

Topological Data Analysis

Part I: Persistent homology

Ulrich Bauer

TUM

February 4, 2015



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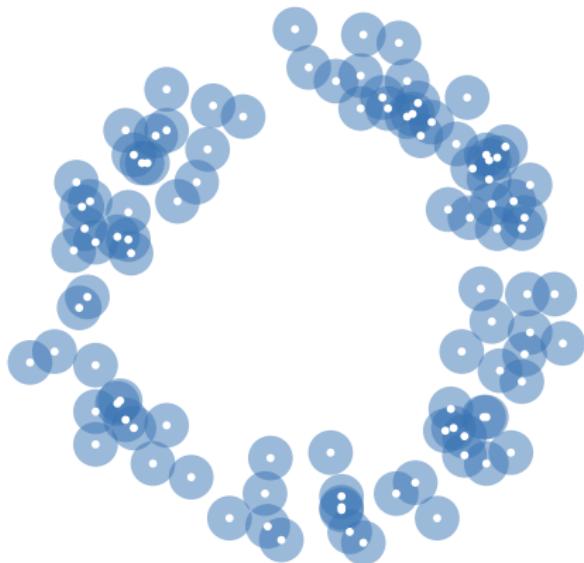
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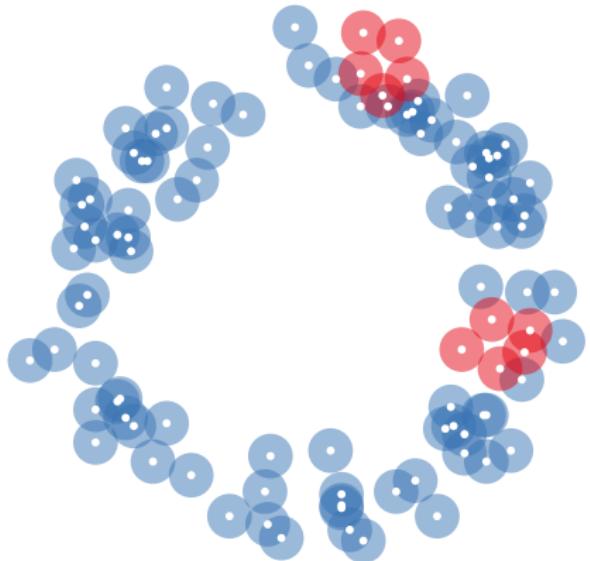


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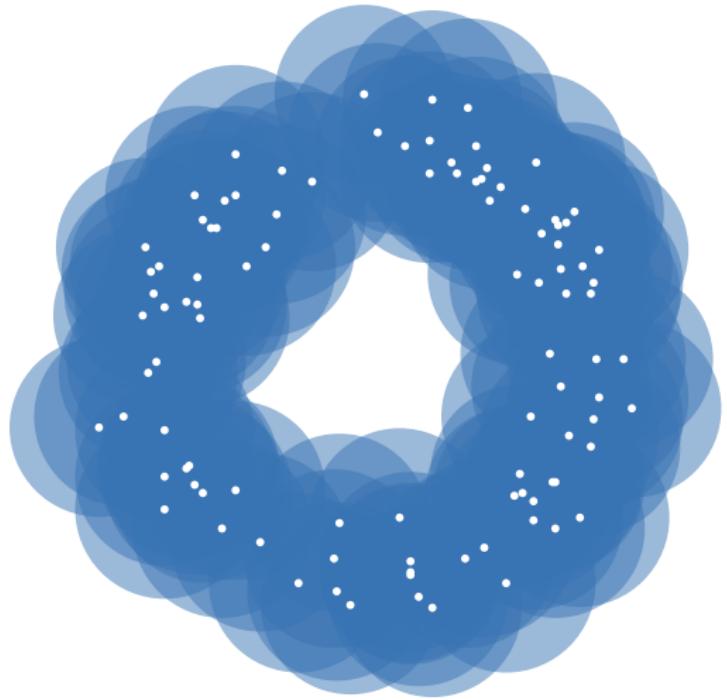
Persistent homology

