

4 - Homology Inference

Clément Maria

EMAp Summer Program 2023

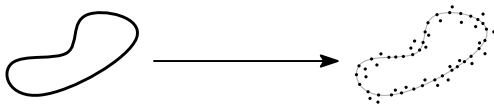
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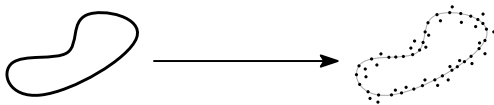
Homology Inference Problem



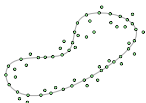
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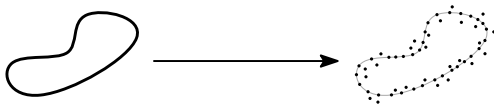
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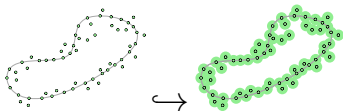
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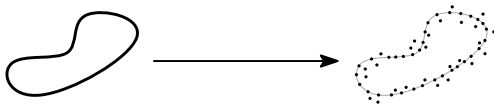
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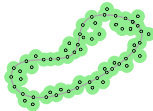
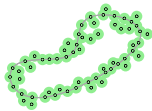
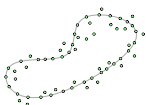
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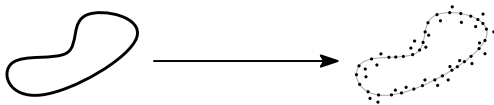
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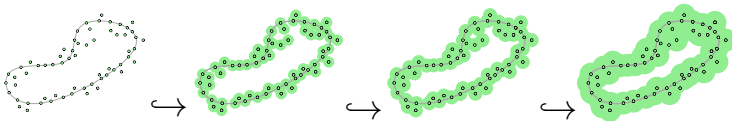
