EP 1] and [EP 2] ...)

Any time Cited by 46 Related articles All 3 versions Cite Save More

Since 2016

Since 2015

Since 2012

Custom range...

Sort by relevance Cited by 24 Related articles All 4 versions Cite Save More

Sort by date

include citations

Create alert

Unstable minimal surfaces of higher topological structure

M Morse, CB Tompkins - Duke Math. J., 1941 - projecteuclid.org

1. Introduction. We are concerned with extending the calculus of variations in the large to multiple integrals. The problem of the existence of minimal surfaces of unstable type contains many of the typical difficulties, especially those of a topological nature. Having studied this ...

Cited by 19 Related articles All 2 versions Cite Save

[PDF] Persistence in discrete Morse theory psu.edu [PDF]

U Bauer - 2011 - Citeseer

The goal of this thesis is to bring together two different theories about critical points of a scalar function and their relation to topology: Discrete Morse theory and Persistent

# When was persistent homology invented first?

BULLETIN (New Series) OF THE  
AMERICAN MATHEMATICAL SOCIETY  
Volume 3, Number 3, November 1980

## MARSTON MORSE AND HIS MATHEMATICAL WORKS

BY RAOUL BOTT<sup>1</sup>

**1. Introduction.** Marston Morse was born in 1892, so that he was 33 years old when in 1925 his paper *Relations between the critical points of a real-valued function of n independent variables* appeared in the Transactions of the American Mathematical Society. Thus Morse grew to maturity just at the time when the subject of Analysis Situs was being shaped by such masters<sup>2</sup> as Poincaré, Veblen, L. E. J. Brouwer, G. D. Birkhoff, Lefschetz and Alexander, and it was Morse's genius and destiny to discover one of the most beautiful and far-reaching relations between this fledgling and Analysis; a relation which is now known as *Morse Theory*.

In retrospect all great ideas take on a certain simplicity and inevitability, partly because they shape the whole subsequent development of the subject. And so to us, today, Morse Theory seems natural and inevitable. However one only has to glance at these early papers to see what a tour de force it was in the 1920's to go from the mini-max principle of Birkhoff to the Morse inequalities, let alone extend these inequalities to function spaces, so that by the early 30's Morse could establish the theorem that for any Riemann

# When was persistent homology invented first?

inequalities between the dimensions of the  $H_i$  and those of  $H(A_i)$ . Thus the Morse inequalities already reflect a certain part of the “Spectral Sequence magic”, and a modern and tremendously general account of Morse’s work on rank and span in the framework of Leray’s theory was developed by Deheuvels [D] in the 50’s.

Unfortunately both Morse’s and Deheuvel’s papers are not easy reading. On the other hand there is no question in my mind that the papers [36] and [44] constitute another tour de force by Morse. Let me therefore illustrate rather than explain some of the ideas of the rank and span theory in a very simple and tame example.

In the figure which follows I have drawn a homeomorph of  $M = S^1$  in the plane, and I will be studying the height function  $F = y$  on  $M$ .

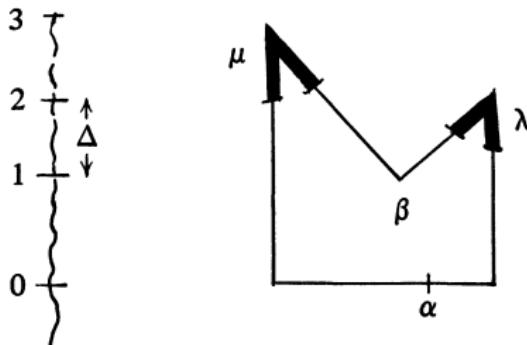


FIGURE 8

The values  $a$  where  $H(a, a^-) \neq 0$  are indicated on the left, and correspond

# *Morse's functional topology*

Key aspects:

- early precursor of persistence and spectral sequences

# *Morse's functional topology*

Key aspects:

- early precursor of persistence and spectral sequences
- uses Vietoris homology with field coefficients

# *Morse's functional topology*

Key aspects:

- early precursor of persistence and spectral sequences
- uses Vietoris homology with field coefficients
- applies to a broad class of functions on metric spaces  
(not necessarily continuous)

# Morse's functional topology

Key aspects:

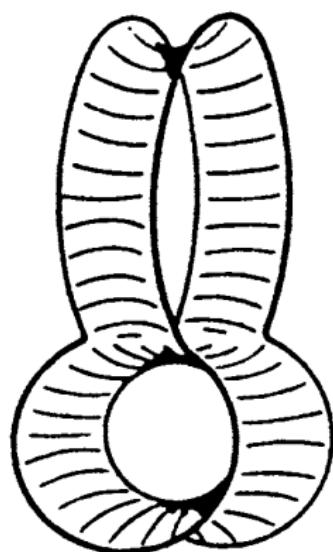
- early precursor of persistence and spectral sequences
- uses Vietoris homology with field coefficients
- applies to a broad class of functions on metric spaces  
(not necessarily continuous)
- inclusions of sublevel sets have finite rank homology  
( $q$ -tame persistent homology)

# Morse's functional topology

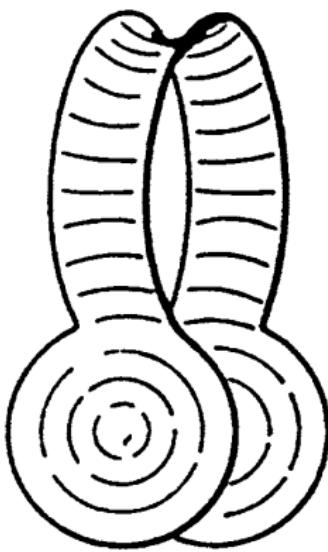
Key aspects:

- early precursor of persistence and spectral sequences
- uses Vietoris homology with field coefficients
- applies to a broad class of functions on metric spaces  
(not necessarily continuous)
- inclusions of sublevel sets have finite rank homology  
( $q$ -tame persistent homology)
- focus on controlled behavior in pathological cases

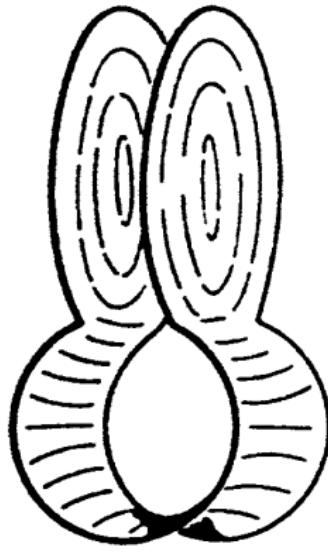
# Motivation and application: minimal surfaces



(a)



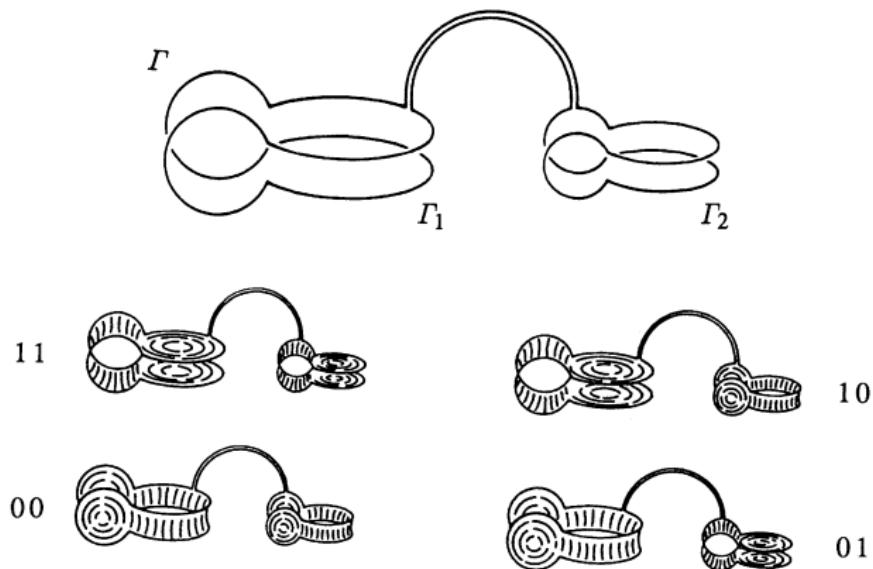
(b)



(c)

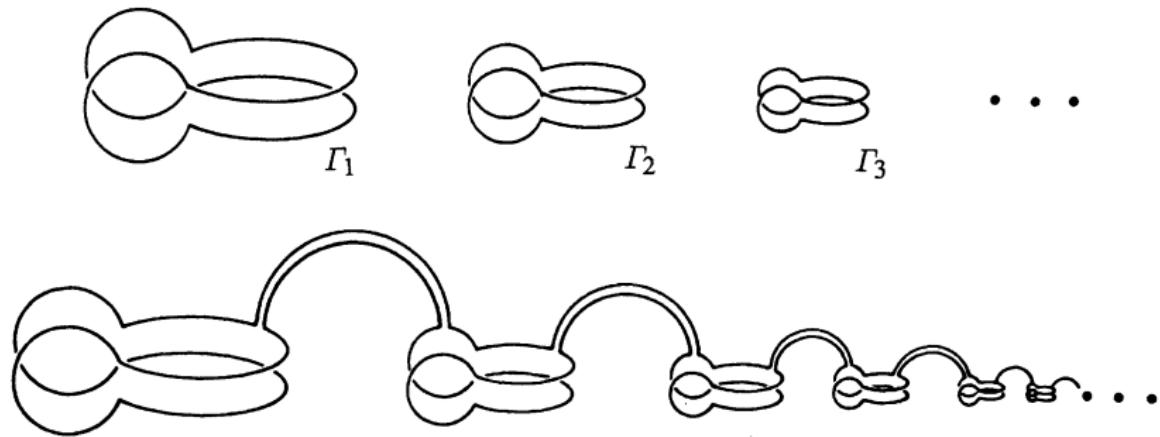
(from Dierkes et al.: Minimal Surfaces, Springer 2010)

# Motivation and application: minimal surfaces



(from Dierkes et al.: Minimal Surfaces, Springer 2010)

# Motivation and application: minimal surfaces



(from Dierkes et al.: Minimal Surfaces, Springer 2010)

# Existence of unstable minimal surfaces

Using persistent homology:

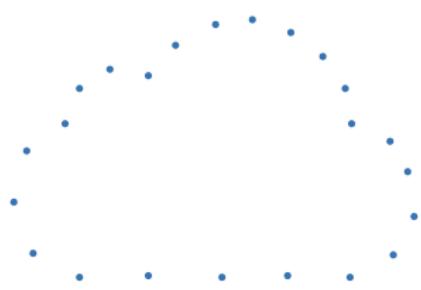
- Number of  $\epsilon$ -persistent critical points (minimal surfaces) is finite for any  $\epsilon > 0$
- Morse inequalities for  $\epsilon$ -persistent critical points

Theorem (Morse, Tompkins 1939)

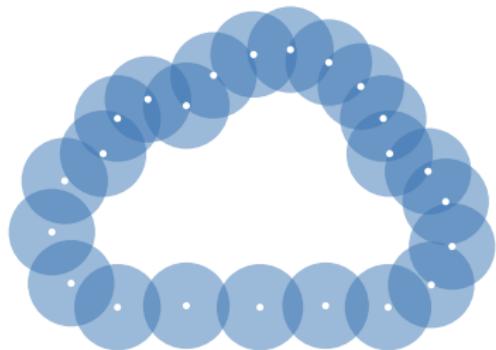
*There is a  $C_1$  curve bounding an unstable minimal surface (a critical point of the area functional with index 1).*