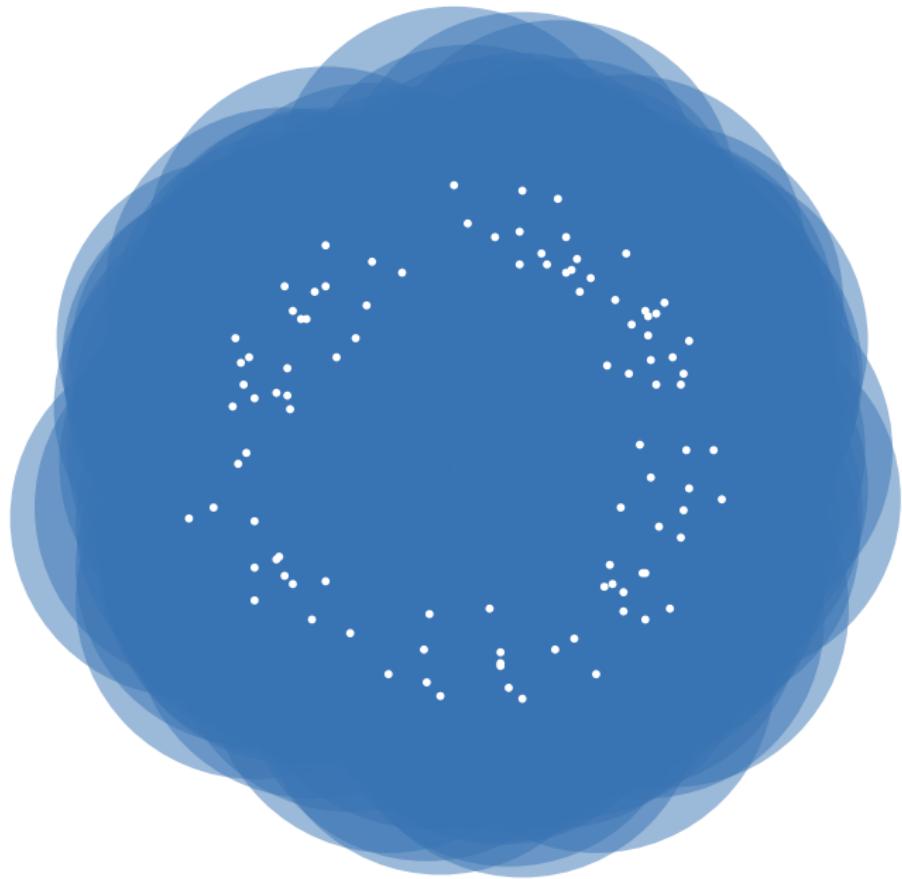


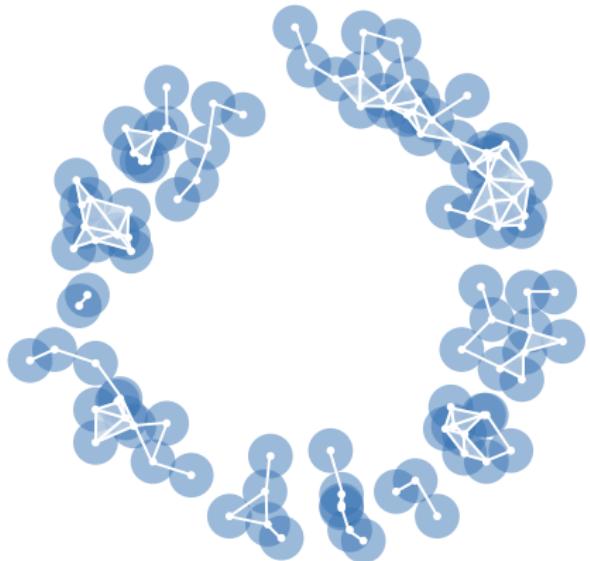


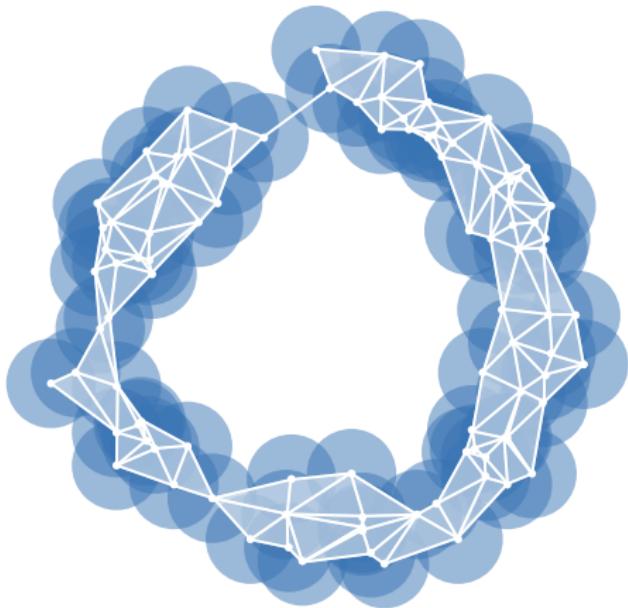


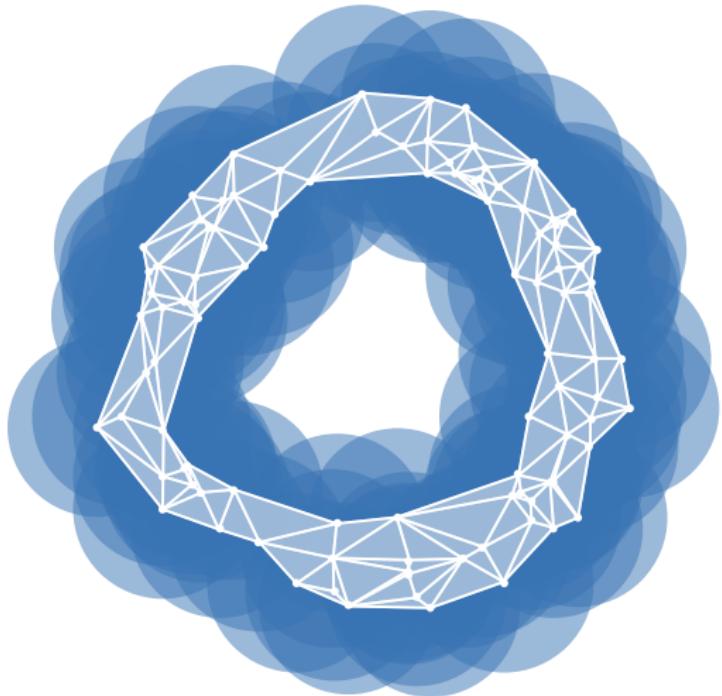


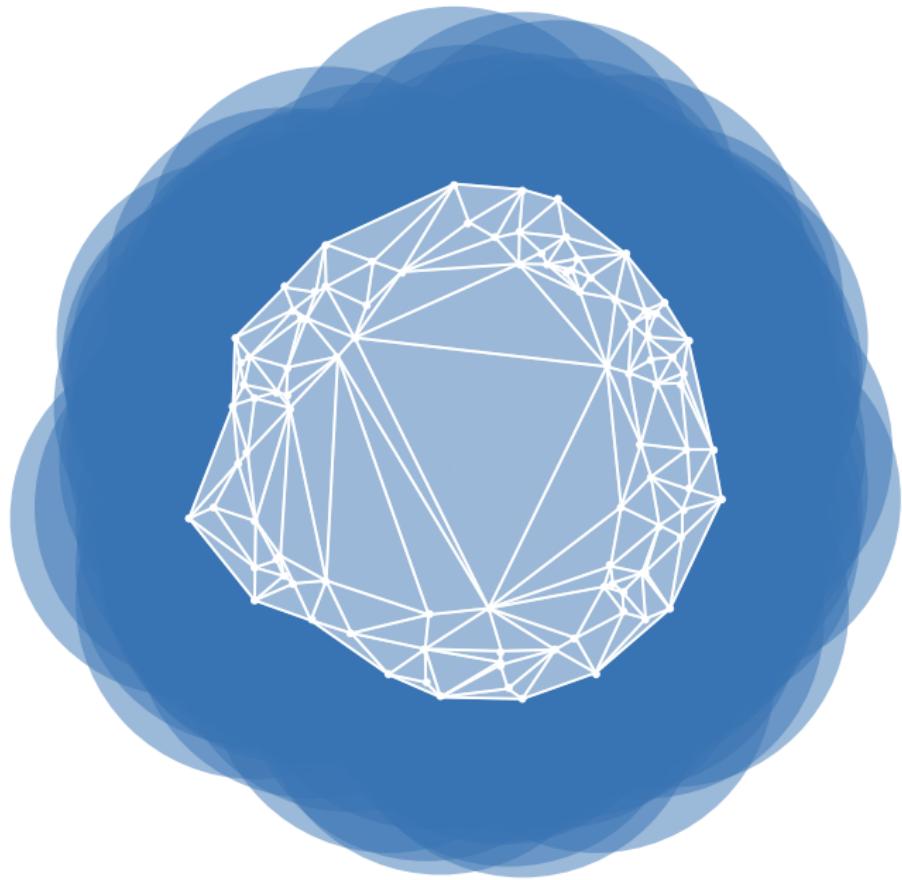




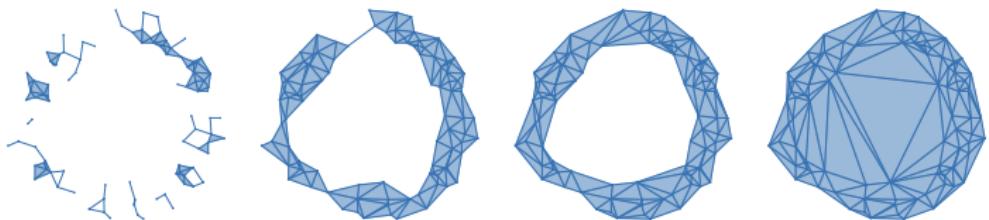




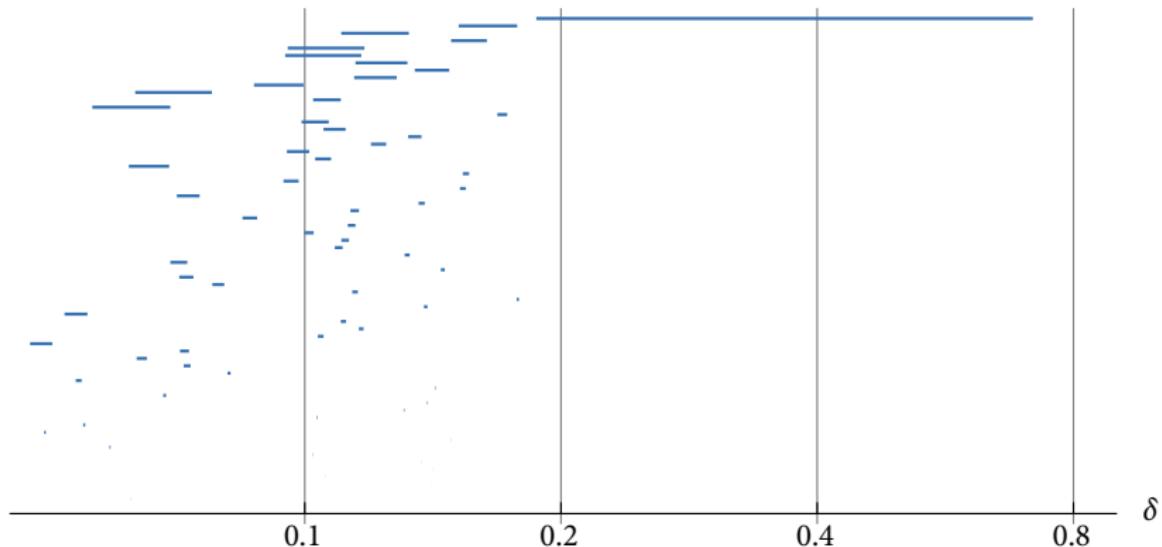
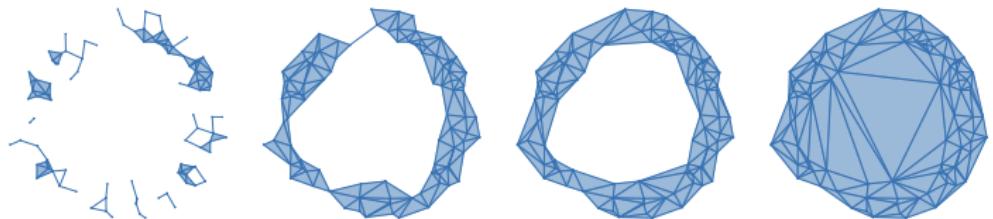




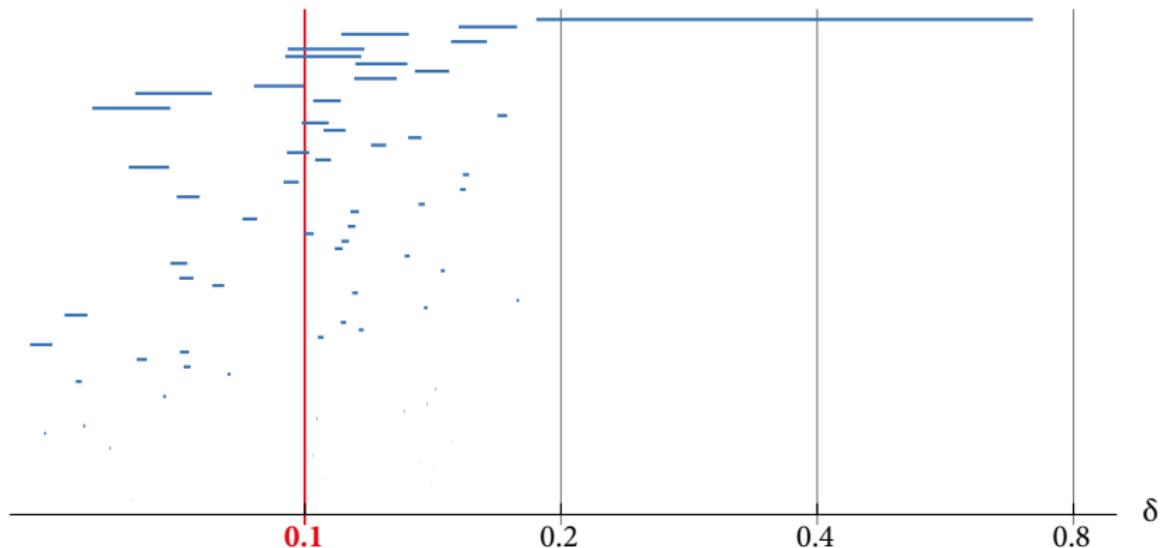
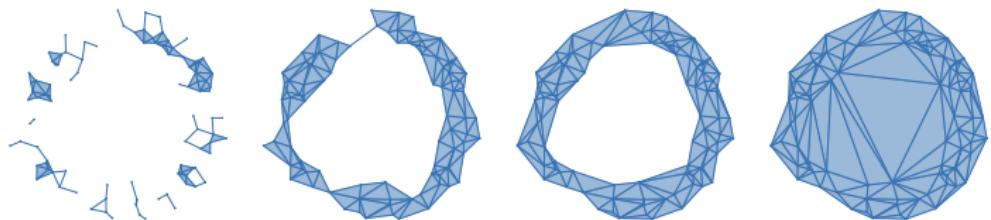
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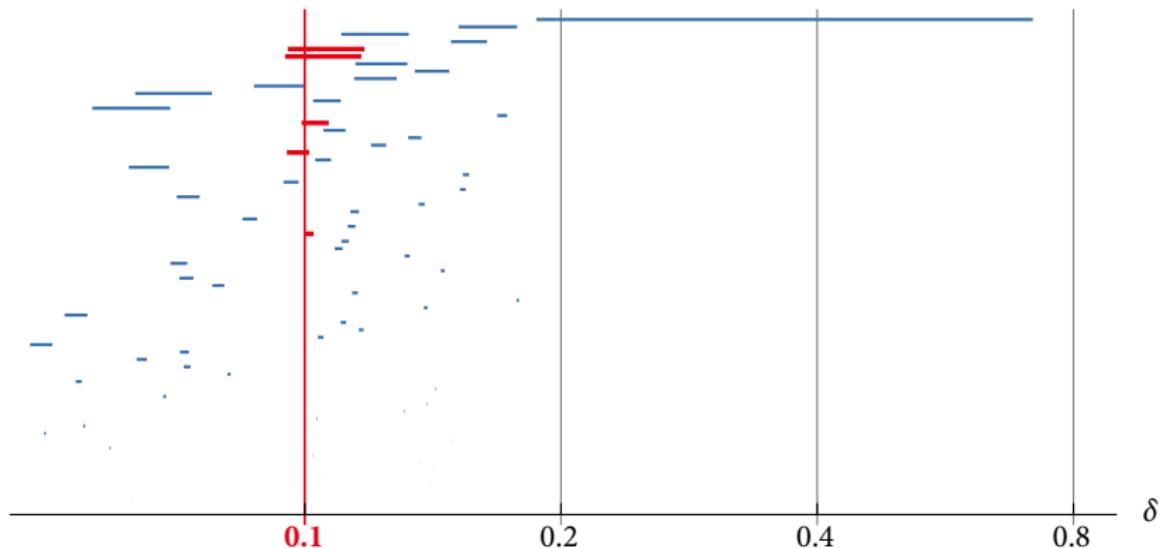
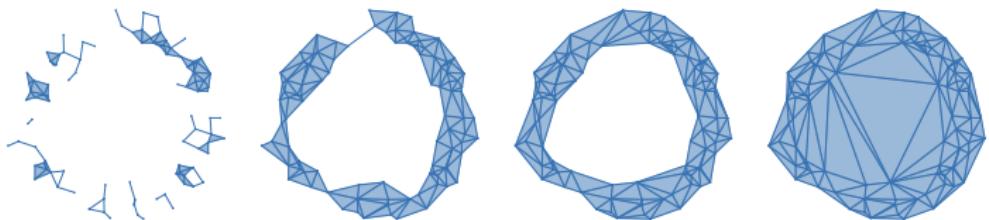
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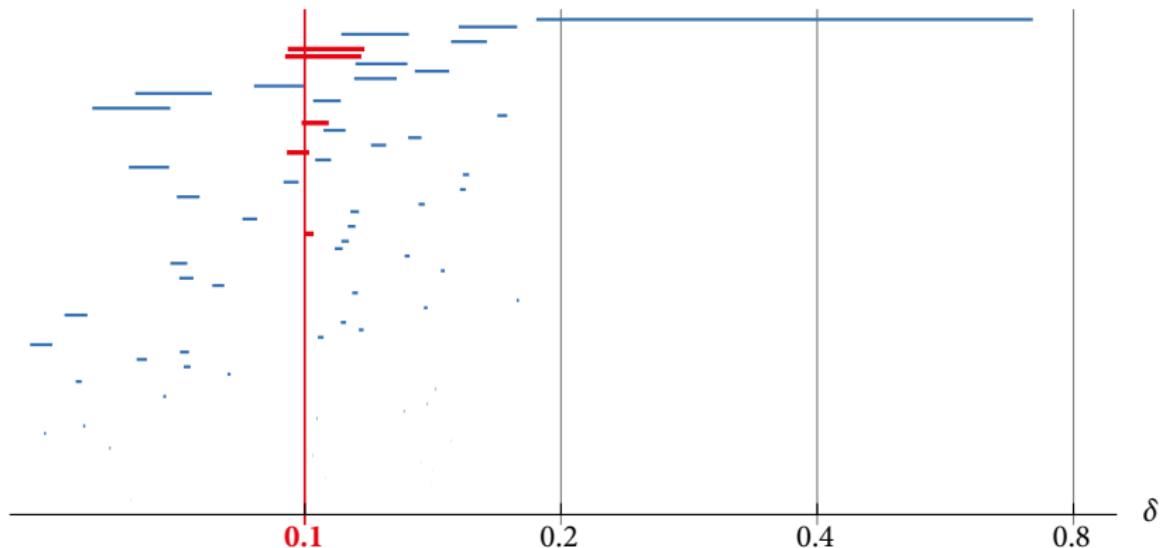
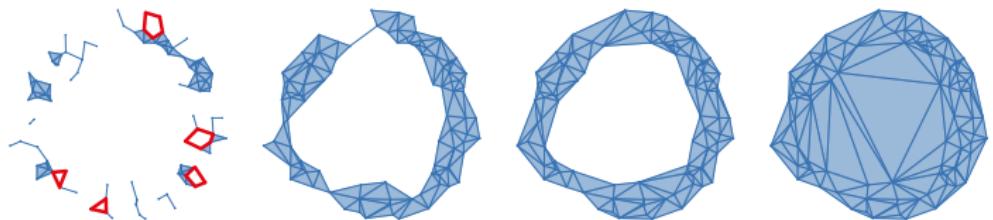
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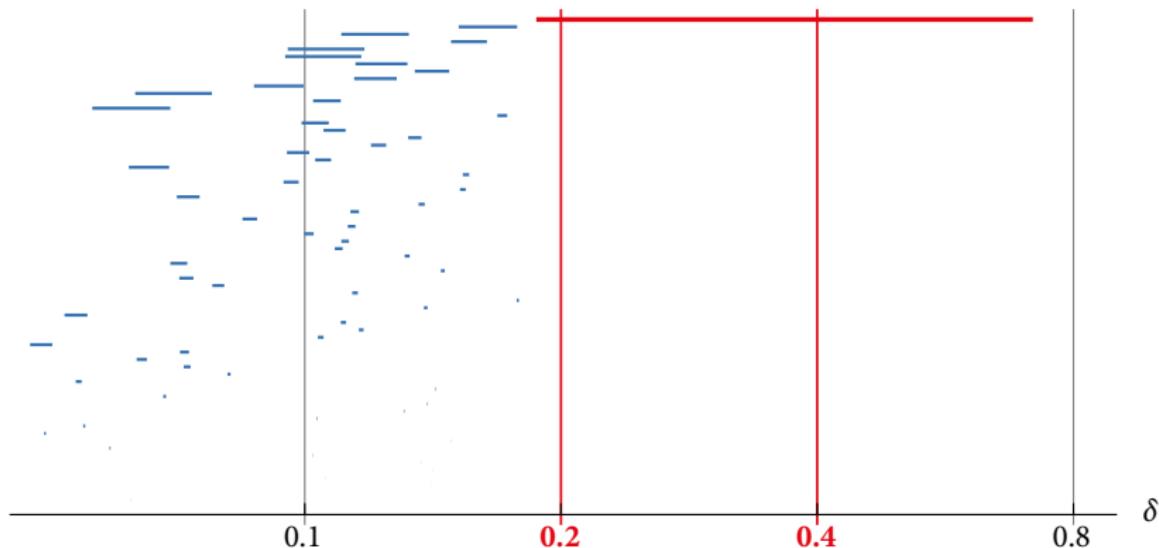
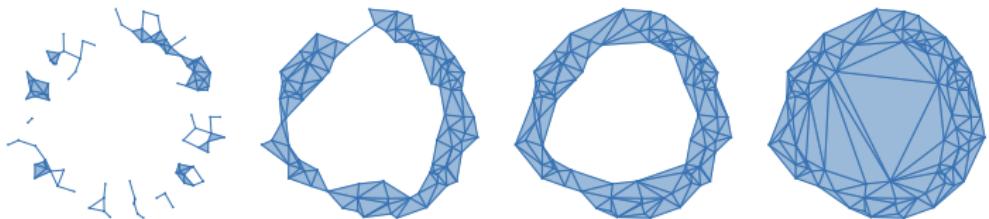
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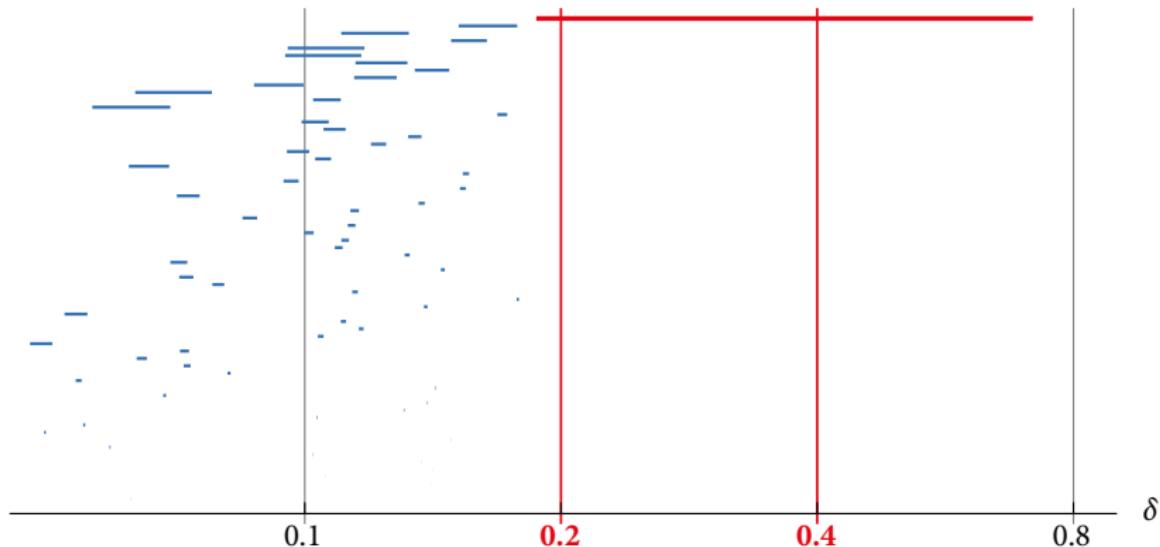
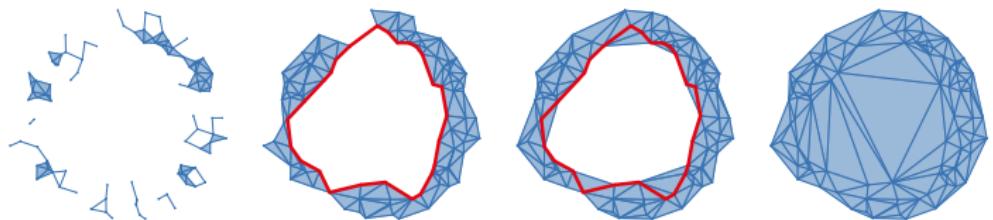
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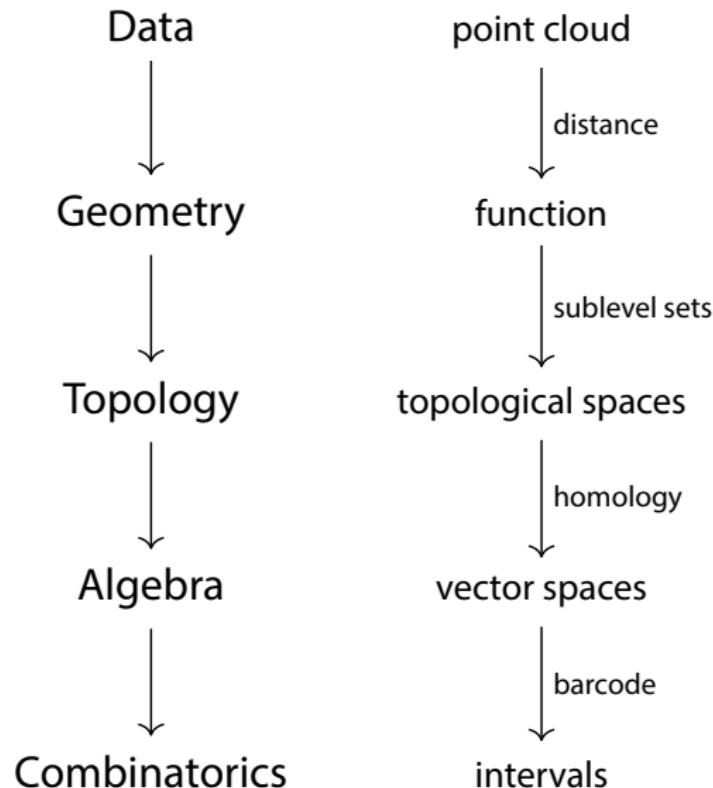
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- ▶ \mathbf{R} is the poset category of (\mathbb{R}, \leq)