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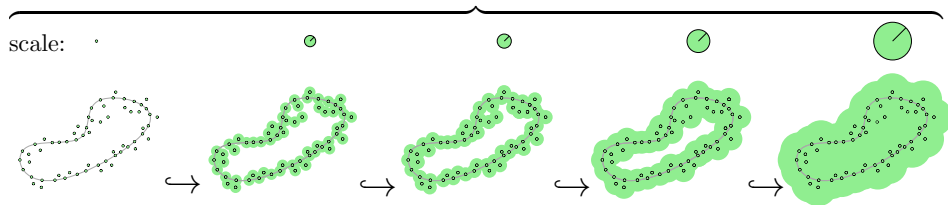
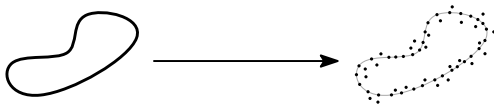
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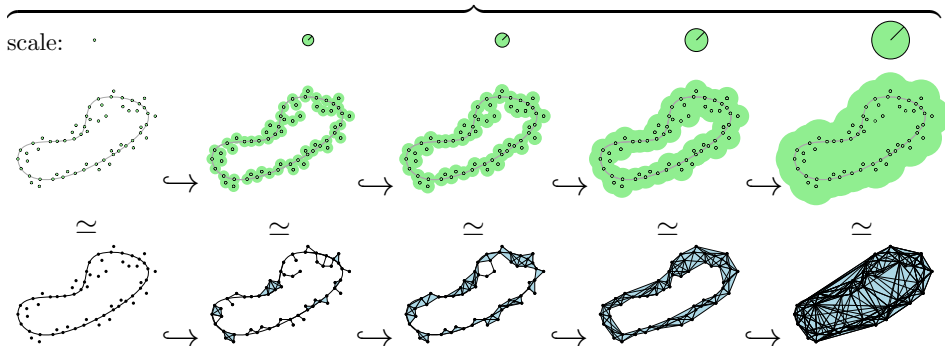
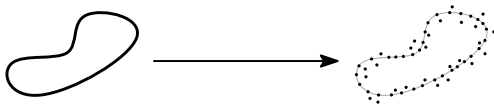
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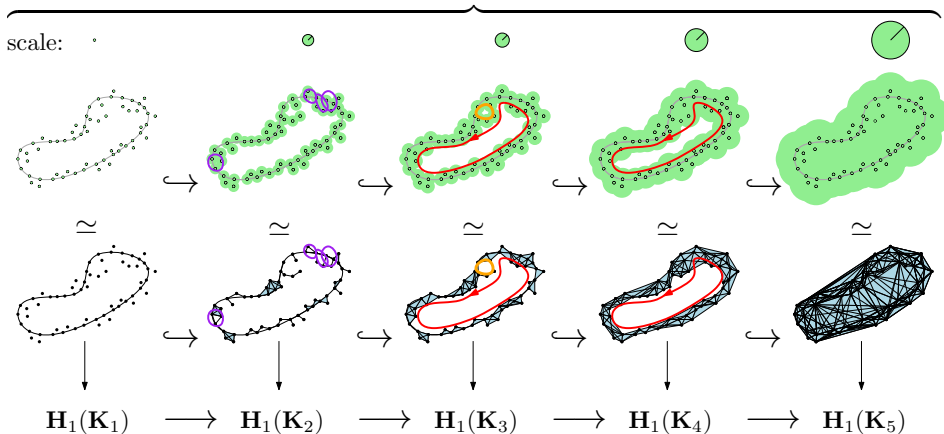
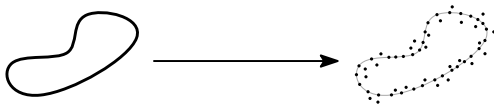
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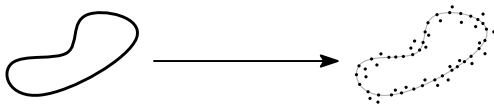
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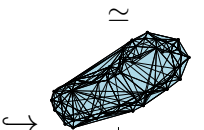
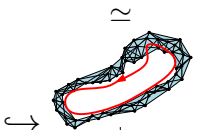
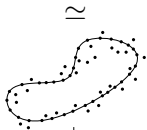
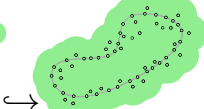
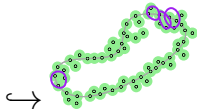
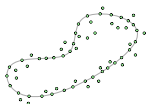
Homology Inference Problem