# Complexes

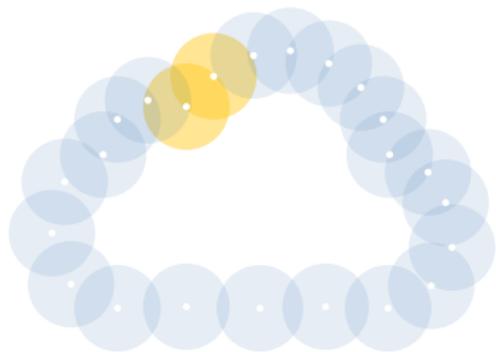
# Čech and Delaunay complexes



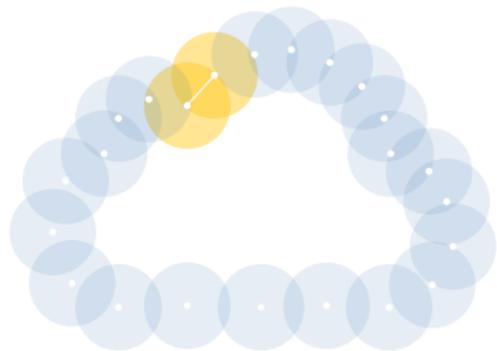
# Čech and Delaunay complexes



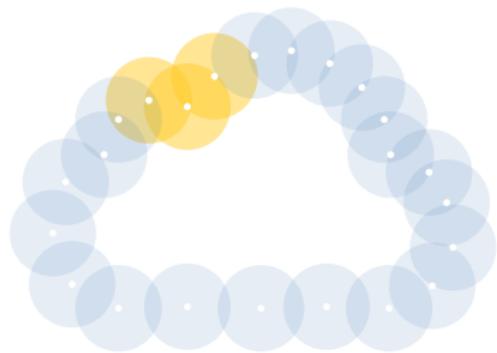
# Čech and Delaunay complexes



# Čech and Delaunay complexes



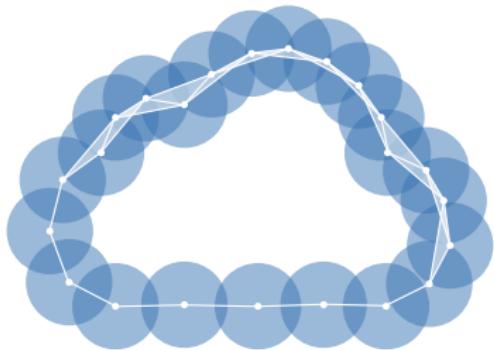
# Čech and Delaunay complexes



# Čech and Delaunay complexes

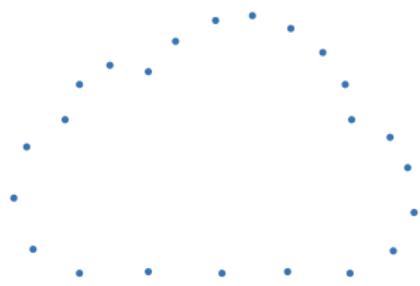


# Čech and Delaunay complexes

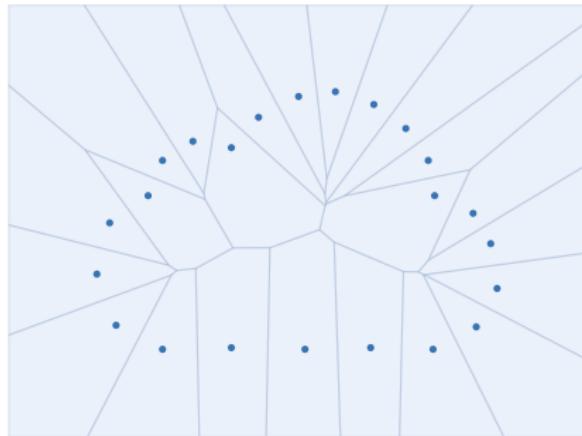


$$\text{Cech}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} B_r(p) \neq \emptyset \right\}$$

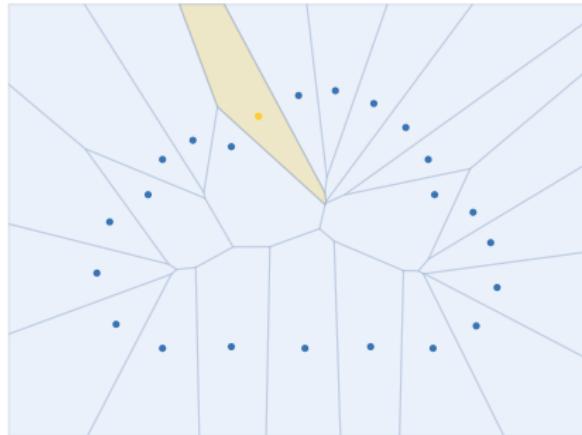
# Čech and Delaunay complexes



# Čech and Delaunay complexes



# Čech and Delaunay complexes



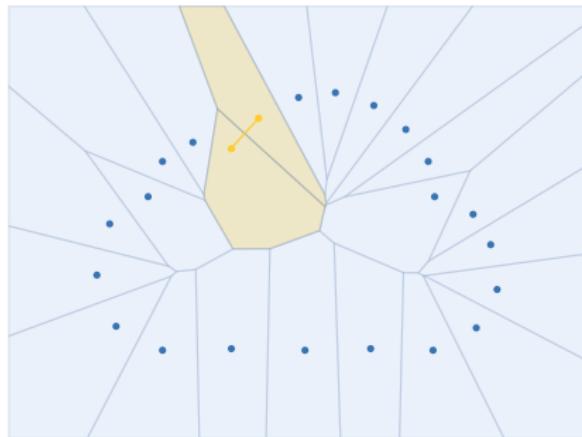
$$\text{Vor}(p, X) = \{a \in \mathbb{R}^n \mid d(a, p) = d(a, X)\}$$

# Čech and Delaunay complexes



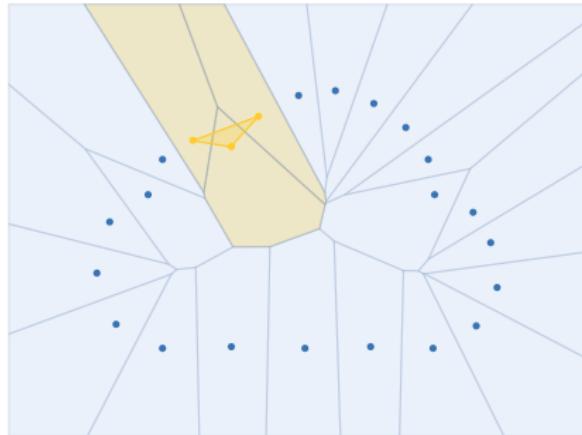
$$\text{Vor}(p, X) = \{a \in \mathbb{R}^n \mid d(a, p) = d(a, X)\}$$

# Čech and Delaunay complexes



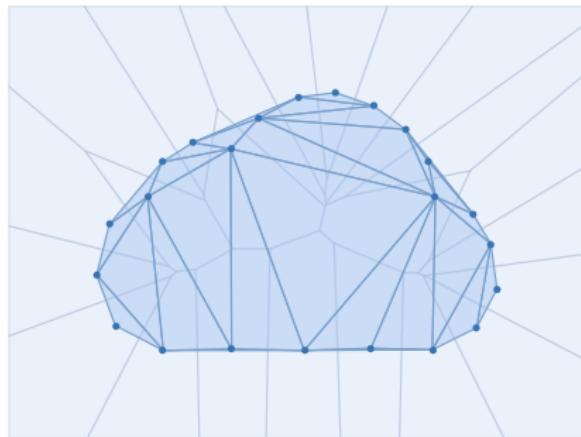
$$\text{Vor}(p, X) = \{a \in \mathbb{R}^n \mid d(a, p) = d(a, X)\}$$

# Čech and Delaunay complexes



$$\text{Vor}(p, X) = \{a \in \mathbb{R}^n \mid d(a, p) = d(a, X)\}$$

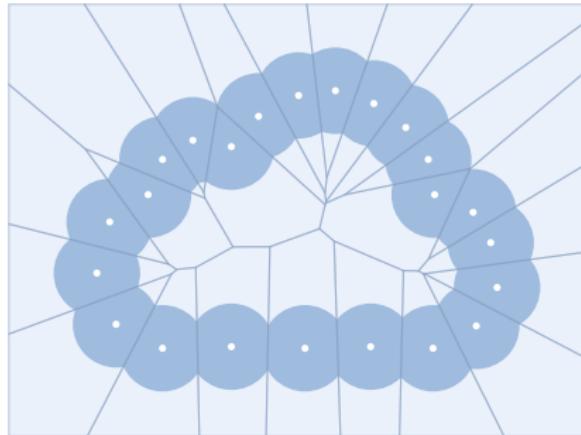
# Čech and Delaunay complexes



$$\text{Vor}(p, X) = \{a \in \mathbb{R}^n \mid d(a, p) = d(a, X)\}$$

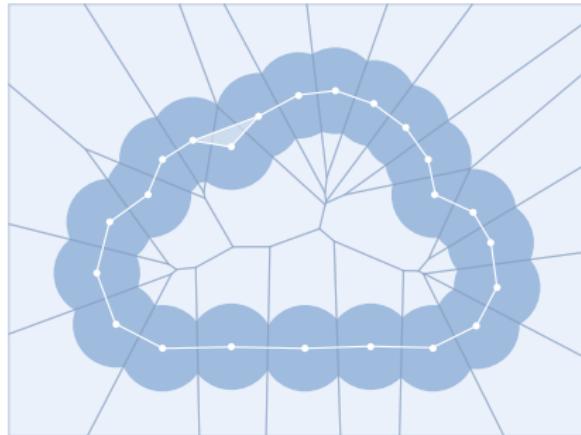
$$\text{Del}(X) = \{Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}(p, X) \neq \emptyset\}$$

# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

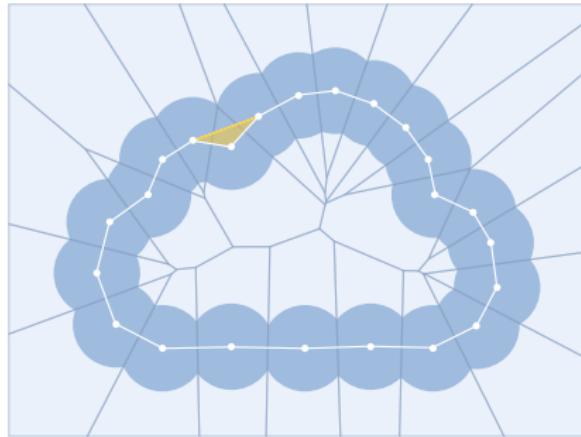
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

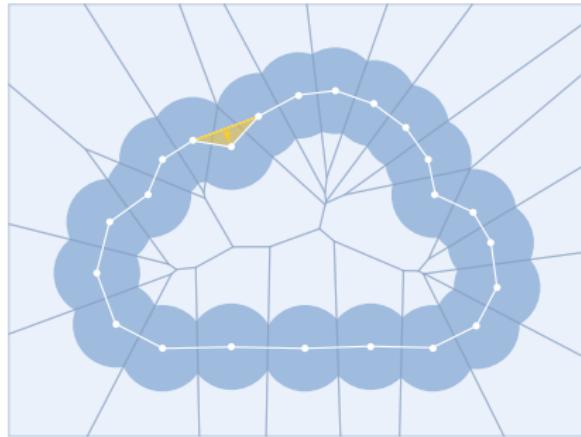
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