The pipeline of topological data analysis



Simplification & Reconstruction

Homology inference

Problem (Homology inference)

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It is sometimes possible to recover the homology of Ω this way, but the assumptions are quite strong:

Homology reconstruction using union of balls

Theorem (Niyogi, Smale, Weinberger 2006)

Let M be a submanifold of \mathbb{R}^d . Let $P \subset M$ be such that $M \subseteq P^\delta$ for some $\delta < \sqrt{3/20} \text{reach}(M)$. Then

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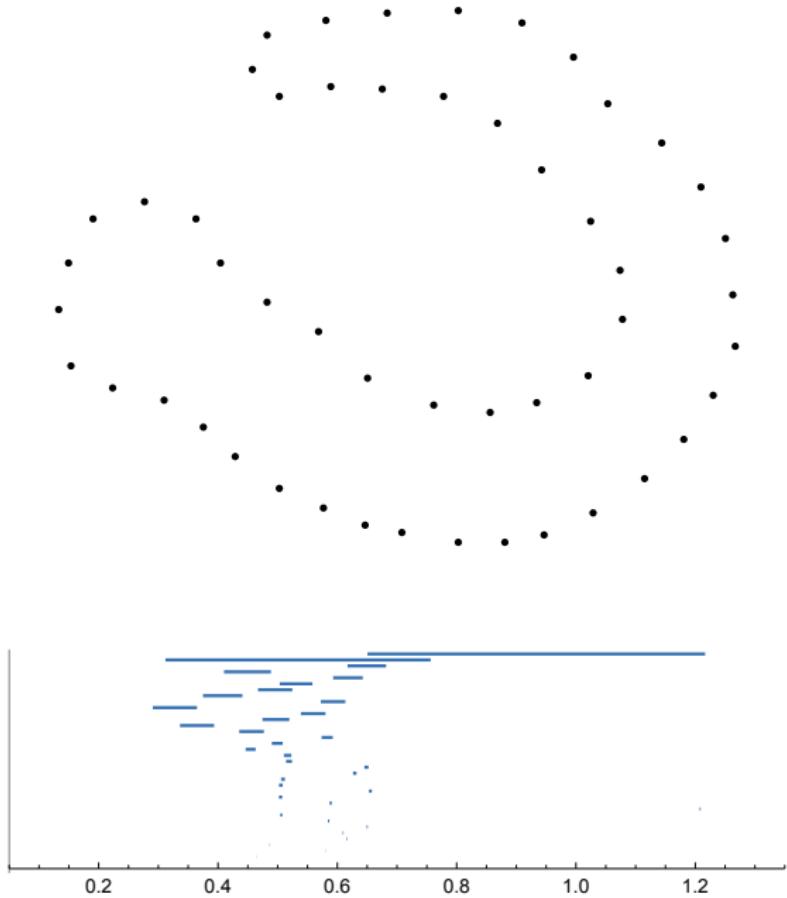
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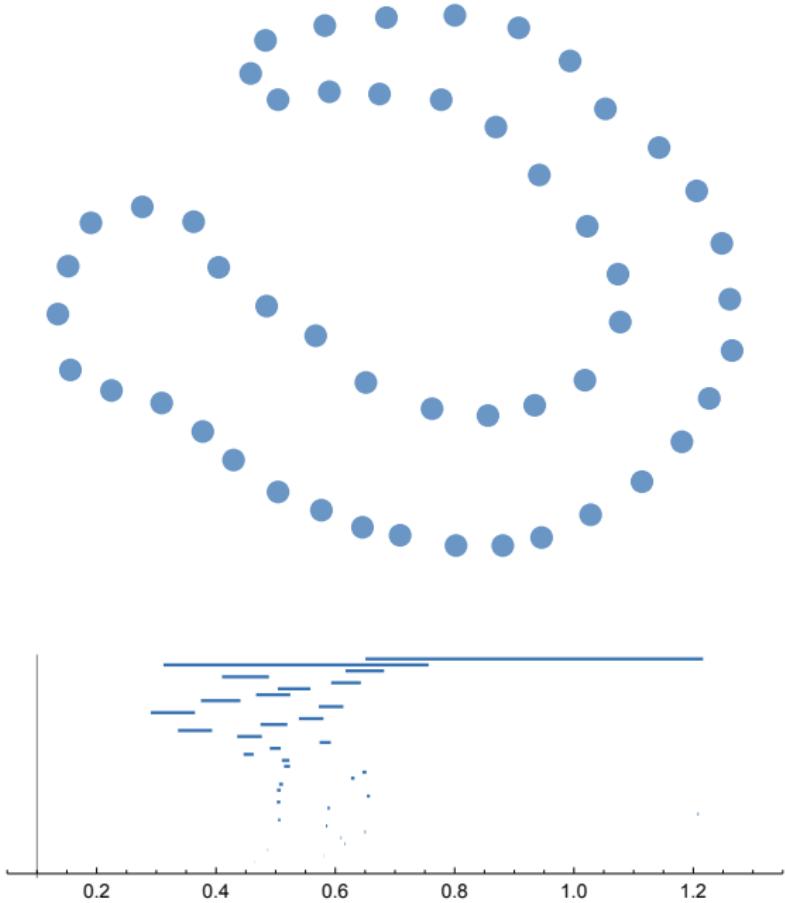
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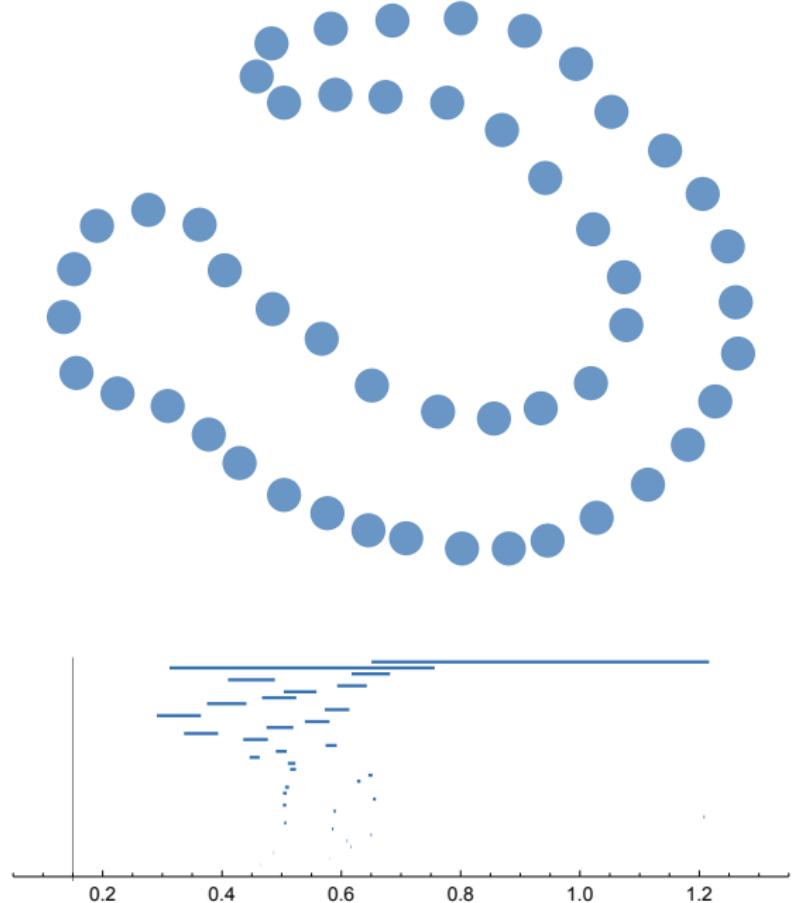
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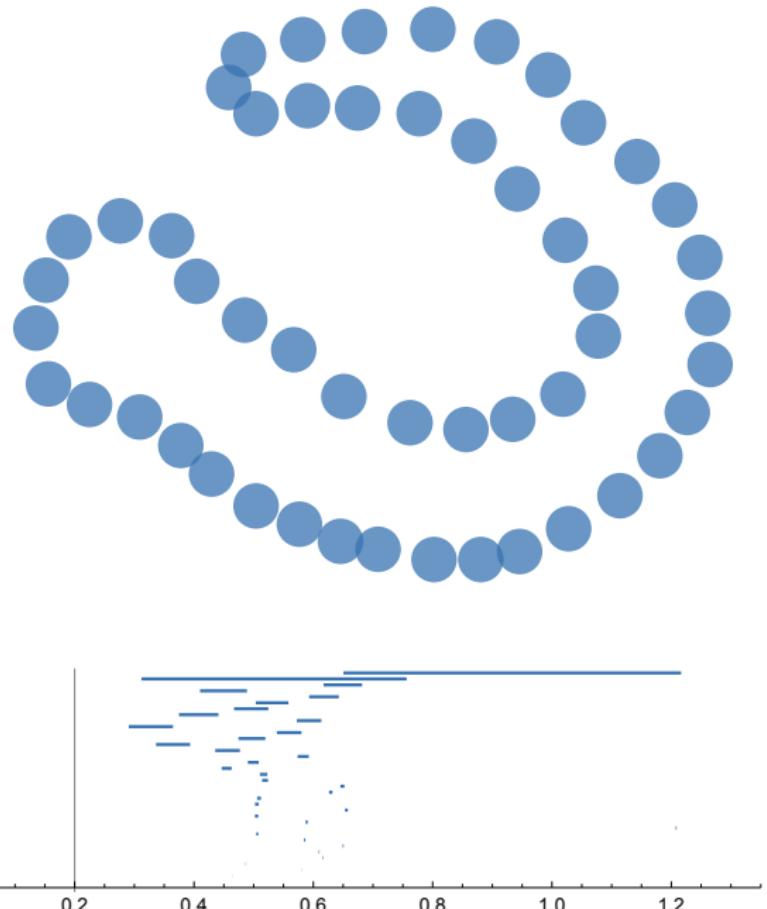
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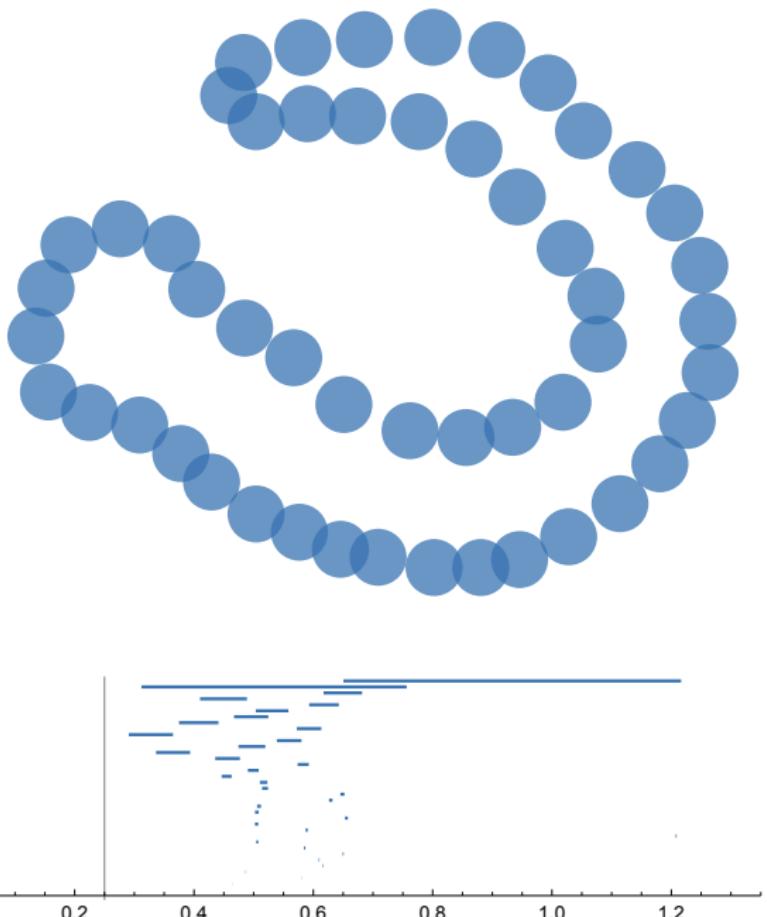
- ▶ $P_\delta = B_\delta(P)$: δ -neighborhood (union of balls) around P .
- ▶ Points with distance $< \text{reach}(M)$ to M have a unique closest point on M
- ▶ The isomorphism is induced by the inclusion $M \hookrightarrow P^{2\delta}$.