Homology Inference Problem



scale: 



$H_1(K_1)$

$\longrightarrow H_1(K_2)$

$\longrightarrow H_1(K_3)$

$\longrightarrow H_1(K_4)$

$\longrightarrow H_1(K_5)$

barcode



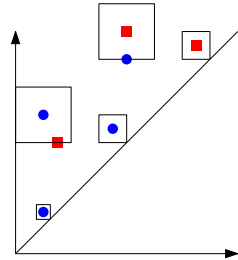
Stability Theorems

Let $D = \{(b_i, d_i)\}_{i \in I} \cup \{x = y\}$ and $D' = \{(b'_j, d'_j)\}_{j \in J} \cup \{x = y\}$ be two persistence diagrams, $b, d \in \mathbb{R}$.

The **bottleneck distance** $d_B(D, D')$ between D and D' is:

$$d_B(D, D') := \inf_{\substack{\Phi: D \rightarrow D' \\ \text{bijection}}} \sup_{p \in D} \|p - \Phi(p)\|_\infty$$

Defined even when $|I| \neq |J|$ by sending points to the diagonal.



Theorem (Stability on a simplicial complex)

Let $f, g: \mathbf{K} \rightarrow \mathbb{R}$ be two functions on a same simplicial complex \mathbf{K} , inducing filtrations $\mathbf{K}_\alpha = f^{-1}((-\infty; \alpha])$ and $\mathbf{K}'_\gamma = g^{-1}((-\infty; \gamma])$. Then

$$d_B(D(f), D(g)) \leq \|f - g\|_\infty.$$

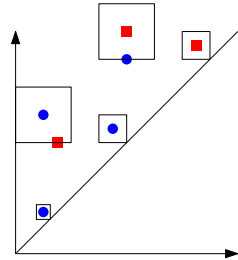
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Theorem (Stability on general space)

Let $f, g: M \rightarrow \mathbb{R}$ be two functions on a same metric space M , satisfying some “tameness” conditions. If $\exists \varepsilon \geq 0$ s.t. $\forall r \in \mathbb{R}$

$f^{-1}(-\infty; r] \subseteq g^{-1}(-\infty; r + \varepsilon]$ and $g^{-1}(-\infty; r] \subseteq f^{-1}(-\infty; r + \varepsilon]$,

then $d_B(D(f), D(g)) \leq \varepsilon$.

Distance between spaces

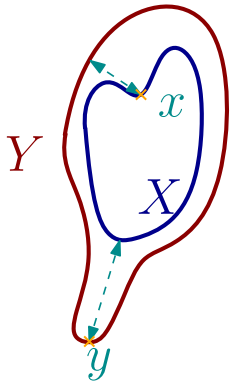
Definition (Hausdorff distance)

The **Hausdorff distance** between two non-empty subsets X, Y of a metric space (M, d) is:

$$d_H(X, Y) := \max \left\{ \sup_{x \in X} d(x, Y), \sup_{y \in Y} d(X, y) \right\},$$

where,

$d(x, Y) = \inf_{y \in Y} d(x, y)$ and $d(X, y) = \inf_{x \in X} d(x, y)$.



Definition (ε -sample)

Let $K \subset \mathbb{R}^D$ be a compact set. An **ε -sample** of K , for some $\varepsilon \geq 0$, is a finite set of points P such that $d_H(P, K) \leq \varepsilon$.



Sampling compacts

Now consider compacts K in Euclidean space \mathbb{R}^D . Any compact K defines a function:

$$d_K: \mathbb{R}^D \rightarrow \mathbb{R}, \quad x \mapsto d_K(x) = d(x, K).$$

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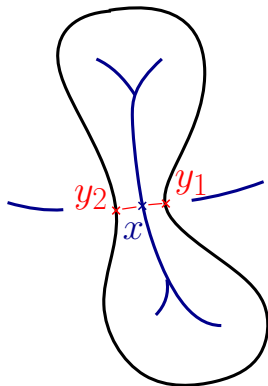
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Definition (Critical point)

A point $x \in \mathbb{R}^D$ is a **critical point** for d_K if there exist distinct points $y_1, y_2 \in K$ such that:

$$d(x, y_1) = d(x, y_2) = d(x, K).$$

The set of critical points is called the **medial axis** of K .



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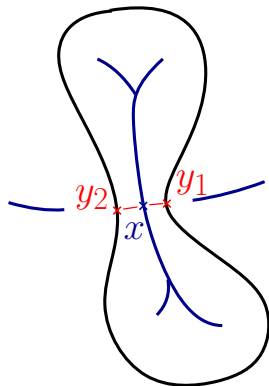
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Definition (Reach of a compact)

The **reach** of a compact K is:

$$\text{reach } K := \inf\{d_K(x) : x \text{ critical point for } d_K\}.$$



Sampling, distance function approach

Lemma

Let P be an ε -sample of a compact K . Then,

$$||d_K - d_P||_{\infty} \leq \varepsilon.$$

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Consequently, by triangle inequality:

$$d_P(x) \leq d(x, p_0) \leq d(x, y_0) + d(y_0, p_0) \leq d_K(x) + \varepsilon.$$

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And vice versa, exchanging K and P .