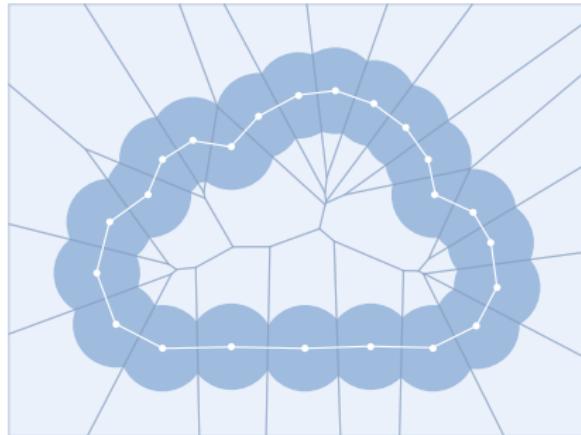
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

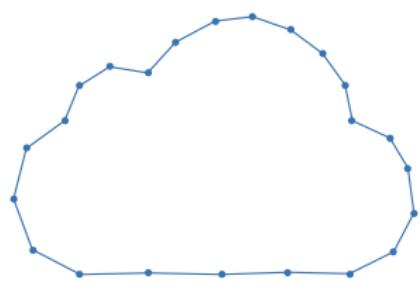
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

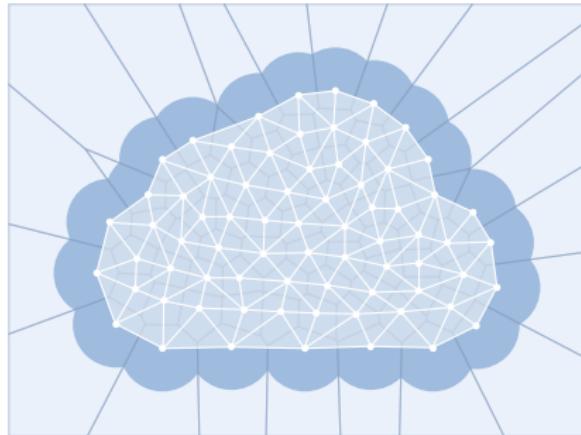
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

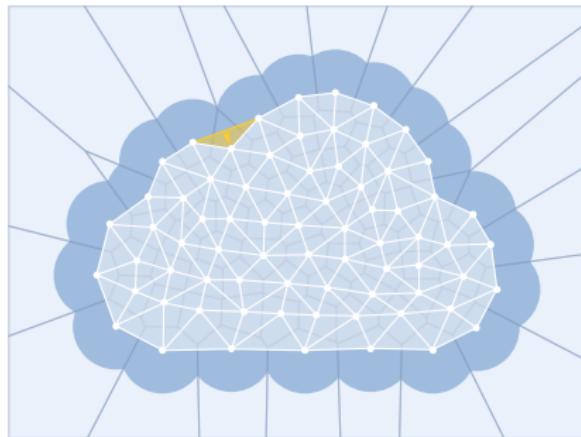
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

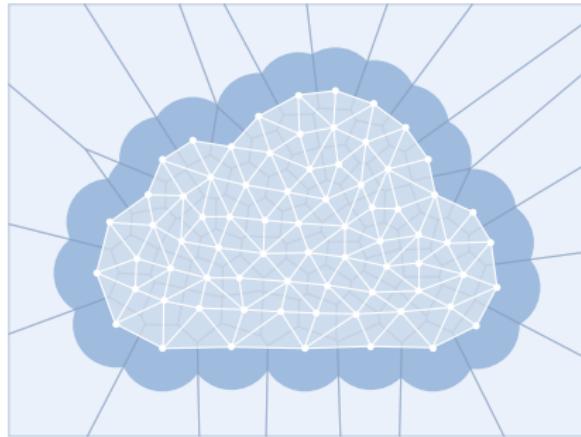
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

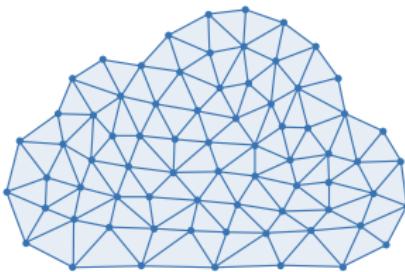
# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

# Čech and Delaunay complexes



$$\text{Vor}_r(p, X) = B_r(p) \cap \text{Vor}(p, X)$$

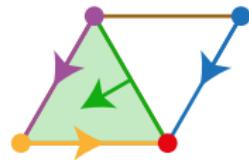
$$\text{Del}_r(X) = \left\{ Q \subseteq X \mid \bigcap_{p \in Q} \text{Vor}_r(p, X) \neq \emptyset \right\}$$

# Discrete Morse theory

## Definition (Forman 1998)

A *discrete vector field* on a cell complex  
is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .

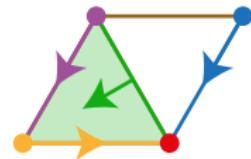


# Discrete Morse theory

## Definition (Forman 1998)

A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .



A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

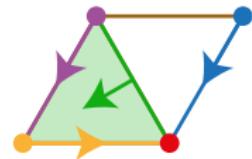
- sublevel sets are subcomplexes, and

# Discrete Morse theory

## Definition (Forman 1998)

A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .



A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

- sublevel sets are subcomplexes, and

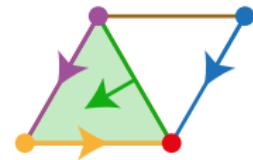


# Discrete Morse theory

## Definition (Forman 1998)

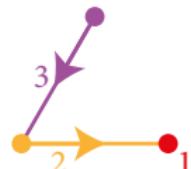
A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .



A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

- sublevel sets are subcomplexes, and

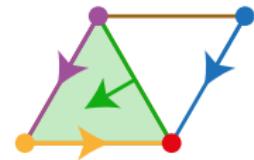


# Discrete Morse theory

## Definition (Forman 1998)

A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .



A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

- sublevel sets are subcomplexes, and



# Discrete Morse theory

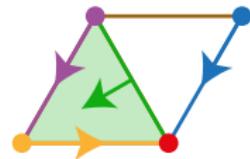
## Definition (Forman 1998)

A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .

A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

- sublevel sets are subcomplexes, and



# Discrete Morse theory

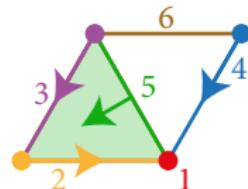
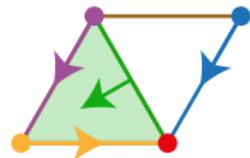
## Definition (Forman 1998)

A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .

A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

- sublevel sets are subcomplexes, and

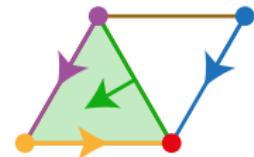


# Discrete Morse theory

## Definition (Forman 1998)

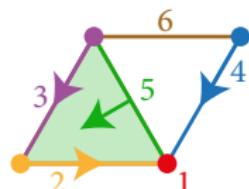
A *discrete vector field* on a cell complex is a partition of the set of simplices into

- singleton sets  $\{\phi\}$  (*critical cells*), and
- pairs  $\{\sigma, \tau\}$ , where  $\sigma$  is a facet of  $\tau$ .



A function  $f : K \rightarrow \mathbb{R}$  on a cell complex is a *discrete Morse function* if

- sublevel sets are subcomplexes, and
- level sets form a discrete vector field.



## Fundamental theorem of discrete Morse theory

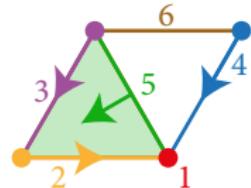
Let  $f$  be a discrete Morse function on a cell complex  $K$ .

# Fundamental theorem of discrete Morse theory

Let  $f$  be a discrete Morse function on a cell complex  $K$ .

## Theorem (Forman 1998)

If  $(s, t]$  contains no critical value of  $f$ ,  
then the sublevel set  $K_t$  collapses to  $K_s$   
(written as  $K_t \searrow K_s$ ).

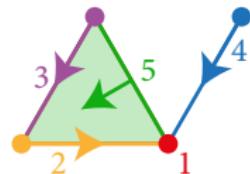


# Fundamental theorem of discrete Morse theory

Let  $f$  be a discrete Morse function on a cell complex  $K$ .

## Theorem (Forman 1998)

If  $(s, t]$  contains no critical value of  $f$ ,  
then the sublevel set  $K_t$  collapses to  $K_s$   
(written as  $K_t \searrow K_s$ ).

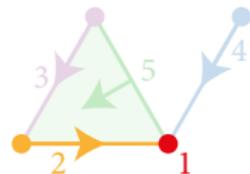


# Fundamental theorem of discrete Morse theory

Let  $f$  be a discrete Morse function on a cell complex  $K$ .

## Theorem (Forman 1998)

If  $(s, t]$  contains no critical value of  $f$ ,  
then the sublevel set  $K_t$  collapses to  $K_s$   
(written as  $K_t \searrow K_s$ ).



# Fundamental theorem of discrete Morse theory

Let  $f$  be a discrete Morse function on a cell complex  $K$ .

## Theorem (Forman 1998)

If  $(s, t]$  contains no critical value of  $f$ ,  
then the sublevel set  $K_t$  collapses to  $K_s$   
(written as  $K_t \searrow K_s$ ).



## Corollary

$K$  is homotopy-equivalent to a cell complex  $M$  built from the critical cells of  $f$ .

# Fundamental theorem of discrete Morse theory

Let  $f$  be a discrete Morse function on a cell complex  $K$ .

## Theorem (Forman 1998)

If  $(s, t]$  contains no critical value of  $f$ ,  
then the sublevel set  $K_t$  collapses to  $K_s$   
(written as  $K_t \searrow K_s$ ).



## Corollary

$K$  is homotopy-equivalent to a cell complex  $M$  built from the critical cells of  $f$ .

This homotopy equivalence is compatible with the filtration.

## Corollary

$K$  and  $M$  have isomorphic persistent homology (with regard to  $f$ ).

# Morse theory for Čech and Delaunay complexes

## Proposition

*The Čech complexes and the Delaunay (alpha) complexes are sublevel set filtrations of generalized discrete Morse functions.*

# Morse theory for Čech and Delaunay complexes

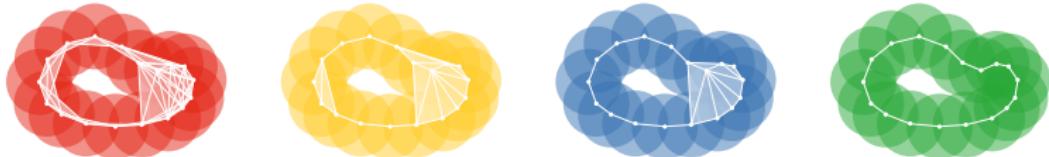
## Proposition

*The Čech complexes and the Delaunay (alpha) complexes are sublevel set filtrations of generalized discrete Morse functions.*

## Theorem (B., Edelsbrunner 2016, Trans. AMS)

*Čech, Delaunay–Čech, Delaunay, and Wrap complexes are naturally homotopy equivalent through a sequence of collapses*

$$\text{Cech}_r X \searrow \text{Cech}_r X \cap \text{Del} X \searrow \text{Del}_r X \searrow \text{Wrap}_r X.$$



## Vietoris–Rips complexes

Consider a finite metric space  $(X, d)$ .

The *Vietoris–Rips complex* is the simplicial complex

$$\text{Rips}_t(X) = \{S \subseteq X \mid \text{diam } S \leq t\}$$

- 1-skeleton: all edges with pairwise distance  $\leq t$
- all possible higher simplices (flag complex)
- Vietoris–Rips complex are *not* generically sublevel sets of discrete Morse functions

## Vietoris–Rips complexes

Consider a finite metric space  $(X, d)$ .