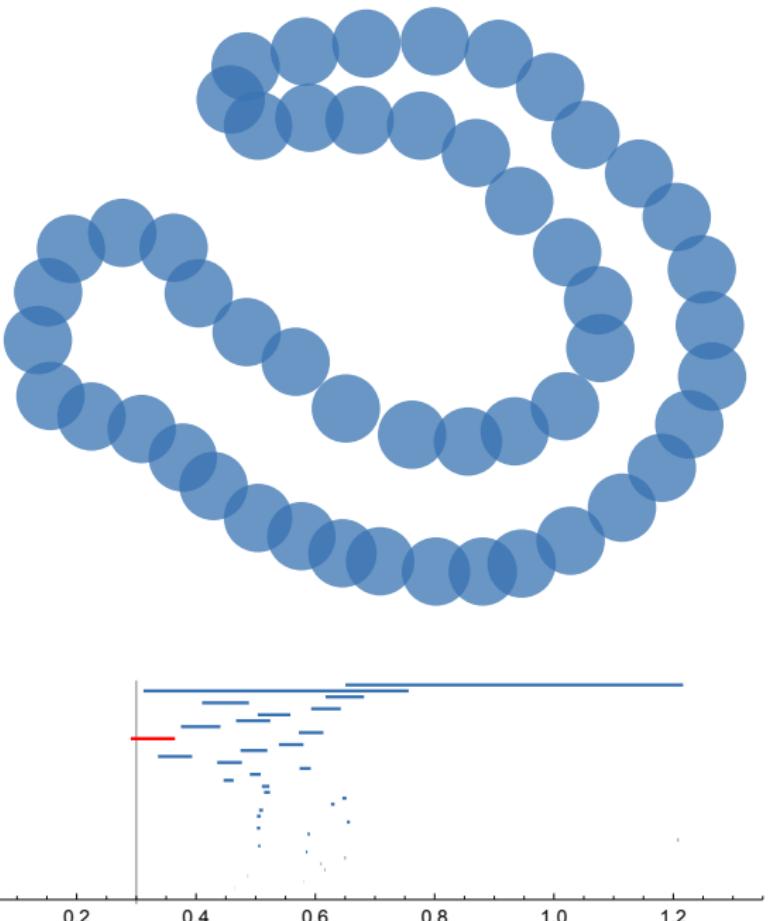


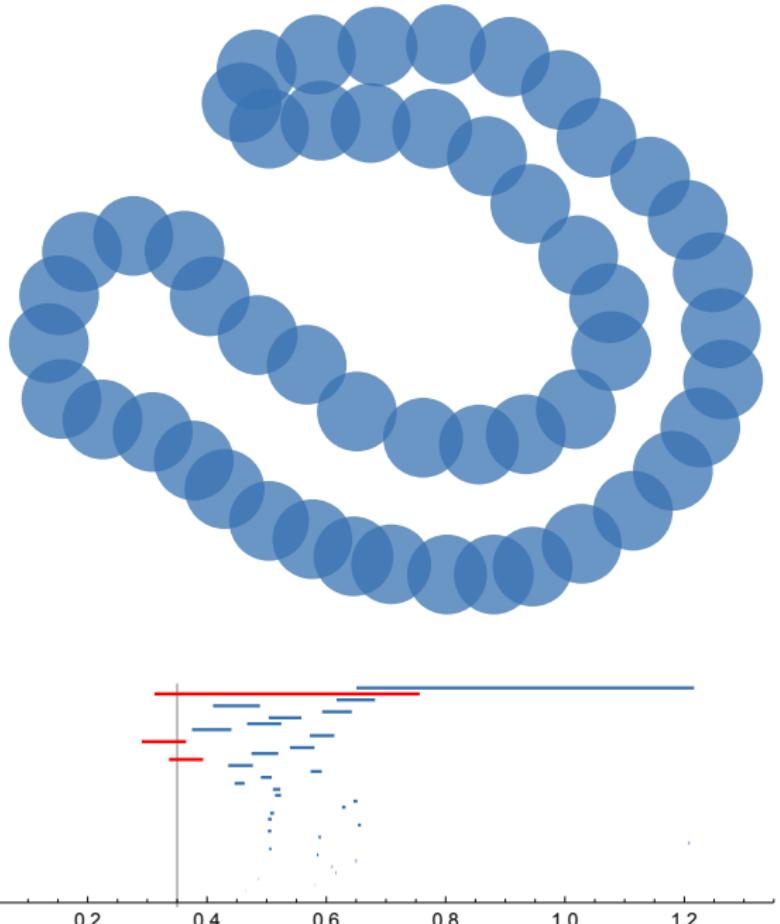


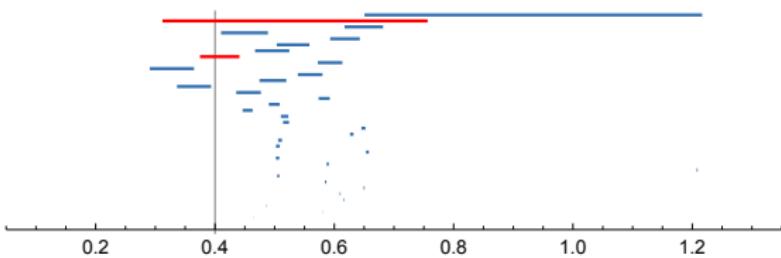
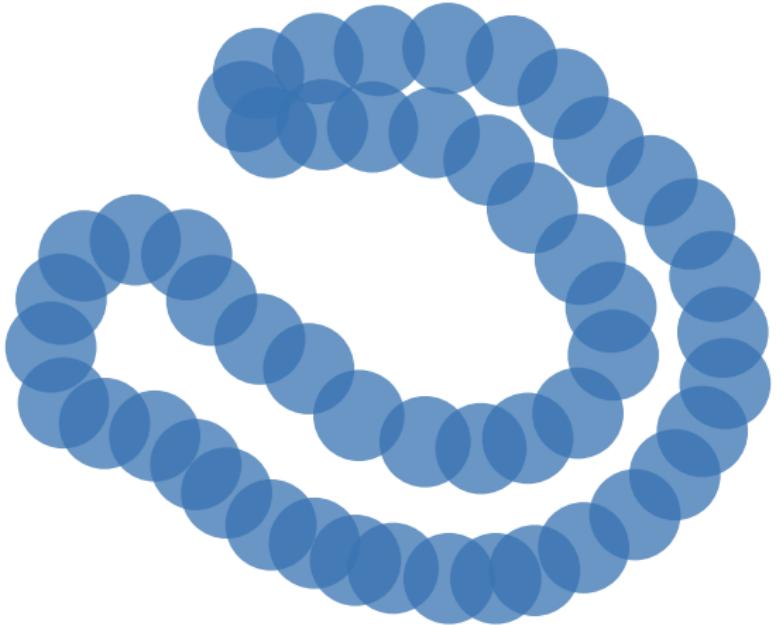


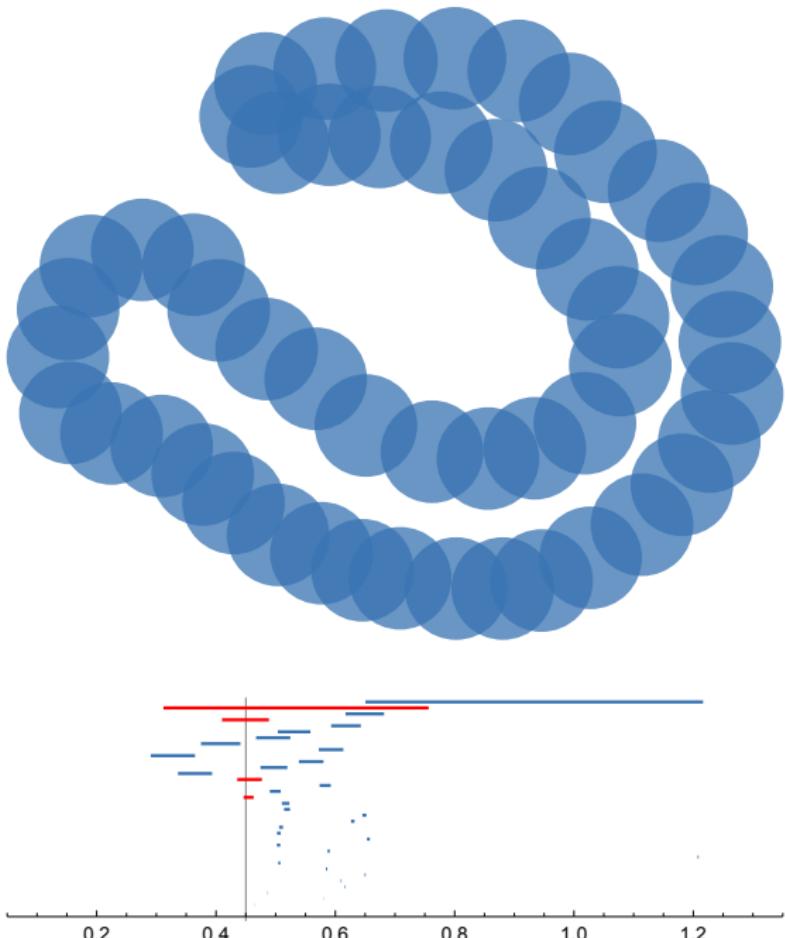


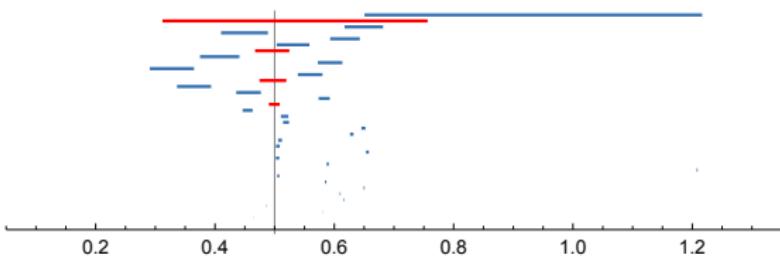
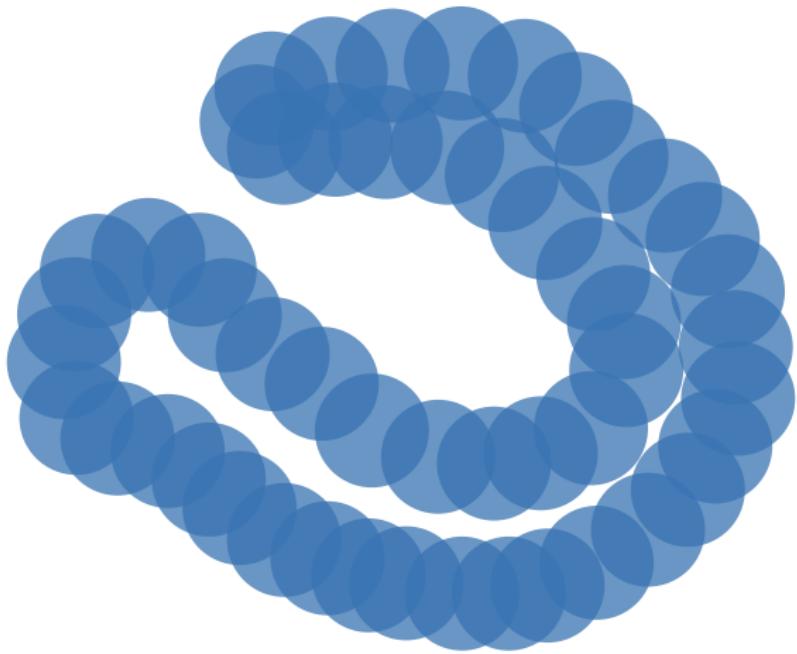