Reconstruction theorem

Theorem (Reconstruction)

Let $K \subset \mathbb{R}^D$ be a compact set, and $\varepsilon > 0$ be such that:

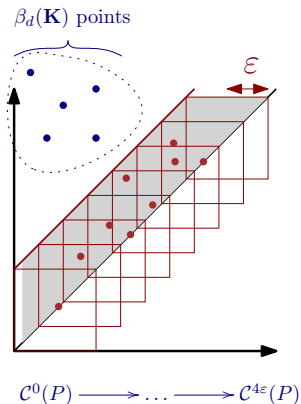
$$0 < 4\varepsilon < \text{reach } \mathbf{K}.$$

Let P be an ε -sample of K . Then,

- there is a *one-to-one correspondence* between the points of the persistence diagram of the Čech filtration:

$$\check{C}^0(P) \longrightarrow \dots \longrightarrow \check{C}^{4\varepsilon}(P)$$

that are *at least ε -away* from the diagonal Δ in ∞ norm, and the homology features of K .



Reconstruction theorem

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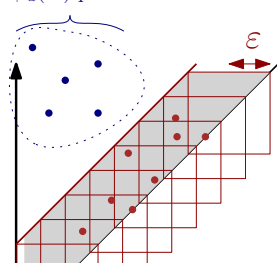
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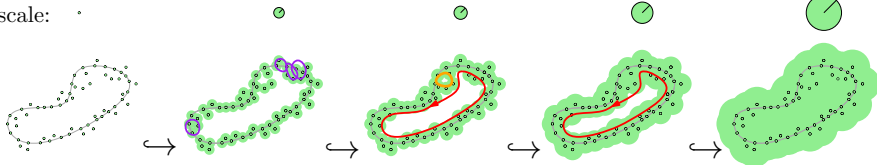
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$\beta_d(\mathbf{K})$ points



scale: \cdot



barcode



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Sketch of proof:

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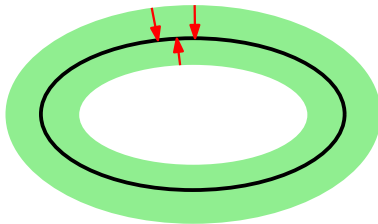
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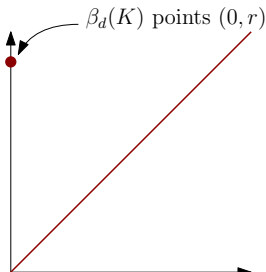
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—→ by projecting each $x \in d_K^{-1}(-\infty; r] \setminus K$ to its **unique** nearest neighbor on K .

Consequently, for all $0 < r_1 \leq r_2 < \text{reach}(K)$ we have

$$\mathbf{H}(d_K^{-1}(-\infty; r_1]) \xrightarrow{\text{id}} \mathbf{H}(d_K^{-1}(-\infty; r_2])$$



Reconstruction theorem

Sketch of proof:

(2) Consider the persistent homology of

$$d_p^{-1}(-\infty; r] = \bigcup_{p \in P} \mathcal{B}_r(p) \simeq |\check{\mathcal{C}}^r(P)| \text{ (Nerve lemma).}$$

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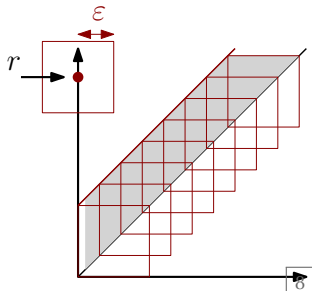
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For $\text{reach}(K) > r > 4\varepsilon > 0$, the radius r $\|\cdot\|_\infty$ -ball in \mathbb{R}^2 around $(0, r)$, and the $\|\cdot\|_\infty$ -band of width ε around the diagonal are disjoint.



Rips complex

Computing intersections of balls in \mathbb{R}^D is *expensive* (high arithmetic complexity): Approximate the Čech complex instead.

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Definition (Flag complex)

Let $G = (V, E)$ be a graph. The **flag complex** induced by G is the abstract simplicial complex $\mathcal{F}(G)$ satisfying:

- the vertices of $\mathcal{F}(G)$ are V ,
- $\sigma = \{v_0, \dots, v_d\} \in \mathcal{F}(G)$ iff v_0, \dots, v_d form a clique in G (i.e., an induced complete subgraph of G).