The *Vietoris–Rips complex* is the simplicial complex

$$\text{Rips}_t(X) = \{S \subseteq X \mid \text{diam } S \leq t\}$$

- 1-skeleton: all edges with pairwise distance  $\leq t$
- all possible higher simplices (flag complex)
- Vietoris–Rips complex are *not* generically sublevel sets of discrete Morse functions

Note: for  $t \geq \text{diam } X$ ,  $\text{Rips}_t(X)$  is the full simplex with vertices  $X$

- Number of simplices grows exponentially in dimension
- Computation is one of the most important challenges in applied topology!

# Computation

# Computing Vietoris–Rips persistence

Goal:

- compute homology  $H_d(\text{Rips}_t(X))$   
(for all  $t$  and  $0 \leq d < k$ )
- together with induced maps  $H_d(\text{Rips}_s(X) \hookrightarrow \text{Rips}_t(X))$   
(for all  $s \leq t$ )

# Demo: Ripser

Example data set:

- 192 points on  $\mathbb{S}^2$
- persistent homology barcodes up to dimension 2
- over 56 mio. simplices in 3-skeleton

# Demo: Ripser

Example data set:

- 192 points on  $\mathbb{S}^2$
- persistent homology barcodes up to dimension 2
- over 56 mio. simplices in 3-skeleton

Some previous software:

- javaplex (Stanford): 3200 seconds, 12 GB
- Dionysus (Duke): 615 seconds, 3.4 GB
- DIPHA (IST Austria): 50 seconds, 6 GB
- GUDHI (INRIA): 60 seconds, 3 GB

# Demo: Ripser

Example data set:

- 192 points on  $\mathbb{S}^2$
- persistent homology barcodes up to dimension 2
- over 56 mio. simplices in 3-skeleton

Some previous software:

- javaplex (Stanford): 3200 seconds, 12 GB
- Dionysus (Duke): 615 seconds, 3.4 GB
- DIPHA (IST Austria): 50 seconds, 6 GB
- GUDHI (INRIA): 60 seconds, 3 GB

Ripser: 1.2 seconds, 160 MB

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies
- support for
  - coefficients in a prime field  $\mathbb{Z}/p\mathbb{Z}$

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies
- support for
  - coefficients in a prime field  $\mathbb{Z}/p\mathbb{Z}$
  - sparse distance matrices for distance threshold

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies
- support for
  - coefficients in a prime field  $\mathbb{Z}/p\mathbb{Z}$
  - sparse distance matrices for distance threshold
- open source (<http://ripser.org>)
  - released in July 2016

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies
- support for
  - coefficients in a prime field  $\mathbb{Z}/p\mathbb{Z}$
  - sparse distance matrices for distance threshold
- open source (<http://ripser.org>)
  - released in July 2016
- online version (<http://live.ripser.org>)
  - launched in August 2016

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies
- support for
  - coefficients in a prime field  $\mathbb{Z}/p\mathbb{Z}$
  - sparse distance matrices for distance threshold
- open source (<http://ripser.org>)
  - released in July 2016
- online version (<http://live.ripser.org>)
  - launched in August 2016
- most efficient software for Vietoris–Rips persistence by far
  - computes  $H^2$  barcode for 50000 random points on a torus in 136 seconds / 9 GB

# Ripser

A software for computing Vietoris–Rips persistence barcodes

- around 1000 lines of C++ code, no external dependencies
- support for
  - coefficients in a prime field  $\mathbb{Z}/p\mathbb{Z}$
  - sparse distance matrices for distance threshold
- open source (<http://ripser.org>)
  - released in July 2016
- online version (<http://live.ripser.org>)
  - launched in August 2016
- most efficient software for Vietoris–Rips persistence by far
  - computes  $H^2$  barcode for 50000 random points on a torus in 136 seconds / 9 GB
- (co-)winner of 2016 ATMCS Best New Software Award

# The four special ingredients

The improved performance is based on 4 insights:

- Skip inessential columns
- Compute cohomology
- Implicit boundary matrix
- Apparent pairs

# Computing homology

Computing homology  $H_* = Z_*/B_*$  (recall:  $B_* \subseteq Z_* \subseteq C_*$ ):

- compute basis for boundaries  $B_* = \text{im } \partial_*$
- extend to basis for cycles  $Z_* = \ker \partial_*$
- new (non-boundary) basis cycles generate quotient  $Z_*/B_*$

# Homology by matrix reduction

Notation:

- $D$ : boundary matrix
- $R_i$ :  $i$ th column of  $R$

# Homology by matrix reduction

Notation:

- $D$ : boundary matrix
- $R_i$ :  $i$ th column of  $R$

Matrix reduction algorithm (variant of Gaussian elimination):

- $R = D, V = I$
- while  $\exists i < j$  with pivot  $R_i = \text{pivot } R_j$ 
  - add  $R_i$  to  $R_j$ , add  $V_i$  to  $V_j$

# Homology by matrix reduction

Notation:

- $D$ : boundary matrix
- $R_i$ :  $i$ th column of  $R$

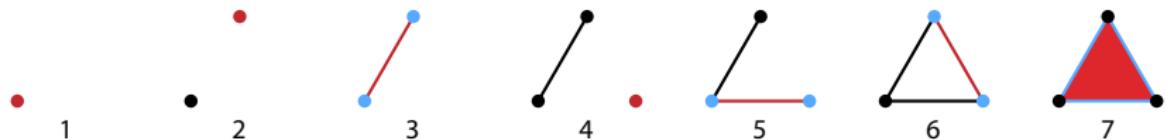
Matrix reduction algorithm (variant of Gaussian elimination):

- $R = D, V = I$
- while  $\exists i < j$  with pivot  $R_i = \text{pivot } R_j$ 
  - add  $R_i$  to  $R_j$ , add  $V_i$  to  $V_j$

Result:

- $R = D \cdot V$  is reduced (each column has a unique pivot)
- $V$  is full rank upper triangular

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1		
2			1			1	
3							1
4				1	1		
5							1
6							1
7							

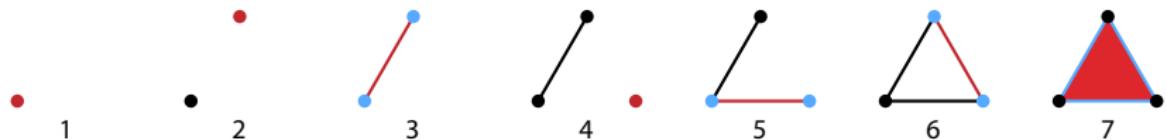
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1				
4				1			
5					1		
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1		
2				1		1	
3							1
4					1	1	
5							1
6							1
7							

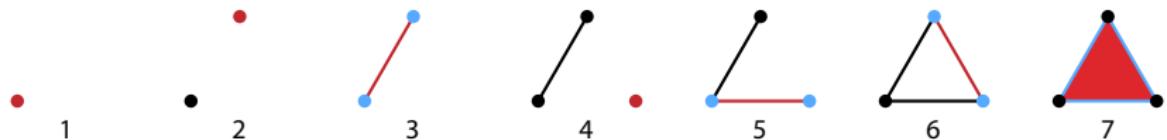
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1				
4				1			
5					1		
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1		
2			1			1	
3							1
4					1	1	
5							1
6							1
7							

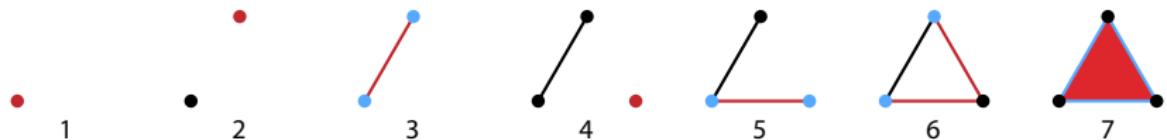
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1				
4				1			
5					1		
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1	1	
2			1			1	
3							1
4				1	0		
5						1	
6						1	
7							

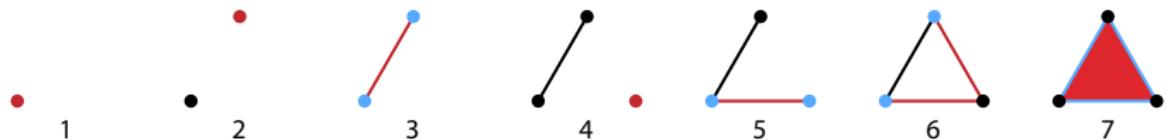
$\underbrace{\hspace{1cm}}$  *R*

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1				
4				1			
5					1	1	
6						1	
7							1

$\underbrace{\hspace{1cm}}$  *V*

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1	1	
2				1			1
3							1
4					1		
5							1
6							1
7							

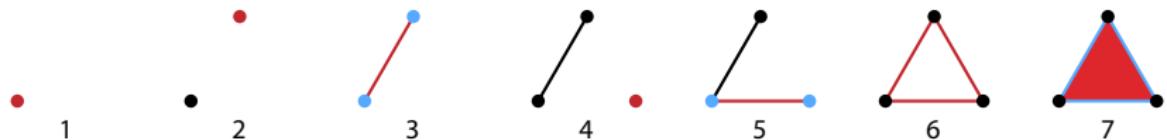
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1				
4				1			
5					1	1	
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



	1	2	3	4	5	6	7
1			1	1	1	0	
2			1			0	
3						1	
4				1			
5						1	
6						1	
7							

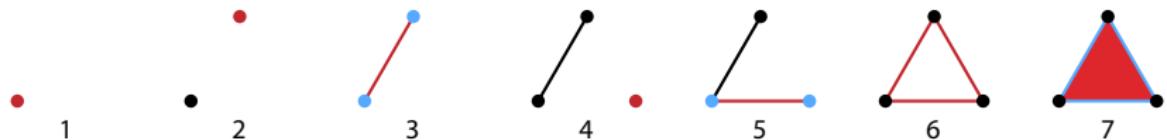
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1				1
4				1			
5					1	1	
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1		
2			1				
3						1	
4				1			
5						1	
6						1	
7							1

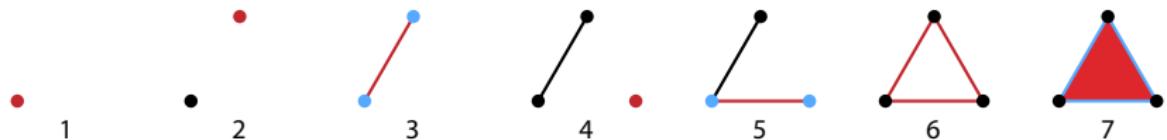
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1			1	
4				1			
5					1	1	
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