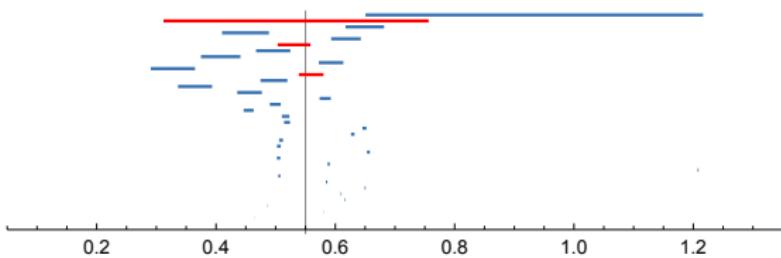
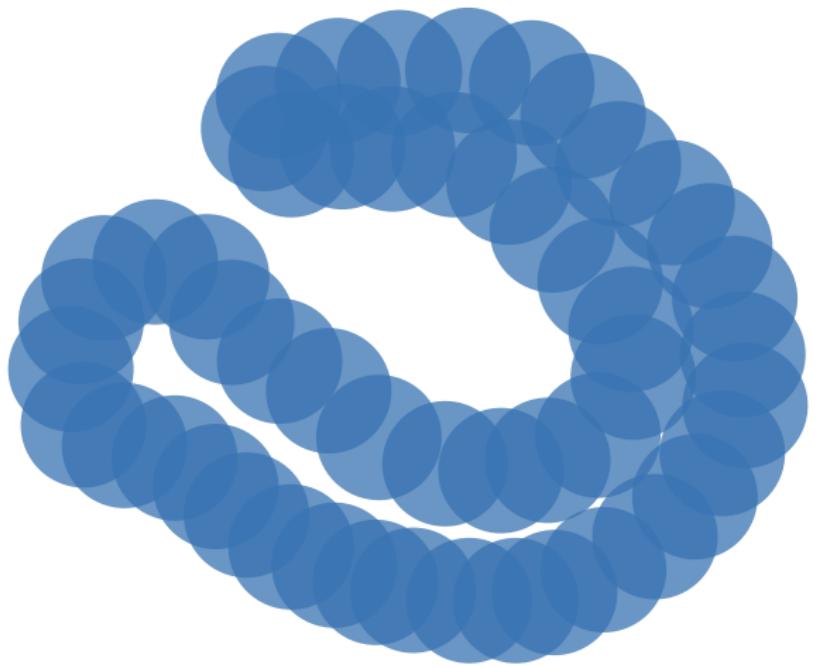


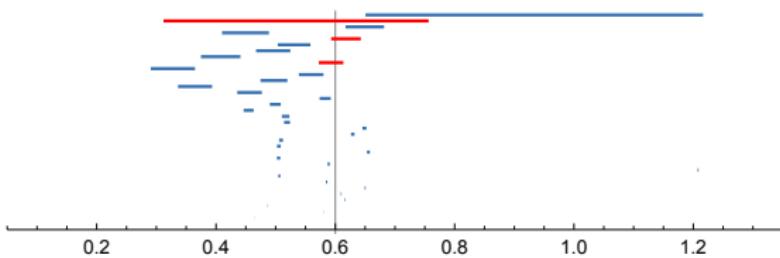
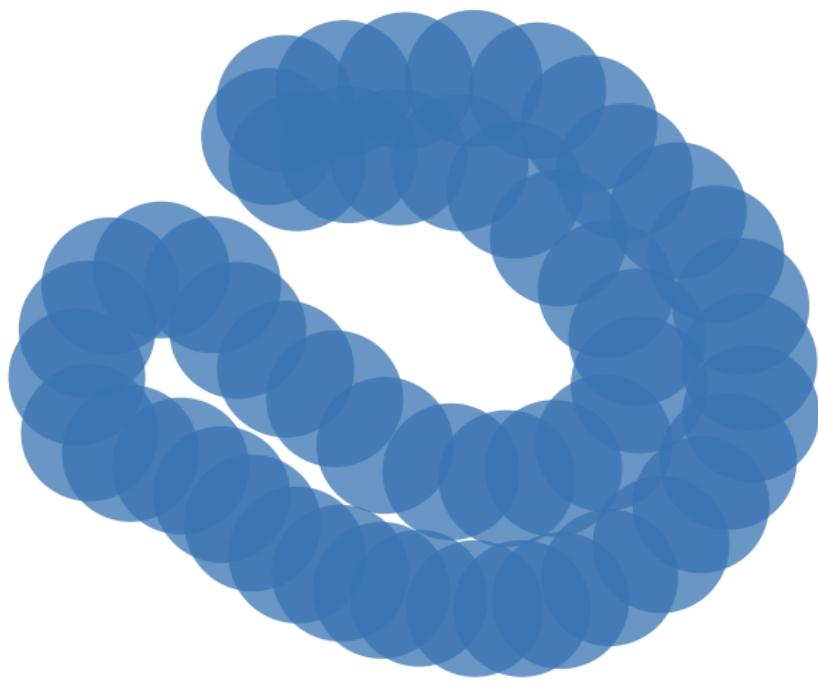


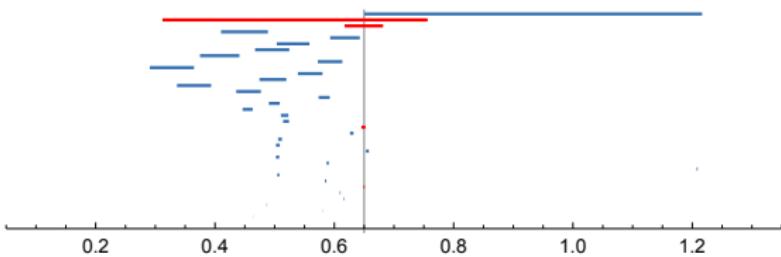
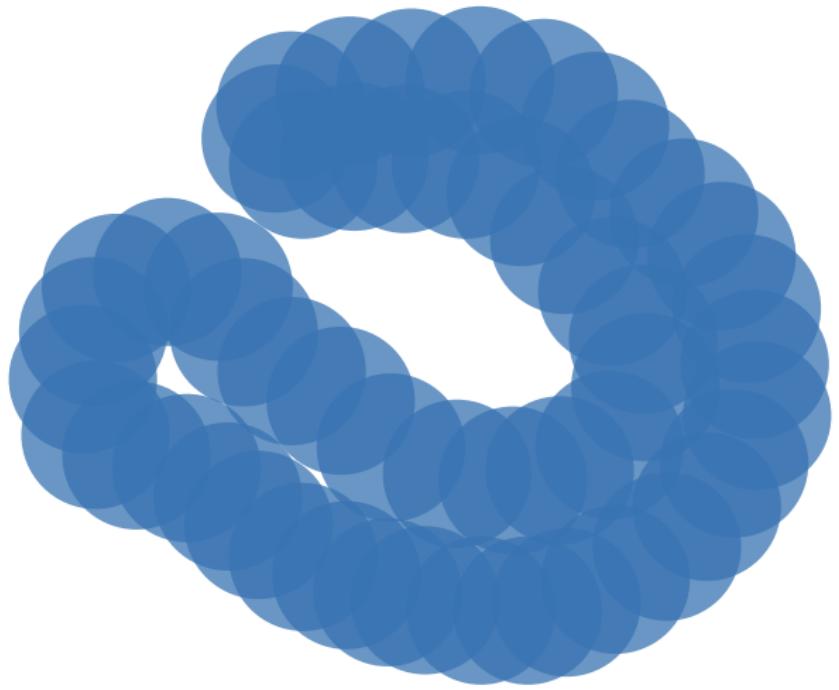