It is the maximal simplicial complex with G as 1-skeleton.

Rips complex

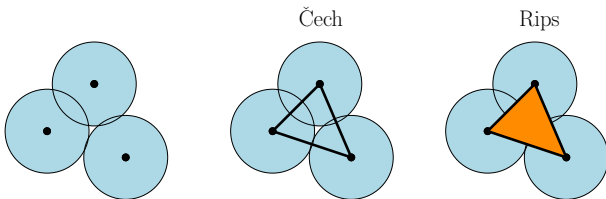
Computing intersections of balls in \mathbb{R}^D is *expensive* (high arithmetic complexity): Approximate the Čech complex instead.

Definition (Rips complex)

Let $P = \{x_1, \dots, x_n\}$ be a set of points in \mathbb{R}^N . The **neighbor graph of threshold r** of P is the graph $N_r(P)$ with:

- vertices the points in P , and
- any two points x_i, x_j , $i \neq j$ are connected by an edge iff $\|x_i - x_j\| \leq r$.

The **Rips complex** of threshold r , denoted by $\mathcal{R}^r(P)$, is the flag complex $\mathcal{F}(N_{2r}(P))$.



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Computation: Simply compute pairwise distances (low arithmetic complexity predicate), then expand the flag complex combinatorial

$\sim O(\#\text{points}^2) + \text{linear in the number of cliques/size of the complex.}$

Rips vs Čech

Lemma

For a point cloud P in Euclidean space, and any $r > 0$:

$$\check{C}^r(P) \subseteq \mathcal{R}^r(P) \subseteq \check{C}^{2r}(P).$$

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$\check{C}^r(P) \subseteq \mathcal{R}^r(P)$. Let $\sigma = \{x_0, \dots, x_k\} \in \check{C}^r(P)$, $k \geq 1$, then for any $i \neq j$, $0 \leq i, j \leq k$ we have $\mathcal{B}_r(x_i) \cap \mathcal{B}_r(x_j) \neq \emptyset \Rightarrow \|x_i - x_j\| \leq 2r$. Hence x_0, \dots, x_k form a clique in $N_{2r}(P)$ and $\sigma \in \mathcal{R}^r(P)$.

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Consequently, for all i :

$$\|z - x_i\| \leq \text{diam}(\text{CH}(x_0, \dots, x_k)) \leq \max_{i,j} \|x_i - x_j\| \leq 2r.$$

In conclusion, $z \in \bigcap_{i=0\dots k} \mathcal{B}_{2r}(x_i) \neq \emptyset$ and $\sigma \in \check{C}^{2r}(P)$.

Ranks in sequence of morphisms

Lemma

Consider the following sequence of vector spaces and morphisms:

$$A \xrightarrow{\alpha} B \xrightarrow{\beta} C \xrightarrow{\gamma} D \xrightarrow{\delta} E \xrightarrow{\eta} F$$

such that $\text{rk} (A \xrightarrow{\eta \circ \dots \circ \alpha} F) = \text{rk} (C \xrightarrow{\gamma} D) = k$, then:

$$\text{rk} (B \xrightarrow{\delta \circ \gamma \circ \beta} E) = k.$$

Ranks in sequence of morphisms

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Consider the following sequence of vector spaces and morphisms:

$$A \xrightarrow{\alpha} B \xrightarrow{\beta} C \xrightarrow{\gamma} D \xrightarrow{\delta} E \xrightarrow{\eta} F$$

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- $\boxed{A \rightarrow B \rightarrow E \rightarrow F}$ $k = \text{rk} (A \rightarrow F) \leq \text{rk} (B \rightarrow E),$
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Cech persistence, a closer look