	1	2	3	4	5	6	7
1			1		1		
2			1				
3						1	
4					1		
5						1	
6						1	
7							

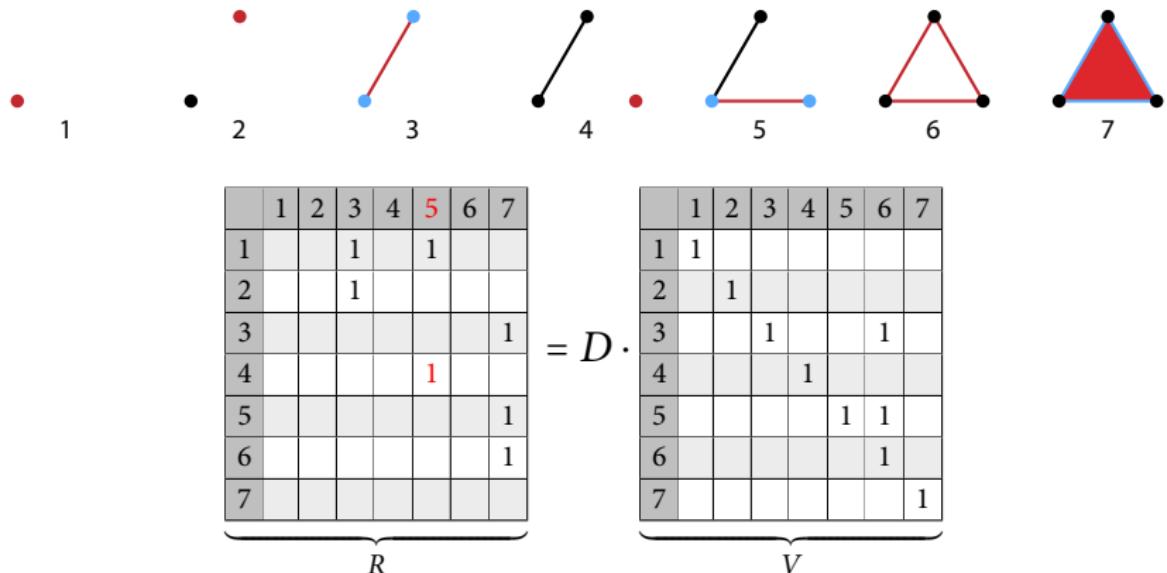
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

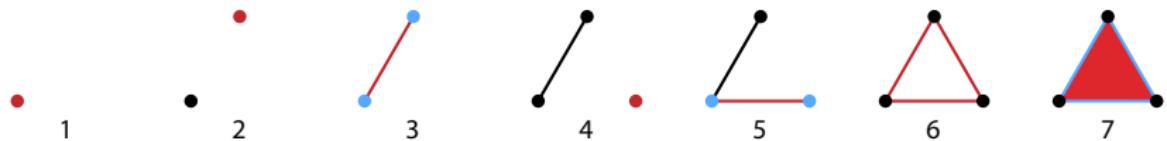
	1	2	3	4	5	6	7
1	1						
2		1					
3			1			1	
4				1			
5					1	1	
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



# Matrix reduction



	1	2	3	4	5	6	7
1			1		1		
2			1				
3						1	
4					1		
5						1	
6						1	
7							

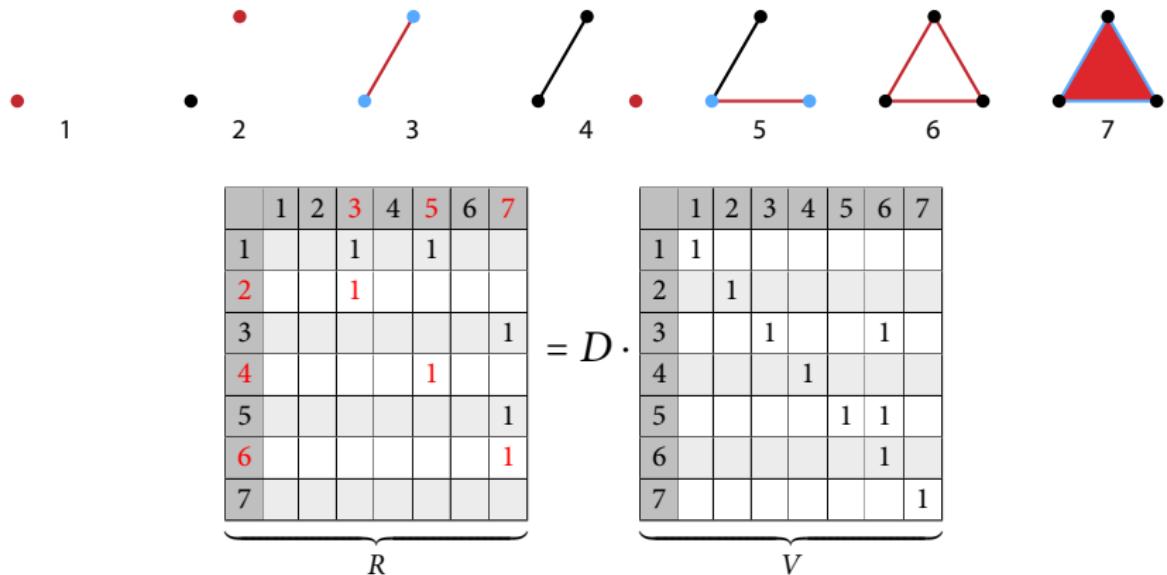
$\underbrace{\hspace{10em}}$   
R

$= D \cdot$

	1	2	3	4	5	6	7
1	1						
2		1					
3			1			1	
4				1			
5					1	1	
6						1	
7							1

$\underbrace{\hspace{10em}}$   
V

# Matrix reduction



## Skipping inessential columns

Assume that  $R = D \cdot V$  is reduced,  $V$  is full rank upper triangular.

Then

$\Sigma_B = \{R_j \mid R_j \neq 0\}$  is a basis for  $B_* = \text{im } D$  (boundaries),

## Skipping inessential columns

Assume that  $R = D \cdot V$  is reduced,  $V$  is full rank upper triangular.

Then

$\Sigma_B = \{R_j \mid R_j \neq 0\}$  is a basis for  $B_* = \text{im } D$  (boundaries),

$\Sigma_Z = \{V_i \mid R_i = 0\}$  is a basis for  $Z_* = \ker D$  (cycles).

## Skipping inessential columns

Assume that  $R = D \cdot V$  is reduced,  $V$  is full rank upper triangular.

Then

$\Sigma_B = \{R_j \mid R_j \neq 0\}$  is a basis for  $B_* = \text{im } D$  (boundaries),

$\Sigma_Z = \{V_i \mid R_i = 0\}$  is a basis for  $Z_* = \ker D$  (cycles).

Extend  $\Sigma_B$  to *another* basis for  $Z_*$  using cycles from  $\Sigma_Z$

(exchange cycle  $V_i \in \Sigma_Z$  with boundary  $R_j \in \Sigma_B$  if  $i = \text{pivot } R_j$ ):

## Skipping inessential columns

Assume that  $R = D \cdot V$  is reduced,  $V$  is full rank upper triangular.

Then

$\Sigma_B = \{R_j \mid R_j \neq 0\}$  is a basis for  $B_* = \text{im } D$  (boundaries),

$\Sigma_Z = \{V_i \mid R_i = 0\}$  is a basis for  $Z_* = \ker D$  (cycles).

Extend  $\Sigma_B$  to *another* basis for  $Z_*$  using cycles from  $\Sigma_Z$

(exchange cycle  $V_i \in \Sigma_Z$  with boundary  $R_j \in \Sigma_B$  if  $i = \text{pivot } R_j$ ):

$$\widetilde{\Sigma}_Z = \Sigma_B \cup \underbrace{\{V_i \mid R_i = 0, i \neq \text{pivot } R_j \text{ for all } j\}}_{\Sigma_E}$$

## Skipping inessential columns

Assume that  $R = D \cdot V$  is reduced,  $V$  is full rank upper triangular.  
Then

$$\begin{aligned}\Sigma_B &= \{R_j \mid R_j \neq 0\} \quad \text{is a basis for } B_* = \text{im } D \text{ (boundaries),} \\ \Sigma_Z &= \{V_i \mid R_i = 0\} \quad \text{is a basis for } Z_* = \ker D \text{ (cycles).}\end{aligned}$$

Extend  $\Sigma_B$  to *another* basis for  $Z_*$  using cycles from  $\Sigma_Z$   
(exchange cycle  $V_i \in \Sigma_Z$  with boundary  $R_j \in \Sigma_B$  if  $i = \text{pivot } R_j$ ):

$$\widetilde{\Sigma}_Z = \Sigma_B \cup \underbrace{\{V_i \mid R_i = 0, i \neq \text{pivot } R_j \text{ for all } j\}}_{\Sigma_E}$$

Homology  $Z_*/B_*$  is generated by the extra basis cycles  $\Sigma_E$

## Skipping inessential columns

Assume that  $R = D \cdot V$  is reduced,  $V$  is full rank upper triangular.  
Then

$$\begin{aligned}\Sigma_B &= \{R_j \mid R_j \neq 0\} \quad \text{is a basis for } B_* = \text{im } D \text{ (boundaries),} \\ \Sigma_Z &= \{V_i \mid R_i = 0\} \quad \text{is a basis for } Z_* = \ker D \text{ (cycles).}\end{aligned}$$

Extend  $\Sigma_B$  to *another* basis for  $Z_*$  using cycles from  $\Sigma_Z$   
(exchange cycle  $V_i \in \Sigma_Z$  with boundary  $R_j \in \Sigma_B$  if  $i = \text{pivot } R_j$ ):

$$\widetilde{\Sigma}_Z = \Sigma_B \cup \underbrace{\{V_i \mid R_i = 0, i \neq \text{pivot } R_j \text{ for all } j\}}_{\Sigma_E}$$

Homology  $Z_*/B_*$  is generated by the extra basis cycles  $\Sigma_E$

- The columns with indices  $i = \text{pivot } R_j$  are never used!

# Persistent cohomology

We have seen: many columns of  $R = D \cdot V$  are not needed

- Skip those inessential columns in matrix reduction

# Persistent cohomology

We have seen: many columns of  $R = D \cdot V$  are not needed

- Skip those inessential columns in matrix reduction

For persistence barcodes in low dimensions  $d < k$ :

- Number of skipped indices for reducing  $D^T$  (cohomology) is much larger than for  $D$  (homology)
  - reducing boundary matrix produces basis for  $H_k(K_k)$ , which is not needed
- The resulting persistence barcode is the same  
[de Silva et al. 2011]

## Implicit boundary matrix

Typical approach:

- Boundary matrix  $D$  with respect to filtration-ordered basis
- Explicitly generated and stored in memory

## Implicit boundary matrix

Typical approach:

- Boundary matrix  $D$  with respect to filtration-ordered basis
- Explicitly generated and stored in memory

Approach for Ripser: *thinking is faster than reading*

## Implicit boundary matrix

Typical approach:

- Boundary matrix  $D$  with respect to filtration-ordered basis
- Explicitly generated and stored in memory

Approach for Ripser: *thinking is faster than reading*

- Boundary matrix  $D$  with respect to fixed ordered basis (lexicographic)
- Implicitly defined by number of vertices, dimension
- Rows/columns of  $D$  recomputed instead of stored

# Implicit boundary matrix

Typical approach:

- Boundary matrix  $D$  with respect to filtration-ordered basis
- Explicitly generated and stored in memory

Approach for Ripser: *thinking is faster than reading*

- Boundary matrix  $D$  with respect to fixed ordered basis (lexicographic)
- Implicitly defined by number of vertices, dimension
- Rows/columns of  $D$  recomputed instead of stored

Matrix reduction in Ripser: store only coefficient matrix  $V$

- recompute columns of  $R = D \cdot V$  from  $D, V$ 
  - Typically,  $V$  is much sparser and smaller than  $R$  (fewer  $k$ -simplices than  $(k+1)$ -simplices)

# Apparent pairs

## Apparent pairs

- provide a close connection between persistence and discrete Morse theory
- are both persistence pairs and Morse pairs
- typically cover almost all simplices in the Rips filtration
- provide a shortcut for computation

# Natural filtration settings

Typical assumptions on the filtration:

- general filtration persistence (in theory)
- filtration by singletons or pairs discrete Morse theory
- simplexwise filtration persistence (computation)

# Natural filtration settings

Typical assumptions on the filtration:

- general filtration persistence (in theory)
- filtration by singletons or pairs discrete Morse theory
- simplexwise filtration persistence (computation)

Conclusion:

- Discrete Morse theory sits in the middle between persistence and persistence (hah!)