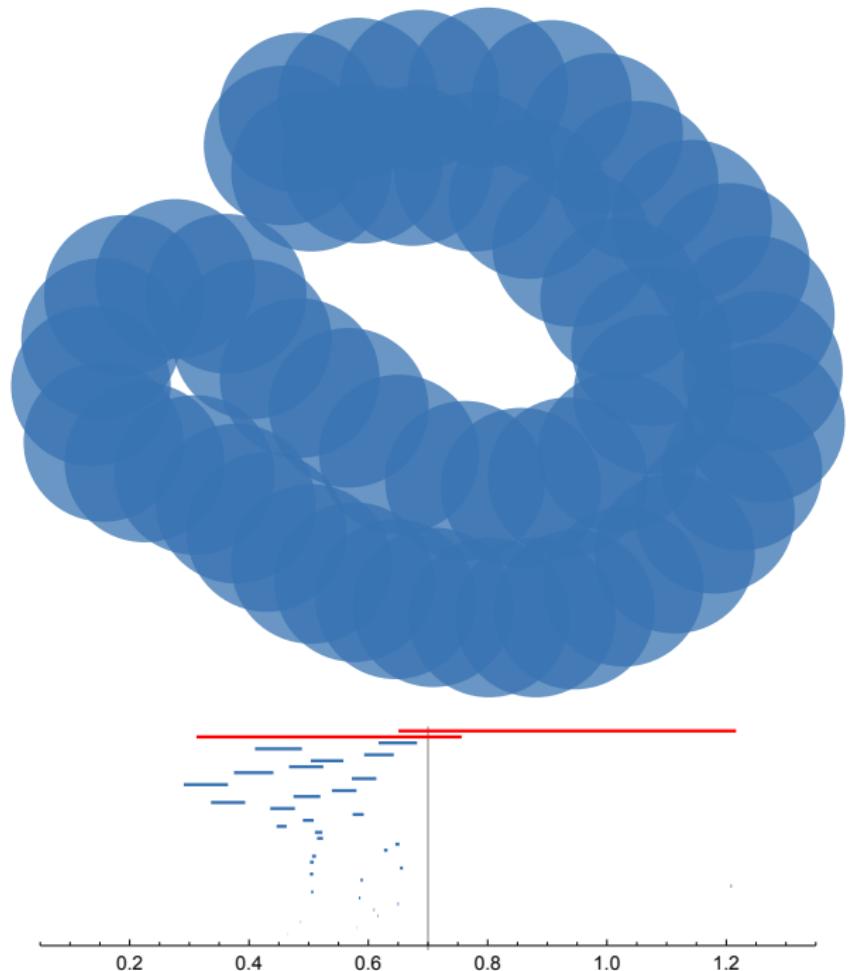


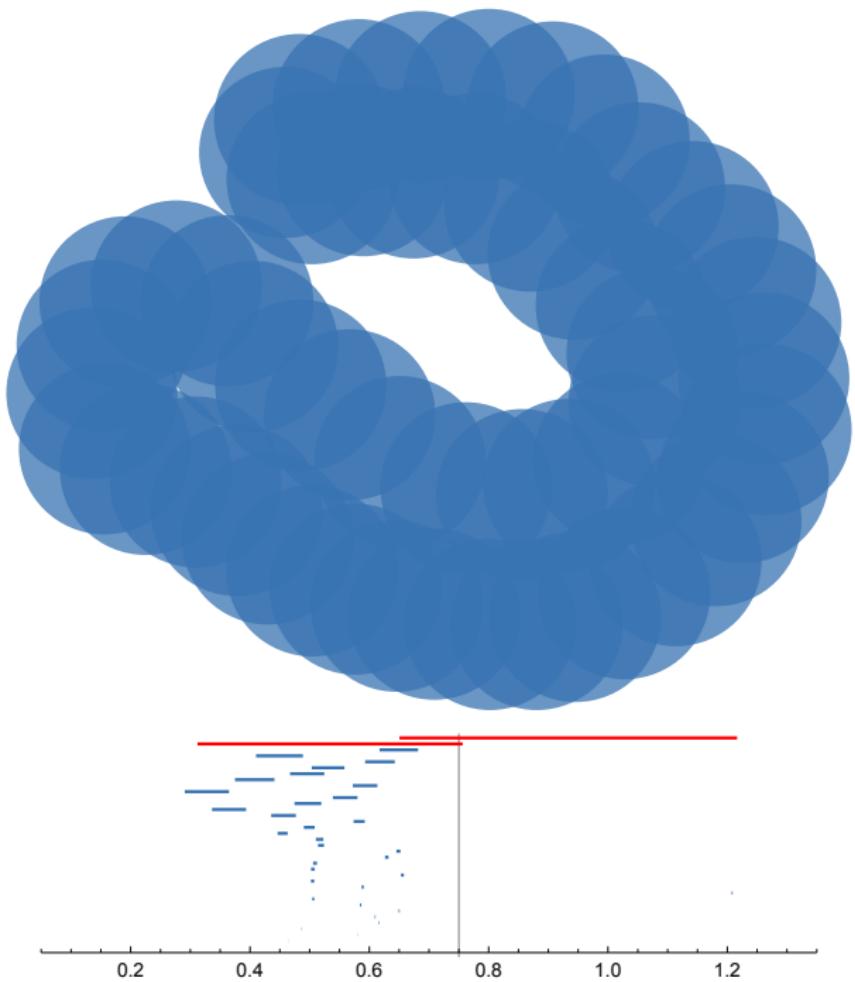


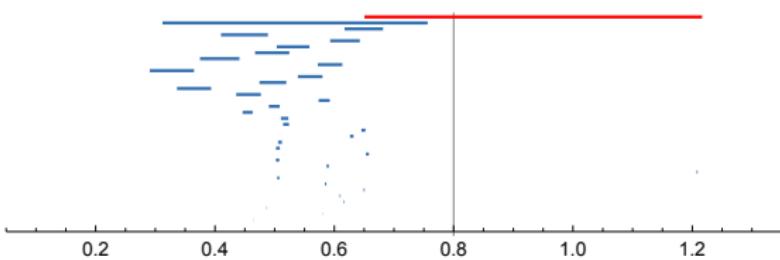
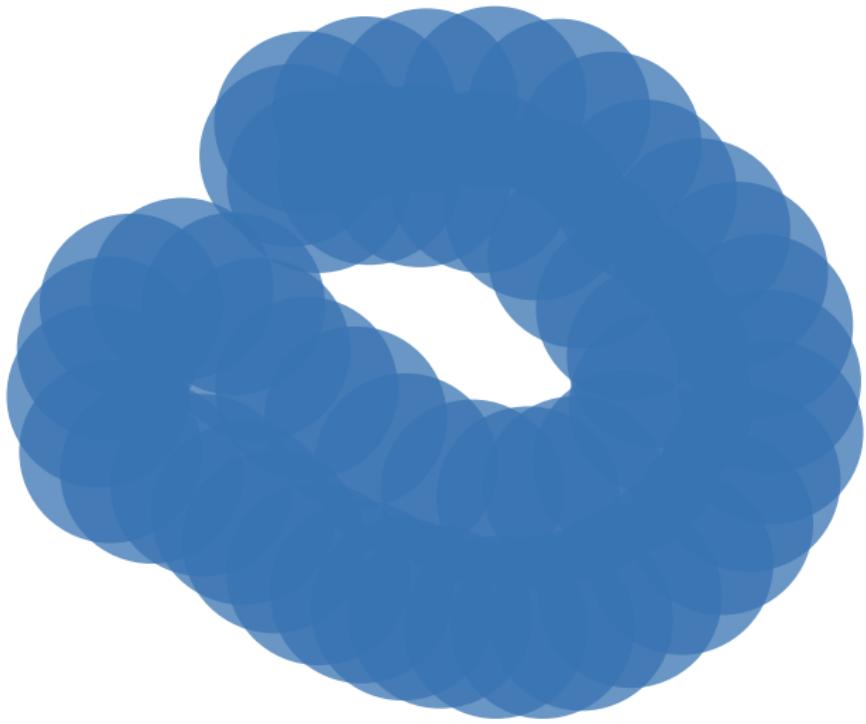


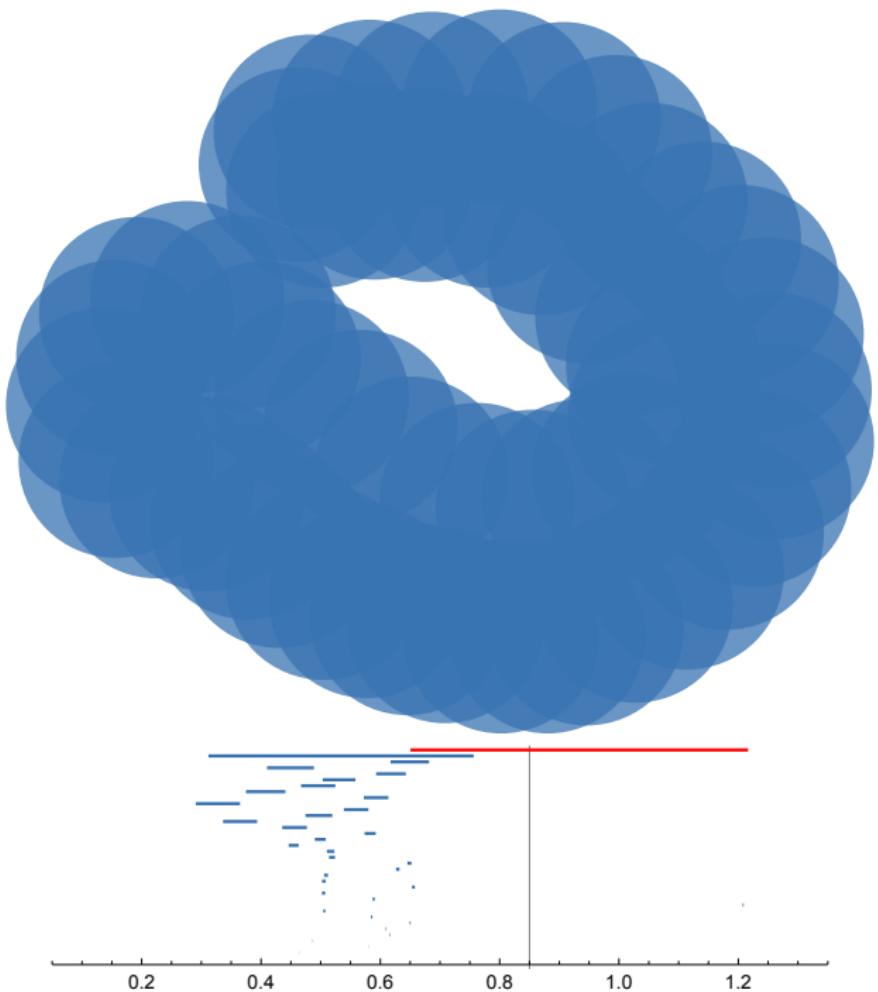


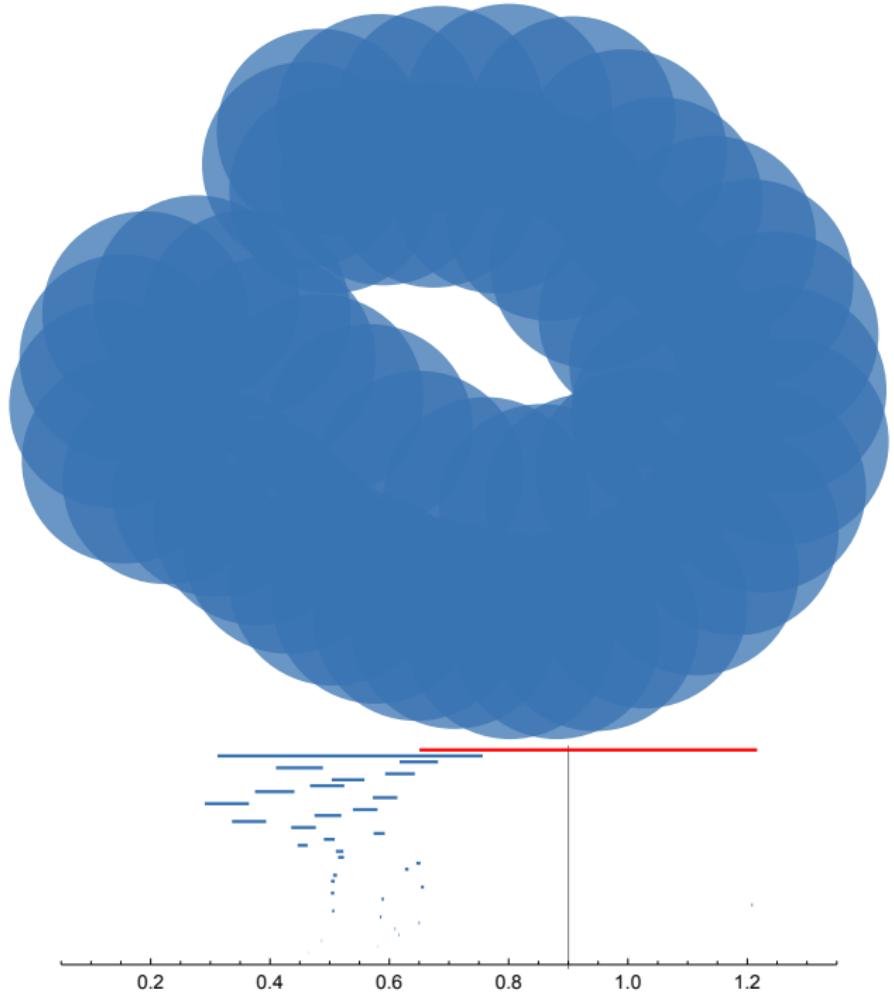












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Let $\Omega \subset \mathbb{R}^d$. Let $P \subset \Omega$ be such that

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- ▶ We say that P is a *homological δ -sample* of Ω .

Homology inference using persistent homology

$P_\delta = B_\delta(P)$: δ -neighborhood (union of balls) around P

Theorem (Cohen-Steiner, Edelsbrunner, Harer 2005)

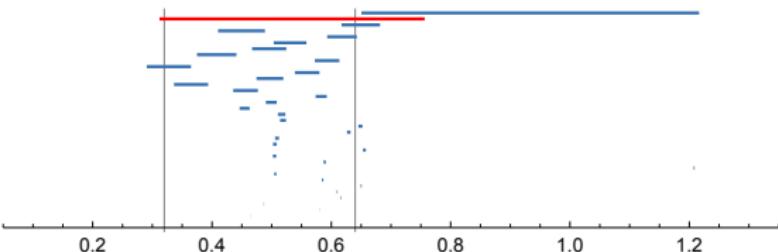
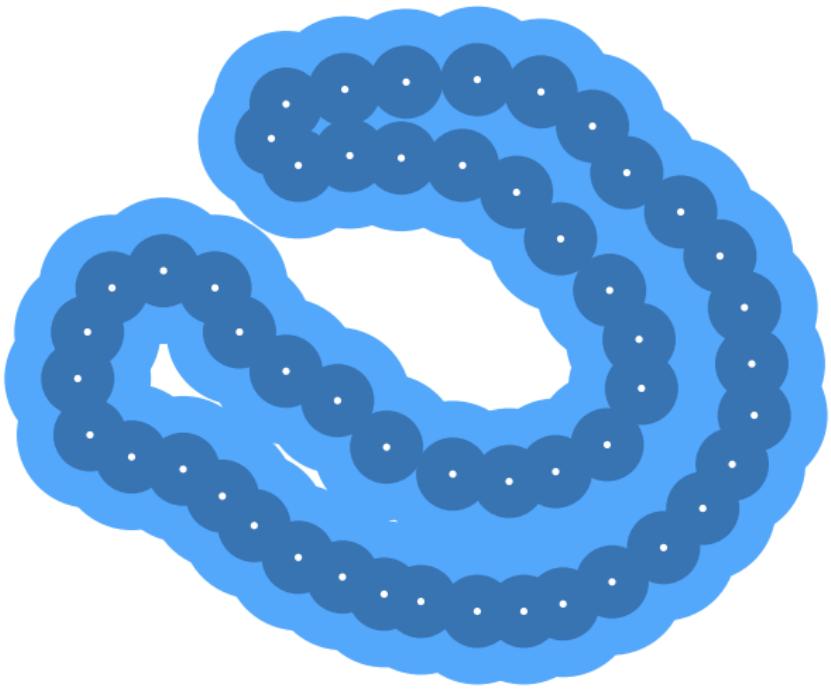
Let $\Omega \subset \mathbb{R}^d$. Let $P \subset \Omega$ be such that

- ▶ $\Omega \subseteq P_\delta$ for some $\delta > 0$ and
- ▶ both $H_*(\Omega \hookrightarrow \Omega_\delta)$ and $H_*(\Omega_\delta \hookrightarrow \Omega_{2\delta})$ are isomorphisms.

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- ▶ We say that P is a *homological δ -sample* of Ω .
- ▶ The image $\text{im } H_*(P_\delta \hookrightarrow P_{2\delta})$ is called a *persistent homology group*.



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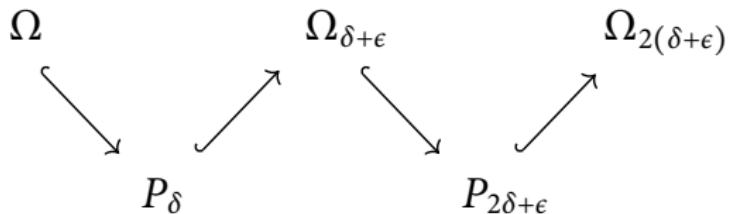
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Homological realization

This motivates the *homological realization problem*:

Problem

Given a simplicial pair $L \subseteq K$, find X with $L \subseteq X \subseteq K$ such that

$$H_*(X) = \text{im } H_*(L \hookrightarrow K).$$

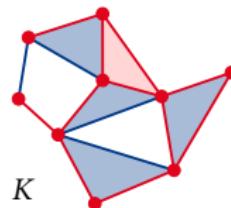
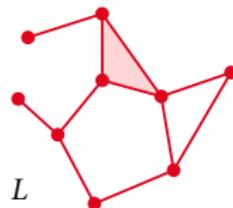
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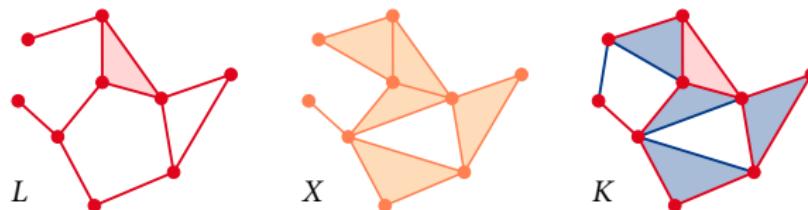
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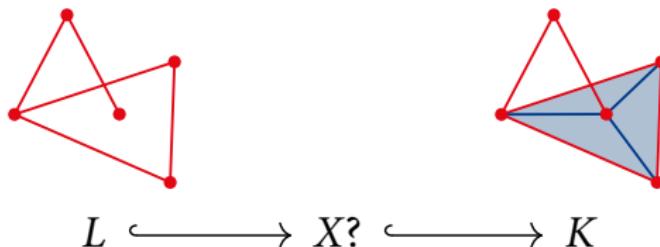
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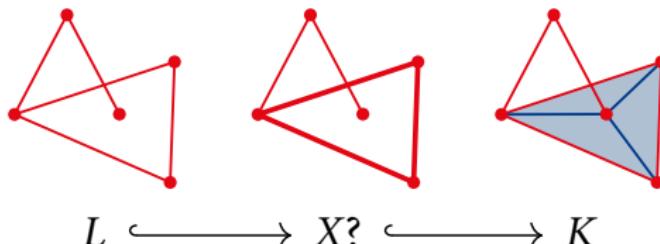
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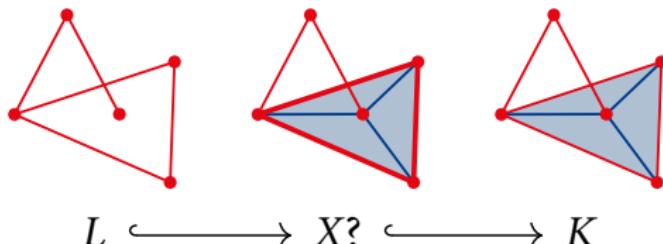
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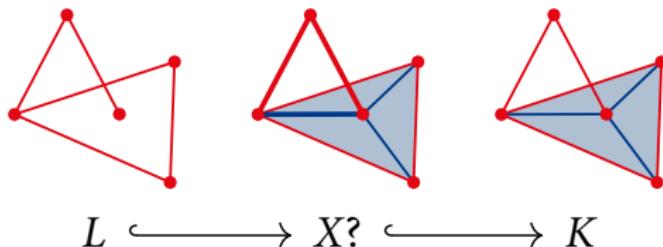
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Homological realization in \mathbb{R}^3