Lemma

Let P be an ε -sample of a compact K of $\text{reach}(K) > 0$ in \mathbb{R}^N . Then, for any $\varepsilon_0 > \varepsilon$, and α , such that $\varepsilon \leq \alpha < \text{reach}(K) - 2\varepsilon_0$, we have:

$$\text{rk} (\mathbf{H}_d(\check{C}^\alpha(P)) \longrightarrow \mathbf{H}_d(\check{C}^{\alpha+\varepsilon_0}(P))) = \dim \mathbf{H}_d(K)$$

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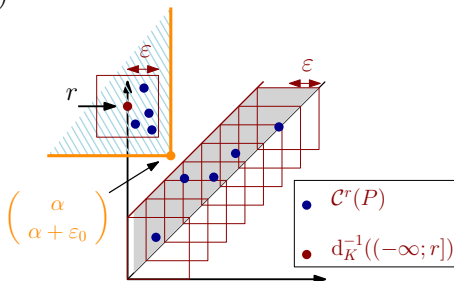
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Theorem

Let P be an ε -sample of a compact K s.t. $0 < \varepsilon < \frac{\text{reach}(K)}{8}$. The homology of K can be read off the persistence diagram of $\mathcal{R}^r(P)$ for $r \geq \varepsilon + 2\varepsilon_0$, for any $\varepsilon < \varepsilon_0 < \frac{\text{reach}(K)}{8}$.

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In conclusion, $\dim \mathbf{H}_d(K)$ is exactly the number of points in the persistence diagram of the Rips filtration (up to $r = \varepsilon + 2\varepsilon_0$) in the upper left quadrant based at $(\varepsilon, \varepsilon + 2\varepsilon_0)$.

Morse filtration

Definition

Let \mathbf{K} be a complex, equipped with a filtration,

$$\emptyset := \mathbf{K}_0 \xrightarrow{\mathfrak{S}_1} \mathbf{K}_1 \xrightarrow{\mathfrak{S}_2} \mathbf{K}_2 \xrightarrow{\mathfrak{S}_3} \dots \xrightarrow{\mathfrak{S}_m} \mathbf{K}_m = \mathbf{K}$$

where \mathfrak{S}_i contains **at least** one simplex.

A **Morse filtration** for the filtration of \mathbf{K} is a collection of Morse matchings $(X_i \sqcup T_i \sqcup S_i, \omega_i: T_i \rightarrow S_i)$ for each \mathbf{K}_i , $i = 0 \dots m$, satisfying:

$$X_i \subseteq X_{i+1}, \quad T_i \subseteq T_{i+1}, \quad S_i \subseteq S_{i+1},$$

$$\omega_{i+1}|_{T_i} = \omega_i, \quad \partial^{X_{i+1}}|_{X_i} = \partial^{X_i}, \quad \forall i = 0 \dots m-1.$$

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and a Morse filtration $(X_i \sqcup T_i \sqcup S_i, \omega_i: T_i \rightarrow S_i)$, $i = 0 \dots m$:

$$\begin{aligned} X_i &\subseteq X_{i+1}, & T_i &\subseteq T_{i+1}, & S_i &\subseteq S_{i+1}, \\ \omega_{i+1}|_{T_i} &= \omega_i, & \partial^{X_{i+1}}|_{X_i} &= \partial^{X_i}, & \forall i &= 0 \dots m-1. \end{aligned}$$

Then the following persistence modules:

$$\mathbf{H}(\mathbf{K}_0, \partial) \xrightarrow{\subseteq} \mathbf{H}(\mathbf{K}_1, \partial) \xrightarrow{\subseteq} \mathbf{H}(\mathbf{K}_2, \partial) \xrightarrow{\subseteq} \dots \xrightarrow{\subseteq} \mathbf{H}(\mathbf{K}_m, \partial)$$

$$\mathbf{H}(X_0, \partial^{X_0}) \xrightarrow{\subseteq} \mathbf{H}(X_1, \partial^{X_1}) \xrightarrow{\subseteq} \mathbf{H}(X_2, \partial^{X_2}) \xrightarrow{\subseteq} \dots \xrightarrow{\subseteq} \mathbf{H}(X_m, \partial^{X_m})$$

are *isomorphic*.