# Morse pairs and persistence pairs

Consider a *Morse filtration* (one or two simplices at a time).

*Morse pair*  $(\sigma, \tau)$ :

- inserting  $\sigma$  and  $\tau$  simultaneously does not change the *homotopy type*

# Morse pairs and persistence pairs

Consider a *Morse filtration* (one or two simplices at a time).

*Morse pair*  $(\sigma, \tau)$ :

- inserting  $\sigma$  and  $\tau$  simultaneously does not change the *homotopy type*

Consider a *simplexwise filtration* (one simplex at a time).

*Persistence pair*  $(\sigma, \tau)$ :

- inserting simplex  $\sigma$  creates a new *homological* feature
- inserting  $\tau$  destroys that feature again

# From Morse theory to persistence and back

## Proposition (from Morse to persistence)

*The pairs of a Morse filtration are apparent 0-persistence pairs for the simplexwise refinement of the filtration.*

*Apparent persistence pair  $(\sigma, \tau)$ :*

- $\sigma$  is the youngest face of  $\tau$
- $\tau$  is the oldest coface of  $\sigma$

# From Morse theory to persistence and back

## Proposition (from Morse to persistence)

*The pairs of a Morse filtration are apparent 0-persistence pairs for the simplexwise refinement of the filtration.*

*Apparent persistence pair  $(\sigma, \tau)$ :*

- $\sigma$  is the youngest face of  $\tau$
- $\tau$  is the oldest coface of  $\sigma$

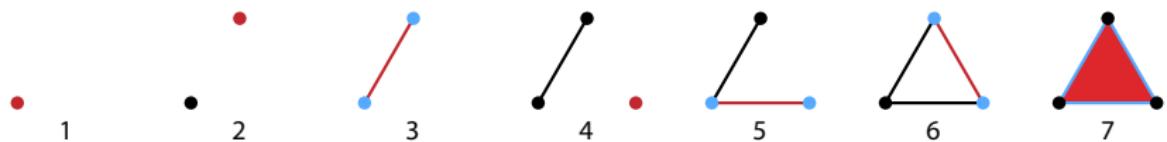
## Proposition (from persistence to Morse)

*Consider an arbitrary filtration with a simplexwise refinement.*

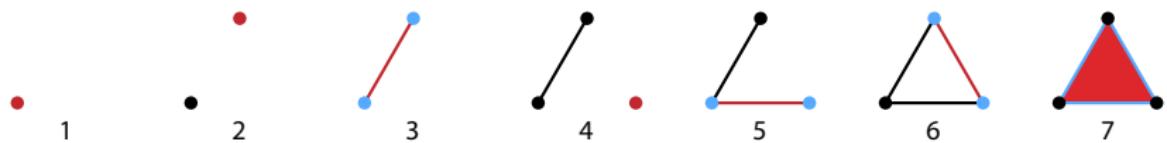
*The apparent 0-persistence pairs yield a Morse filtration*

- *refining the original one, and*
- *coarsening the simplexwise one.*

## Example: apparent pairs



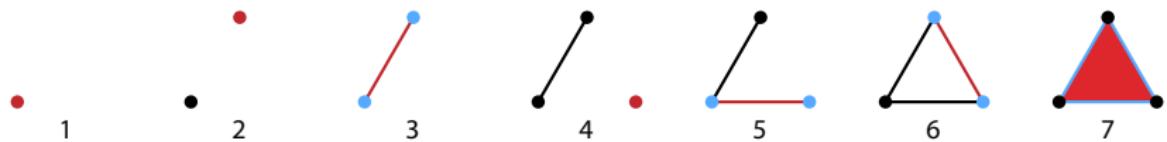
## Example: apparent pairs



$D =$

	1	2	3	4	5	6	7
1			1		1		
2			1			1	
3							1
4					1	1	
5							1
6							1
7							

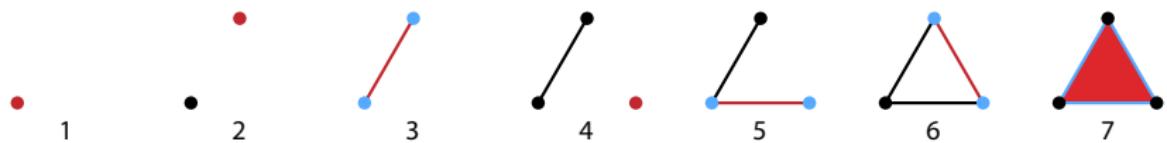
## Example: apparent pairs



$D =$

	1	2	3	4	5	6	7
1			1		1		
2			1			1	
3							1
4					1	1	
5							1
6							1
7							

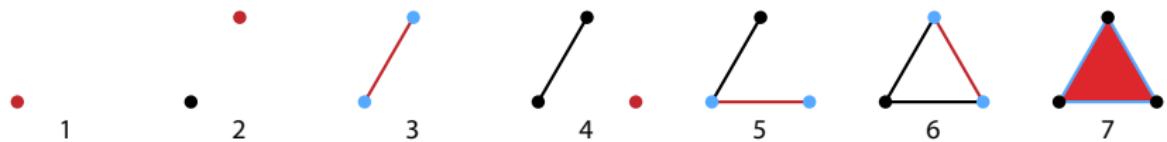
## Example: apparent pairs



$D =$

	1	2	3	4	5	6	7
1			1		1		
2			1			1	
3							1
4					1	1	
5							1
6							1
7							

## Example: apparent pairs



$D =$

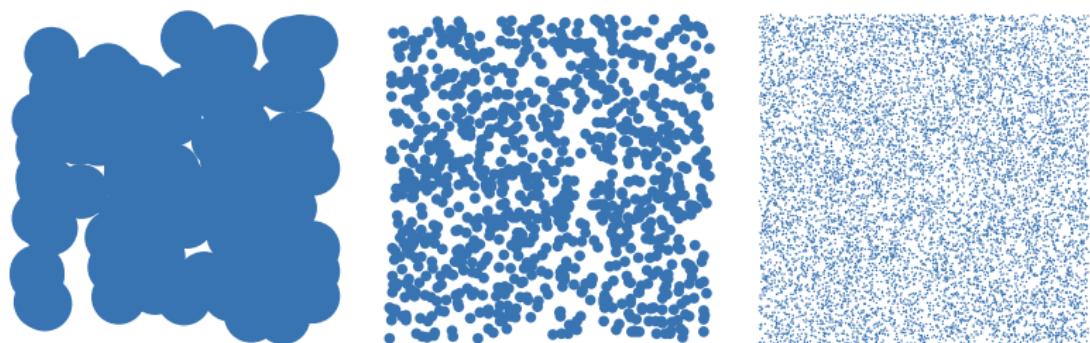
	1	2	3	4	5	6	7
1			1		1		
2			1			1	
3							1
4					1	1	
5							1
6							1
7							

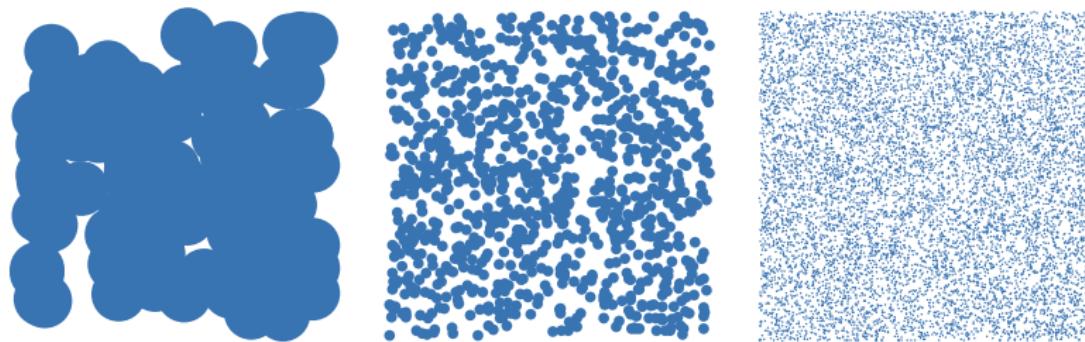
# Discrete Morse functions from total orders

The upshot:

- Total order on simplices induces discrete Morse function
- Simple and important example: lexicographic order, given by total order on vertices
- Often, this discrete Morse function has good properties (few critical points)
- Special cases have appeared in literature before
  - random Rips complex in supercritical regime [Kahle 2010]

# Stochastics



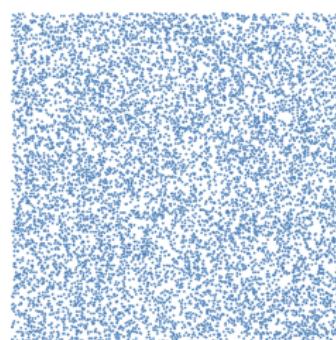
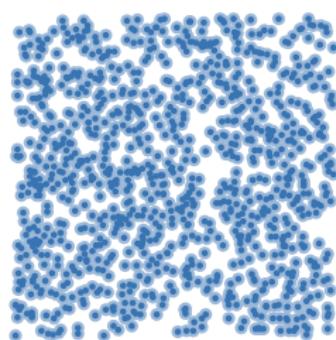
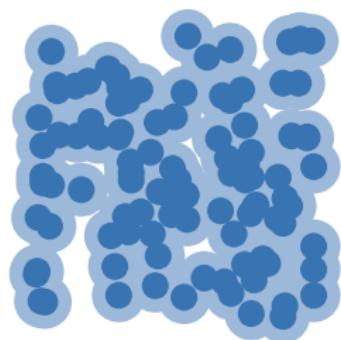


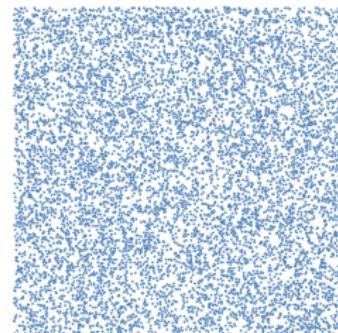
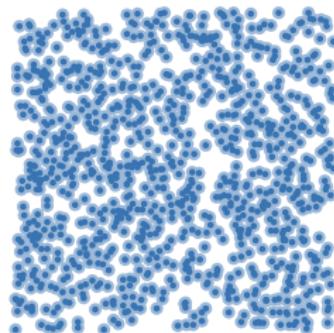
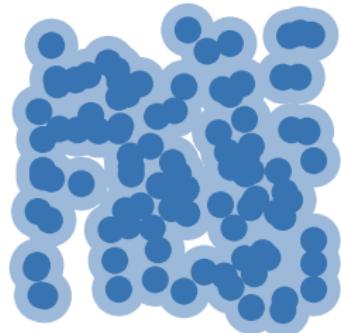
## Theorem (Kahle 2011)

*The expectation of the  $k$ th Betti number of a random Čech complex  $\text{Cech}_r(\mathcal{P}_n)$  satisfies*

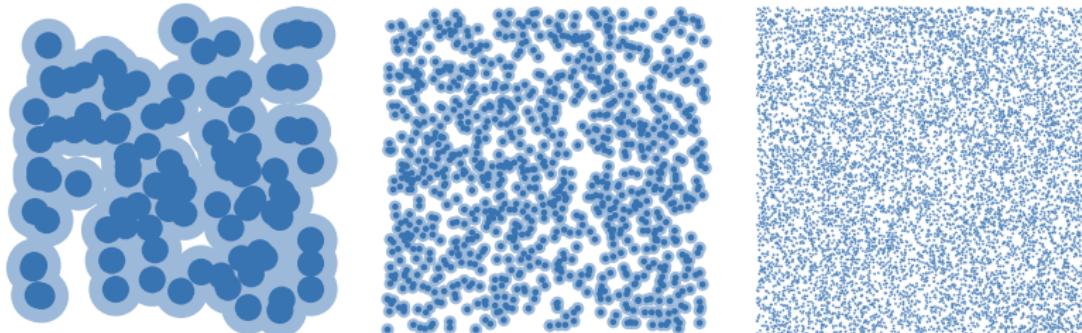
$$E(\beta_k(\mathcal{P}_n, r_n)) \in \Theta\left(n(r_n^d n)^{k+1}\right)$$

*in the subcritical regime  $r_n^d n \rightarrow 0$  as  $n \rightarrow \infty$ .*





Let  $m(\theta, k)$  be the smallest number of points required to form a persistent cycle with death/birth ratio  $\geq \theta$ .