Theorem (Attali, B, Devillers, Glisse, Lieutier 2012)

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Then the homological realization problem for the pair

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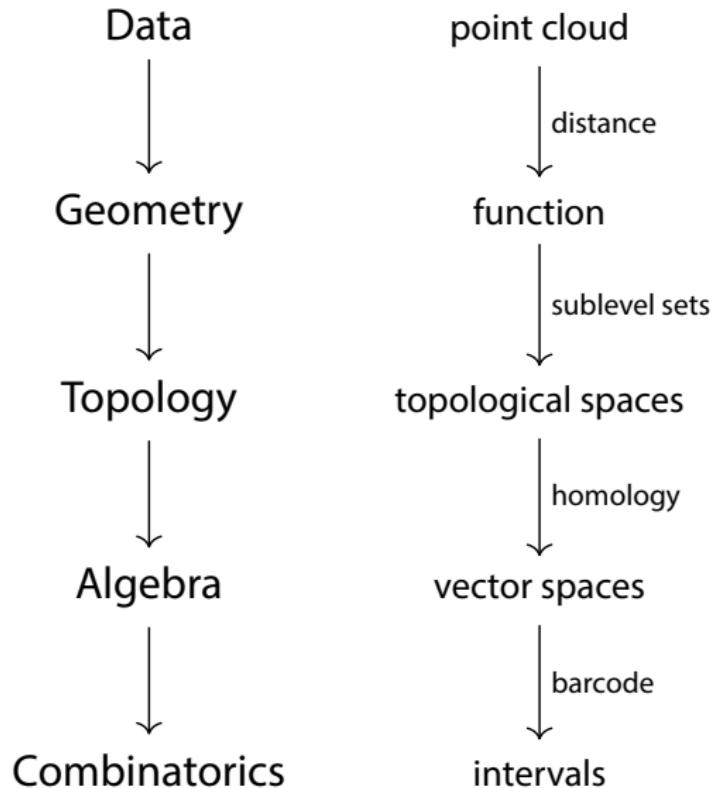
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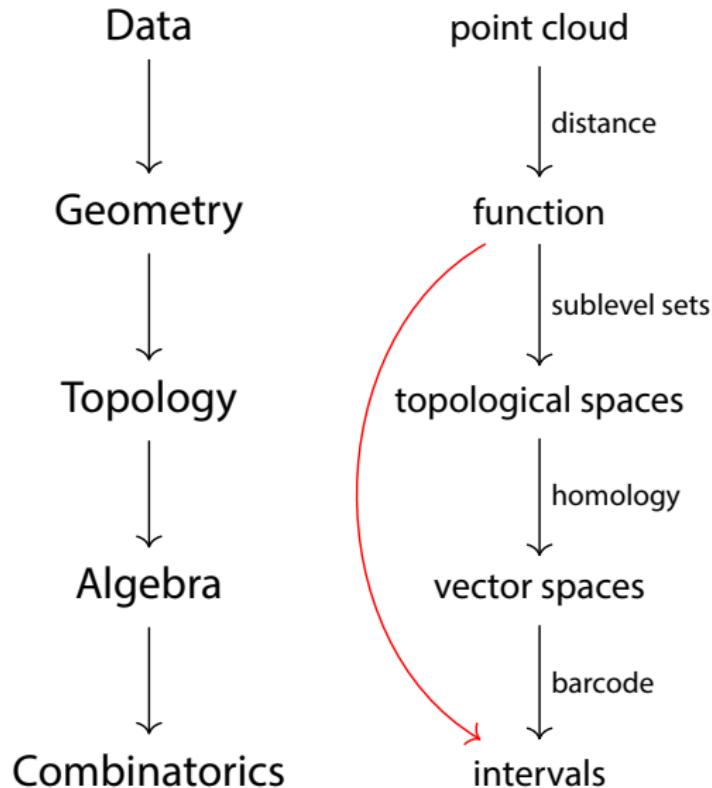
- ▶ If a solution exists, it is a homological reconstruction of Ω .
- ▶ Provides homological reconstruction under much weaker assumptions
- ▶ Even though the pair $P_\delta \subseteq P_{2\delta}$ has the reconstruction Ω_δ , the pair $\text{Del}_\delta(P) \subseteq \text{Del}_{2\delta}(P)$ might not have a reconstruction

Computation

Persistent homology of sublevel sets



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Computational assumptions

For simplicity:

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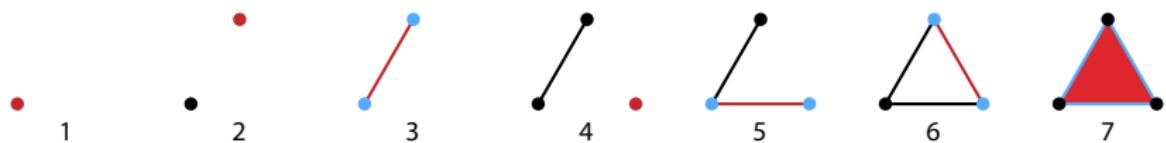
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Computational assumptions

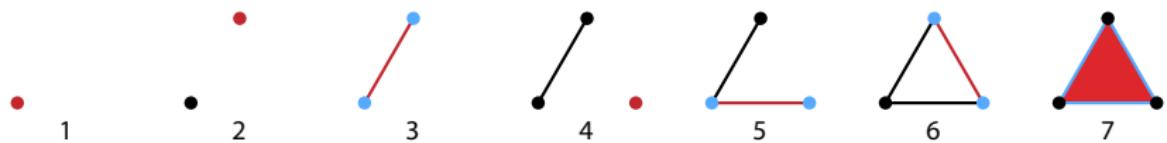
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Example: filtration and boundary matrix



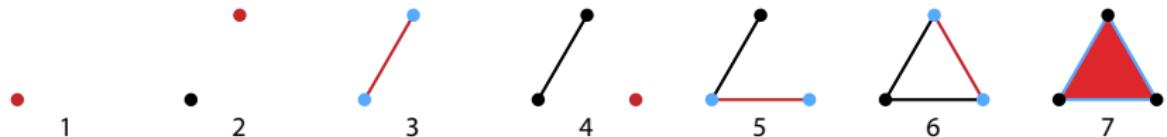
Example: filtration and boundary matrix



$D =$

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

Matrix reduction

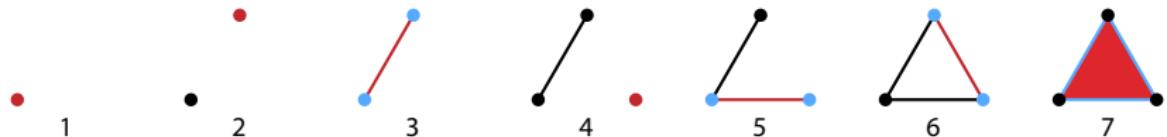


	1	2	3	4	5	6	7
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$= D \cdot$

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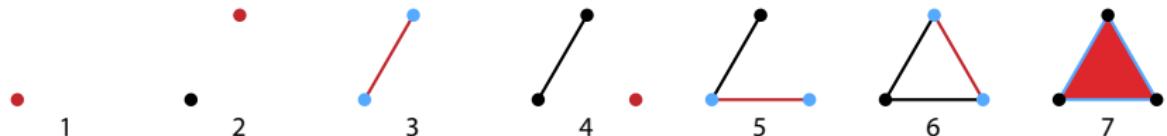
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1	1						
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Pivot of column m_j :

- ▶ largest index with nonzero entry

Matrix reduction



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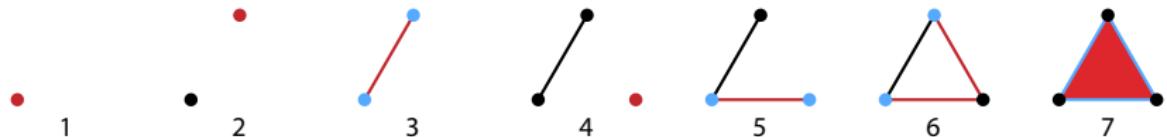
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Matrix reduction algorithm:

- ▶ while there are $i < j$ with pivot $m_i =$ pivot m_j
 - ▶ add m_i to m_j

Matrix reduction



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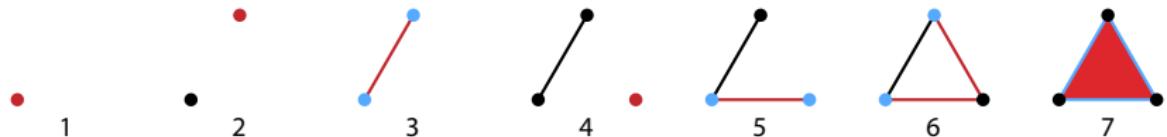
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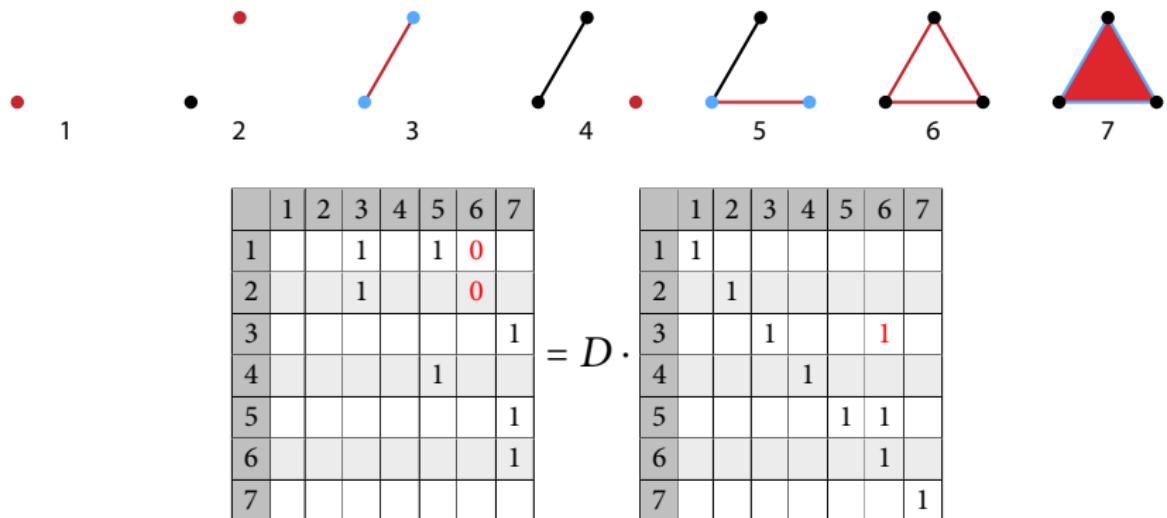
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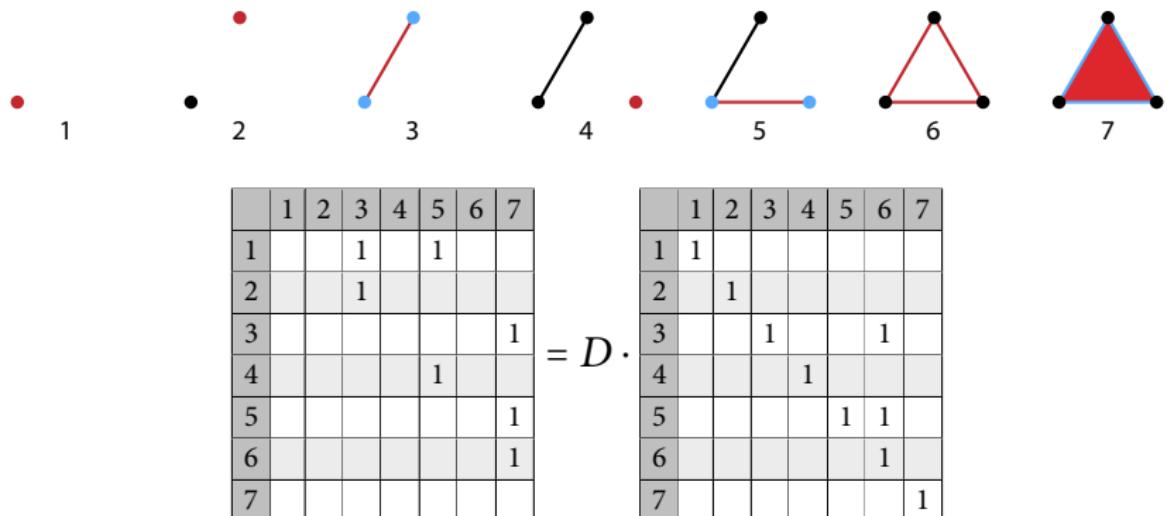
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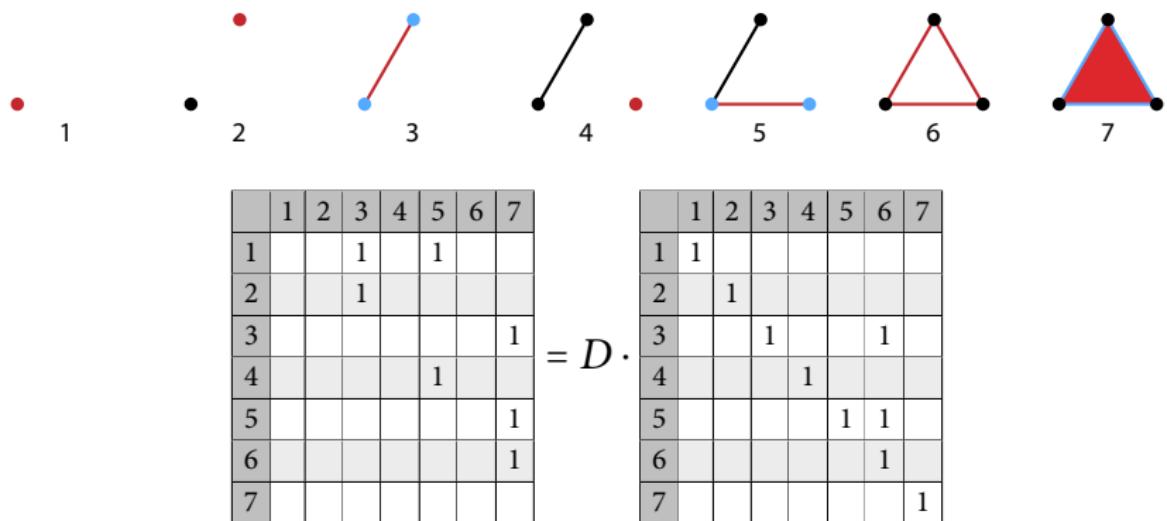
Matrix reduction