Morse filtration

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The following persistence modules are isomorphic:

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Sketch of proof:

$$X_i \subseteq X_{i+1}, \quad T_i \subseteq T_{i+1}, \quad S_i \subseteq S_{i+1}, \quad \omega_{i+1}|_{T_i} = \omega_i, \quad \partial^{X_{i+1}}|_{X_i} = \partial^{X_i},$$

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implies that:

$$(\mathbf{C}(X_0), \partial^{X_0}) \xrightarrow{\subseteq} (\mathbf{C}(X_1), \partial^{X_1}) \xrightarrow{\subseteq} (\mathbf{C}(X_2), \partial^{X_2}) \xrightarrow{\subseteq} \dots \xrightarrow{\subseteq} (\mathbf{C}(X_m), \partial^{X_m})$$

is **well-defined** filtration of complexes

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is **well-defined** filtration of complexes inducing the persist. module:

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Morse filtration

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Sketch of proof:

For each i there is a chain equivalence:

$$(\mathbf{C}(\mathbf{K}_i), \partial) \xrightarrow{\phi_i} (\mathbf{C}(\chi_i), \partial^{\chi_i}) \xrightarrow{\psi_i} (\mathbf{C}(\mathbf{K}_i), \partial^{\chi_i})$$

Morse filtration

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where each chain map **commutes with inclusion**:

$$\begin{array}{ccc} \mathbf{C}(\mathbf{K}_i) & \xrightarrow{\subseteq} & \mathbf{C}(\mathbf{K}_{i+1}) \\ \phi_i \downarrow & & \downarrow \phi_{i+1} \\ \mathbf{C}(X_i) & \xrightarrow{\subseteq} & \mathbf{C}(X_{i+1}) \end{array}$$

$$\begin{array}{ccc} \mathbf{C}(\mathbf{K}_i) & \xrightarrow{\subseteq} & \mathbf{C}(\mathbf{K}_{i+1}) \\ \psi_i \uparrow & & \uparrow \psi_{i+1} \\ \mathbf{C}(X_i) & \xrightarrow{\subseteq} & \mathbf{C}(X_{i+1}) \end{array}$$

Morse filtration

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→ enough prove commutativity for addition of a Morse pair (τ, σ) :

$$(\mathbf{C}(X), \partial^X) \xrightarrow{\phi} (\mathbf{C}(X'), \partial^{X'}) \xrightarrow{\psi} (\mathbf{C}(X), \partial^X)$$

with $X' = X - \{\sigma, \tau\}$ as used previously (same chain maps).

Topology Inference Problem

compact K in Euclidean space



P ε -sample with

$$0 < \varepsilon < \frac{\text{reach}(K)}{8}$$

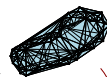


scale:



$$O(|P|^2) + \tilde{O}(n)$$

Rips filtration



$$\mathcal{R}^{3.1\varepsilon}$$

$$n \text{ simplices}$$

Morse filtration

$$(X_1, \partial^{X_1})$$

$$(X_2, \partial^{X_2})$$

$$(X_3, \partial^{X_3})$$

$$(X_4, \partial^{X_4})$$

$$(X_5, \partial^{X_5})$$

$$\tilde{O}(n)$$

m cells

Persistent homology $\mathbf{H}_1(\mathbf{K}_1)$

$\longrightarrow \mathbf{H}_1(\mathbf{K}_2)$

$\longrightarrow \mathbf{H}_1(\mathbf{K}_3)$

$\longrightarrow \mathbf{H}_1(\mathbf{K}_4)$

$\longrightarrow \mathbf{H}_1(\mathbf{K}_5)$

$$O(m^3)$$

barecode



ε

3ε