Let  $m(\theta, k)$  be the smallest number of points required to form a persistent cycle with death/birth ratio  $\geq \theta$ .

### Theorem (B, Pausinger 2016)

*The expectation of the  $k$ th  $\theta$ -persistent Betti number of random Čech complexes  $\text{Cech}_r(\mathcal{P}_n) \subseteq \text{Cech}_{\theta r}(\mathcal{P}_n)$  satisfies*

$$E(\beta_k(\mathcal{P}_n, r_n)) \in \Theta(n(r_n^d n)^{m(\theta, k)-1})$$

*in the subcritical regime  $r_n^d n \rightarrow 0$  as  $n \rightarrow \infty$ .*

# Numerics

# Cauchy's differentiation formula

Differentiation by integration:

- Let  $f : U \subseteq \mathbb{C} \rightarrow \mathbb{C}$  be holomorphic,
- $D \subseteq U$  a closed disk, and
- $a \in \text{int } D$ .

Then

$$f^{(n)}(a) = \frac{n!}{2\pi i} \oint_{\gamma} \frac{f(z)}{(z-a)^{n+1}} dz$$

for any cycle  $\gamma$  in the homology class  $[\partial D] \in H_1(U \setminus \{a\})$ .

# Cauchy's differentiation formula

Differentiation by integration:

- Let  $f : U \subseteq \mathbb{C} \rightarrow \mathbb{C}$  be holomorphic,
- $D \subseteq U$  a closed disk, and
- $a \in \text{int } D$ .

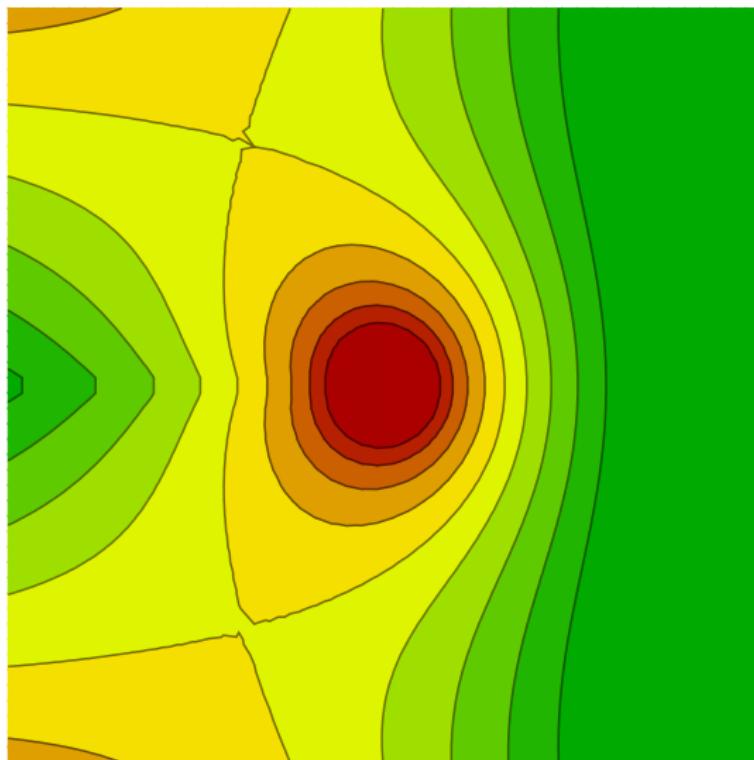
Then

$$f^{(n)}(a) = \frac{n!}{2\pi i} \oint_{\gamma} \frac{f(z)}{(z-a)^{n+1}} dz$$

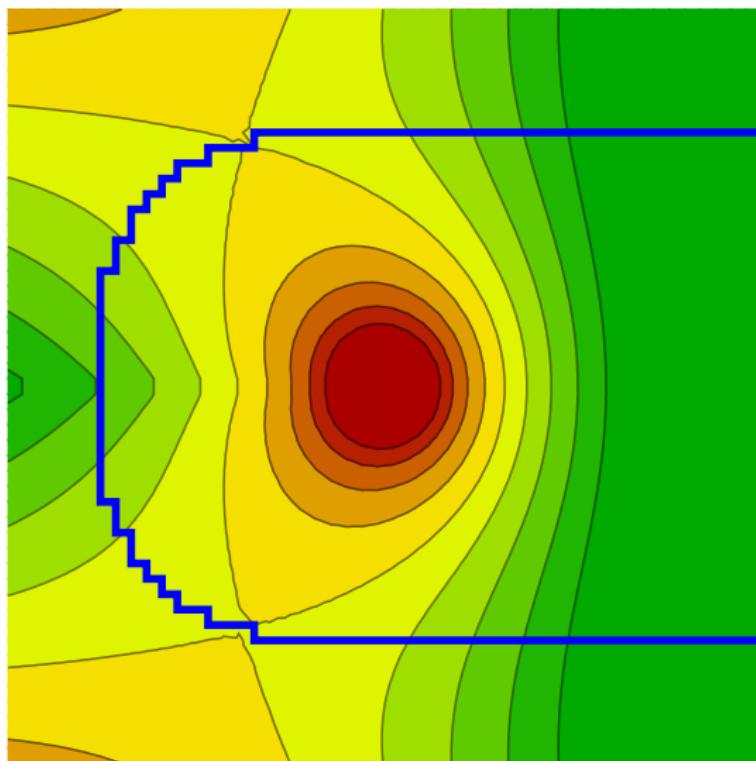
for any cycle  $\gamma$  in the homology class  $[\partial D] \in H_1(U \setminus \{a\})$ .

- But not all cycles are created equal  
(for numerical integration)!

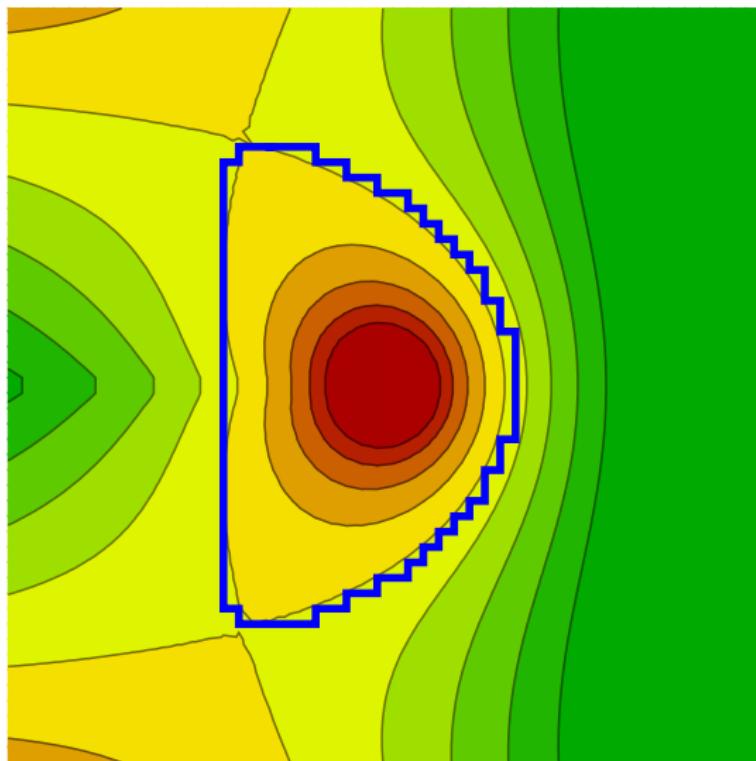
# Optimizing condition of contour integration



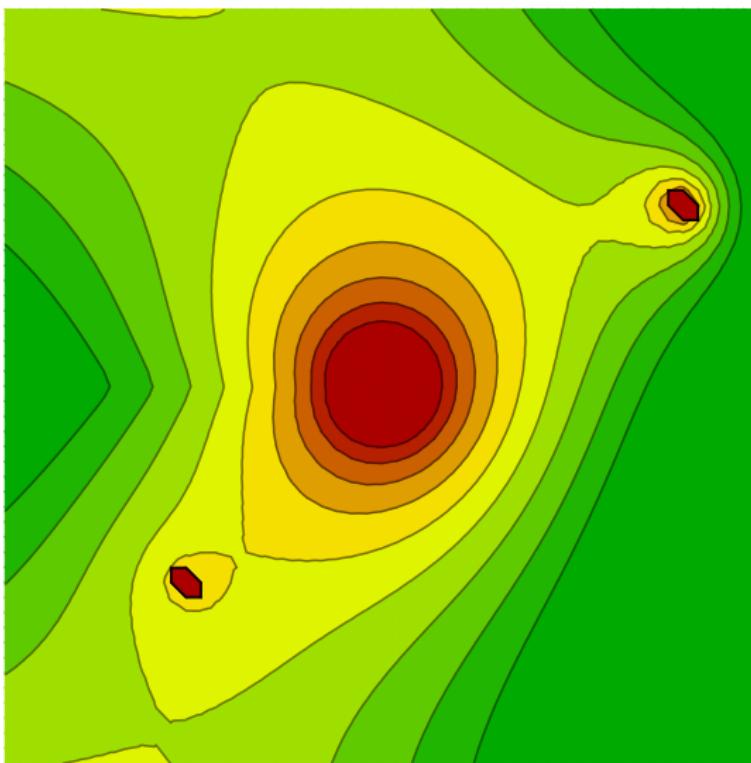
# Optimizing condition of contour integration



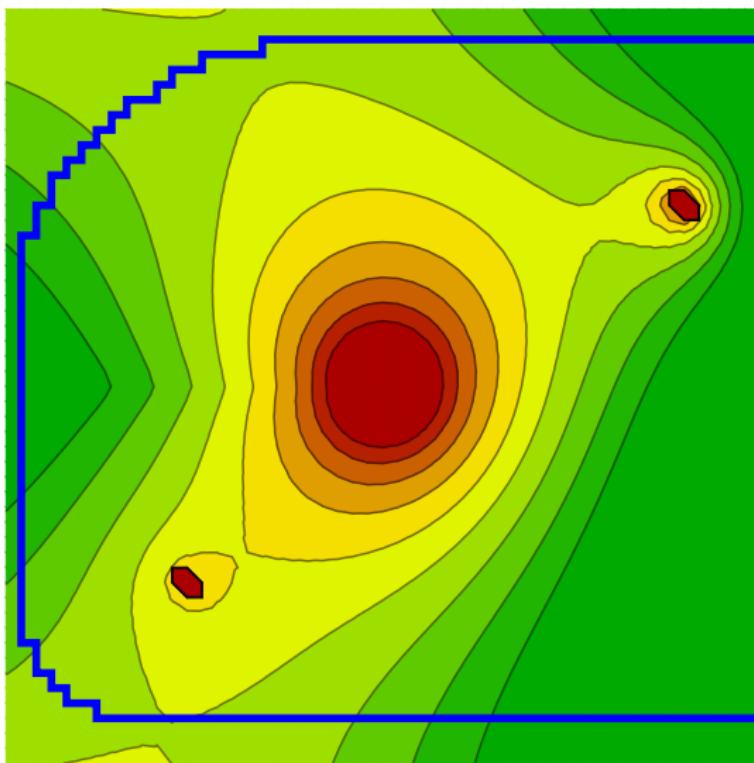
# Optimizing condition of contour integration