Column m_j is reduced:

- ▶ pivot of col m_j minimal under left-to-right column additions

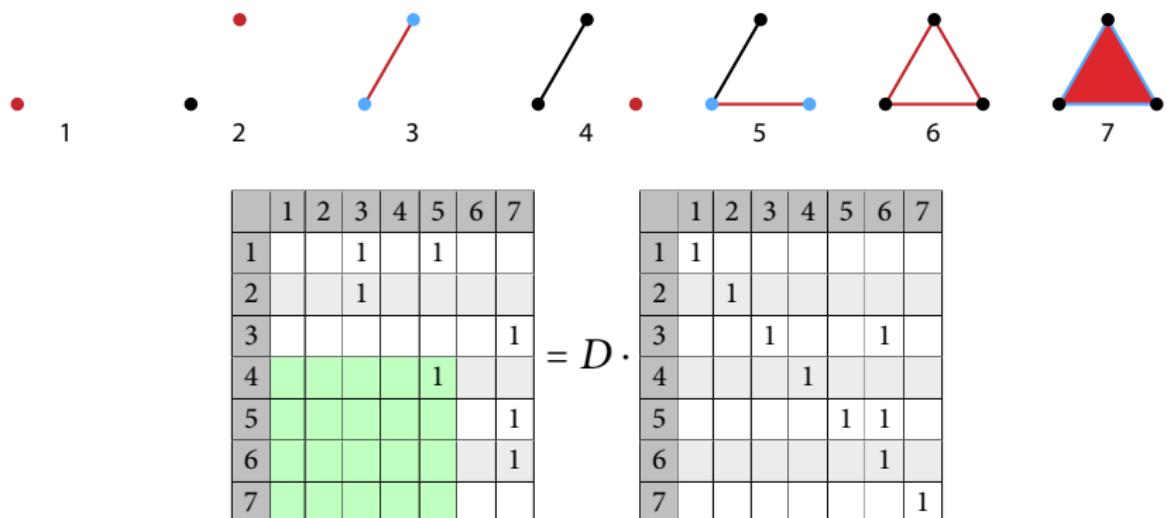
Matrix reduction



Matrix M is *reduced*:

- ▶ all columns are reduced (equivalently: pivots are unique)

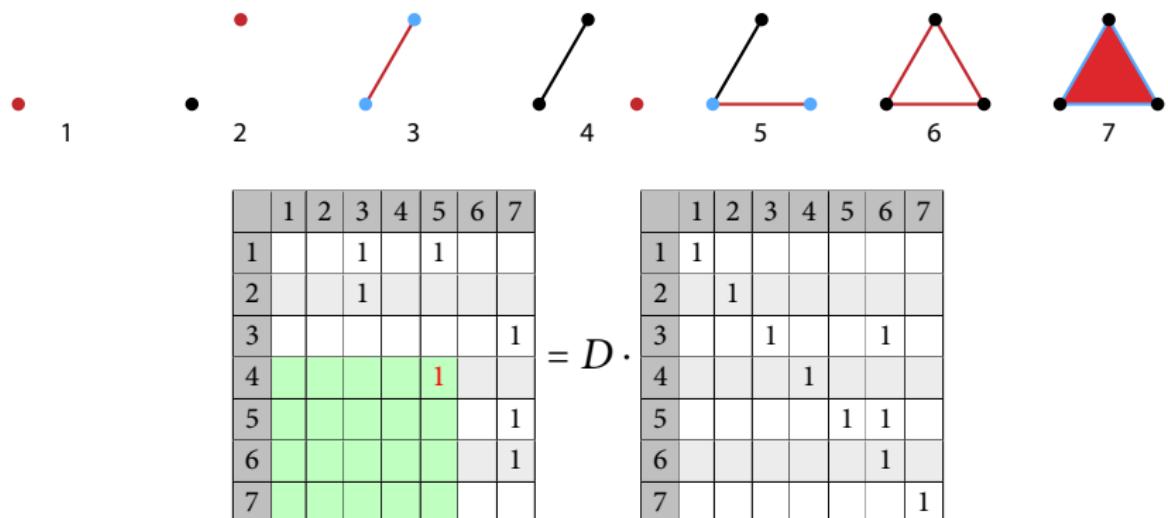
Matrix reduction



Matrix M is *reduced at index* (i, j) :

- submatrix with rows $\geq i$ and cols $\leq j$ (lower left) is reduced

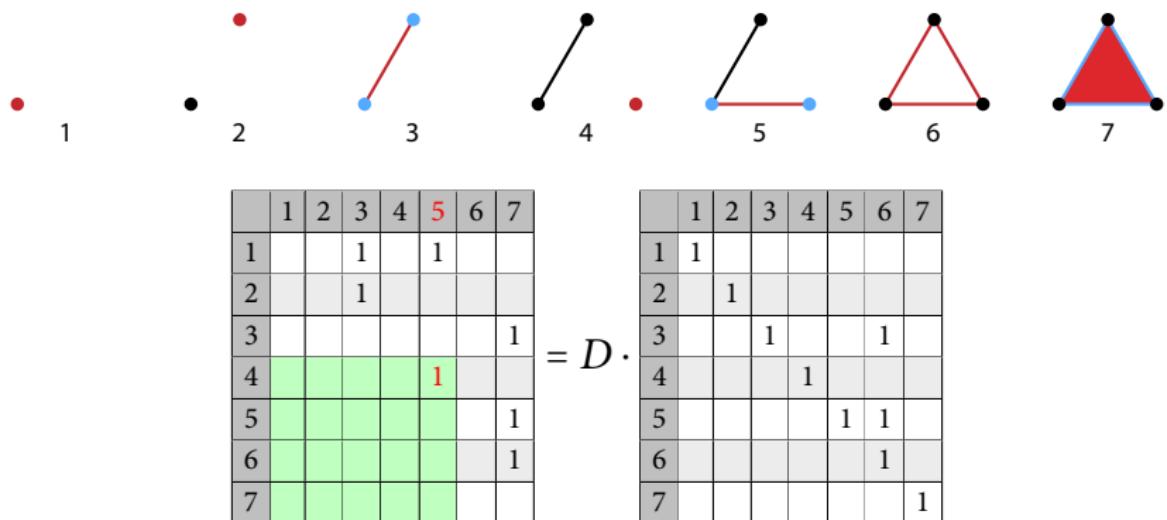
Matrix reduction



$i = \text{pivot } m_j$ and M is reduced at index $(i, j) \Rightarrow$

- ▶ column m_j is reduced
- ▶ (i, j) is a *persistence pair*:
homology is created at step i and killed at step j

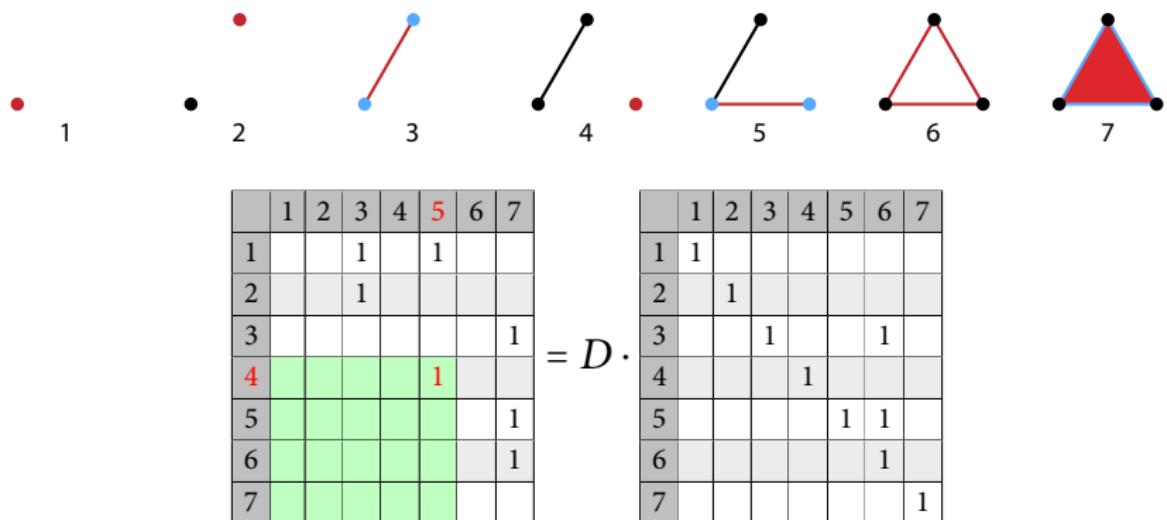
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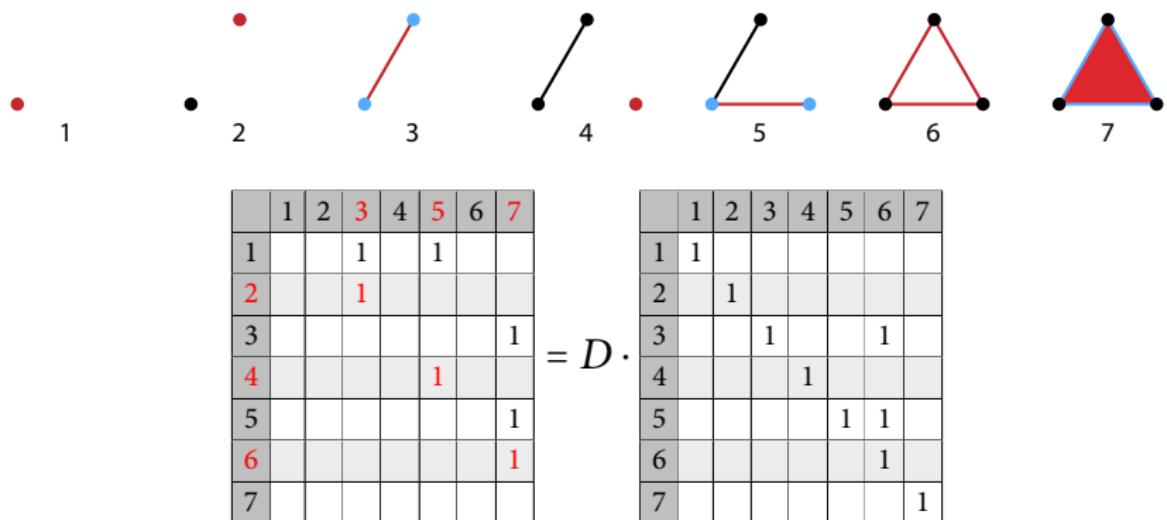
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Then the persistence barcode of $H_(C_n)$ consists of*

$$\{[i, j) : i = \text{pivot } r_j\} \cup \{[i, \infty) : r_i = 0, i \neq \text{pivot } r_j \text{ for any } j\},$$

where r_j is the j th column of R .

Proof.

Let v_i denote the i th column of V and r_j the j th column of R .

For each k :

- ▶ Basis for cycles of C_k : $b_Z = \{v_i : r_i = 0, i \leq k\}$

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- ▶ The additional cycles generate a basis for homology:

$$b_H = \tilde{b}_Z \setminus b_B = \{r_j \neq 0 : i \leq k < j, i = \text{pivot } r_j\} \cup \\ \{v_i : r_i = 0, i \leq k, i \neq \text{pivot } r_j \text{ for all } j\}$$

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Let v_i denote the i th column of V and r_j the j th column of R .

For each k :

- ▶ Basis for cycles of C_k : $b_Z = \{v_i : r_i = 0, i \leq k\}$
- ▶ Basis for boundaries of C_k : $b_B = \{r_j \neq 0 : j \leq k\}$
- ▶ Extend this basis to another basis for cycles:

$$\tilde{b}_Z = \{r_j \neq 0 : \text{pivot } r_j \leq k\} \cup \{v_i : r_i = 0, i \leq k, i \neq \text{pivot } r_j \text{ for all } j\}$$

- ▶ The additional cycles generate a basis for homology:

$$b_H = \tilde{b}_Z \setminus b_B = \{r_j \neq 0 : i \leq k < j, i = \text{pivot } r_j\} \cup \\ \{v_i : r_i = 0, i \leq k, i \neq \text{pivot } r_j \text{ for all } j\}$$