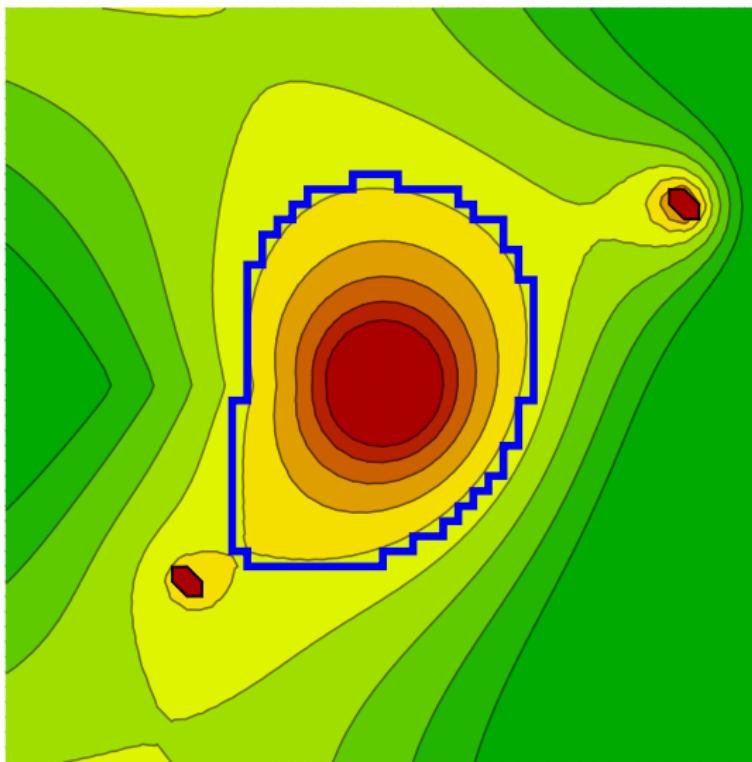
# Optimizing condition of contour integration



# Optimizing condition of contour integration



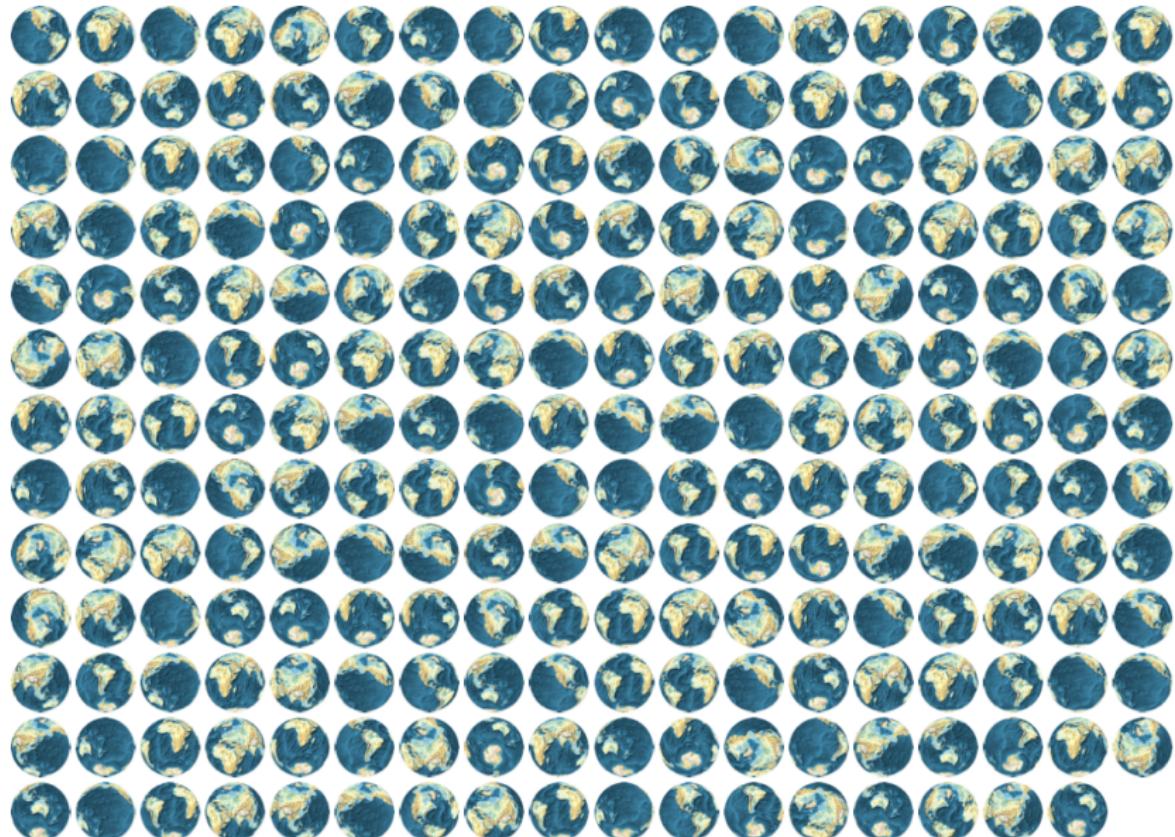
# Optimizing condition of contour integration

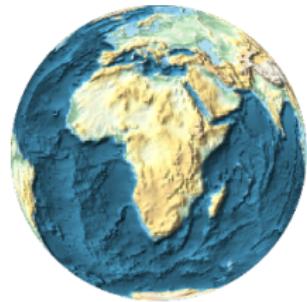


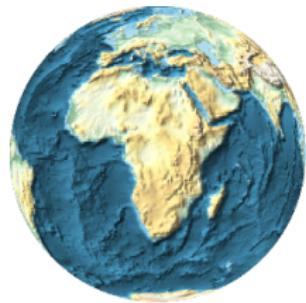
# Around the world

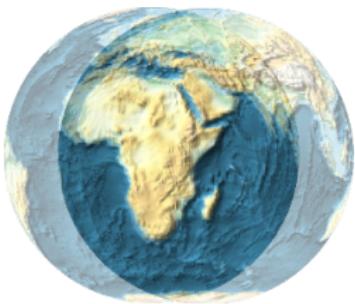
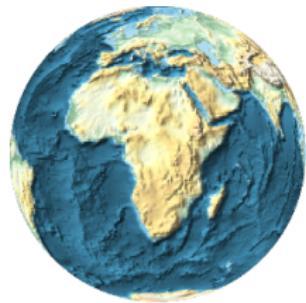


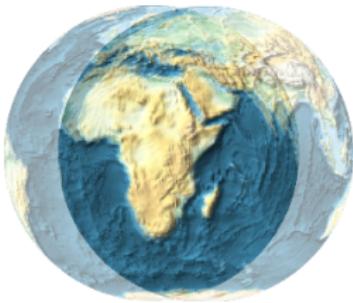
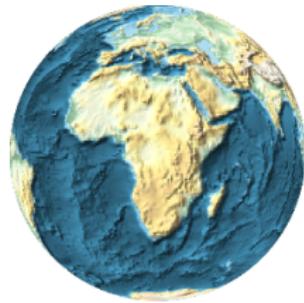




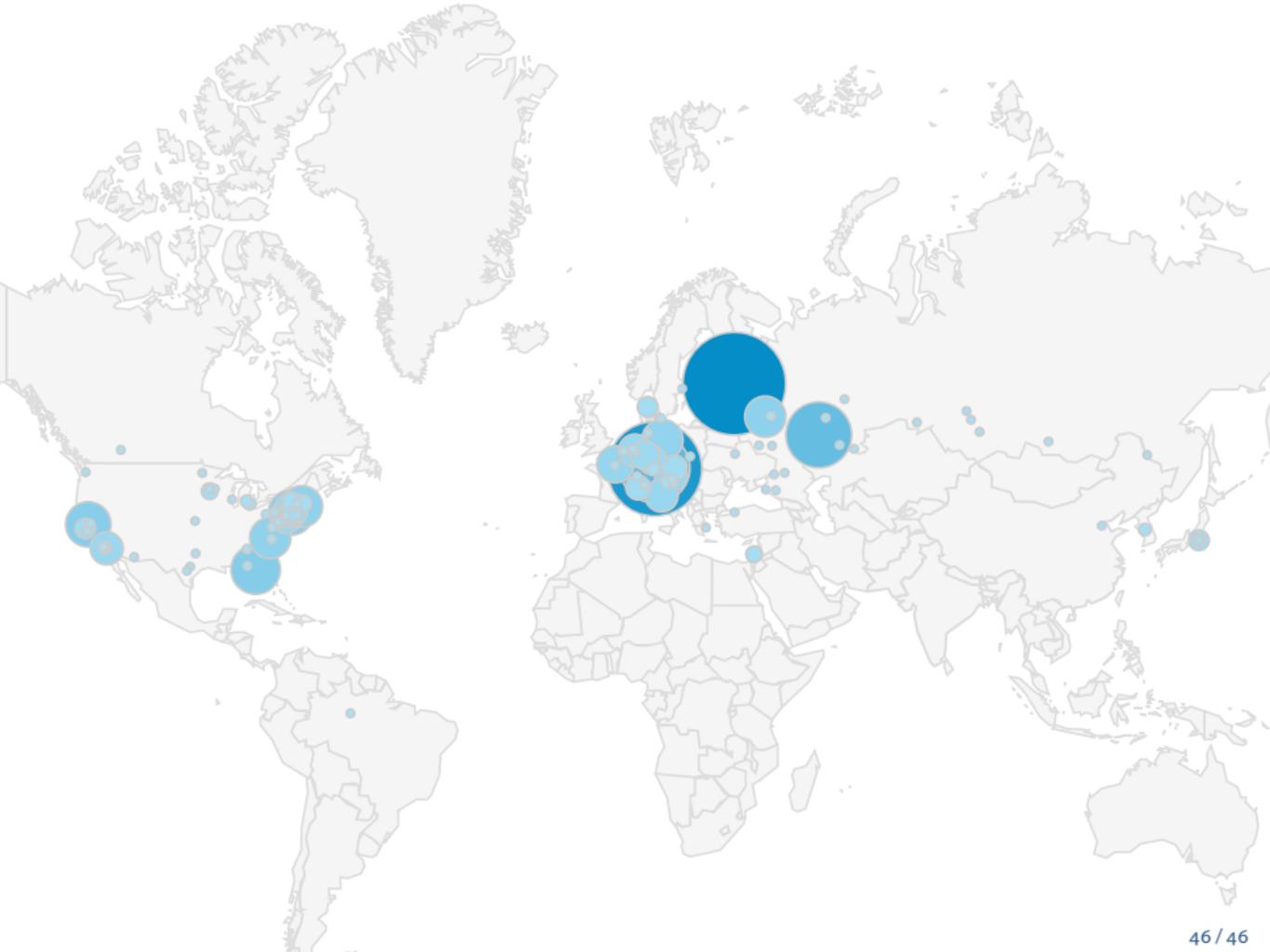








Demo (Ripser)



# Users of Ripser Live

