

No HW 3. (Focus on paper review/project)

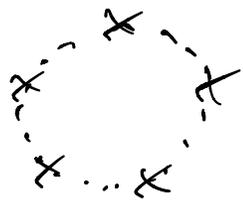
Today: Zigzag ZOO, Linear-~~S~~ approx for Rips.

2 weeks ago: sweet range for Čech/Rips

Last week: optimizations for PH. $O(n^4)$

Problem: # simplices in Rips/Čech
in k skeleton is $O(n^k)$. (can be very expensive.

$O(n^3)$
simplices



Can we use smaller # of pts
to get topology?

Idea: subsampling Recall: we have an interleaving
bound $d_H(PH(R(X)), PH(R(Y))) \leq 2d_H(X, Y)$

so we can select $Y \subseteq X$ with $d_H(Y, X) \leq \frac{1}{2}\epsilon$
to get ϵ -approximation.

Very similar to landmarking for witness cpx.

Here's a greedy procedure

input: pairwise distances on X : matrix D
tolerance ϵ

output set of indices $Y \subseteq X$

⋮
⋮
⋮

$i_0 = \underset{i}{\operatorname{argmax}} \{ \max_{x \in X} d(x_i, x) \}$ // find pt w/ highest
dc-centrality.

$d = D[\{i_0, \dots\}]$, $I \in \{i_0\}$ // distances to set I

while $\max_i d_i > \epsilon$:

→ $i \leftarrow \underset{i}{\operatorname{argmax}} d_i$ // furthest pt from set

$I \leftarrow I \cup \{i\}$ // add pt to set

for $j = 1:n$

$d_j \leftarrow \min \{ d_j, D[\{i, j\}] \}$ // update distances
to set

return I

let $Y = \{x_i \mid i \in I, x \in X\}$. by construction,

$$d_H(Y, X) \leq \epsilon \quad \alpha(|Y|^{k+1}) \text{ } k\text{-simplices}$$

Iterative subsampling:

n -point set X , total order on X (index order)

compute scales $\epsilon_i = d_H(X_i, X)$ $X_i = \{x_1, \dots, x_i\}$

$$\epsilon_1 \geq \epsilon_2 \geq \dots \geq \epsilon_n = 0$$

Choose parameters $\rho > \eta > 0$. Then

$$R(X_i; \eta \epsilon_i) \leq R(X_i; \rho \epsilon_i)$$

Chazal + Oudot '08: compute ranks

$$R_H(R(X_i; \eta \epsilon_i)) \rightarrow R(X; \rho \epsilon_i)$$

Corollary to thm on Rips inference:

Let $p > 8$, $2 < \eta \leq \frac{p}{4}$, suppose $K \subseteq \mathbb{R}^d$ cpl.

Let $d_{\text{R}}(X, K) = \epsilon \leq \frac{\eta - 2}{2p + \eta} \text{wfs}(K)$

then for any $l > m$ s.t.

$$\frac{2\epsilon}{\eta - 2} < \epsilon_l \leq \epsilon_m < \frac{\text{wfs}(K) - \epsilon}{p + 1}$$

$$\forall i \in [m, l], \text{rank } H_{\eta}(R(x_i; \eta\epsilon_i) \rightarrow R(x_i, \rho\epsilon_i)) \\ = \dim H_{\eta}(R(K)) \quad \forall \epsilon \in (0, \text{wfs}(K))$$

Let $m_i =$ doubling dimension of x_i at scale ϵ_i .

= smallest positive integer m s.t. any ball of radius ϵ_i in x_i can be covered by 2^m balls of radius $\epsilon_i/2$

\Rightarrow every x is connected to $2^{O(m_i)}$ other vertices

in Rips cplx $R(x_i; \epsilon_i)$

\Rightarrow # of k -simplices in Rips cplx $R(x_i; \rho\epsilon_i)$ is $2^{O(k m_i)}$

constant

"linear in size of vertex set"

note: m_i close to manifold dimension

$$X \subseteq \mathbb{R}^d, \quad m_i \leq d$$

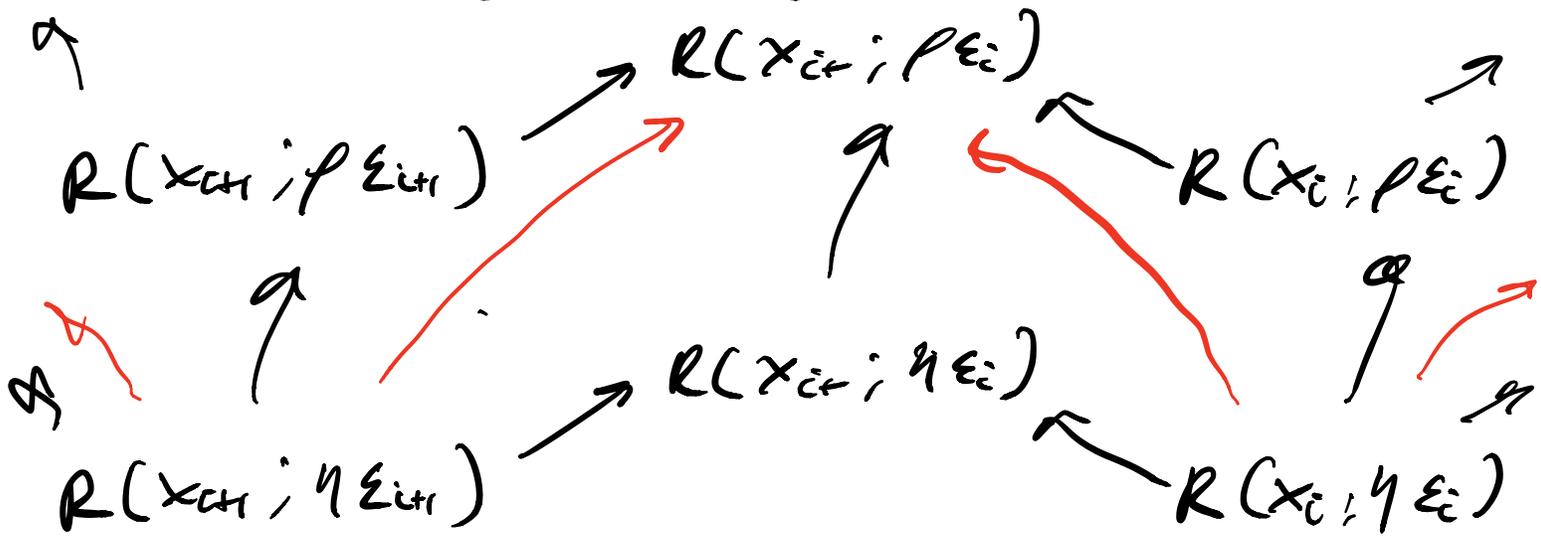
Now, we'd like to connect everything together.

$x_n \in \epsilon_n$, work backwards.

$$R(x_n; \rho \epsilon_n) \rightarrow R(x_n; \rho \epsilon_{n-1}) \leftarrow R(x_{n-1}; \rho \epsilon_{n-1})$$

This is called "Morozov zigzag" (decompos)

can stitch together diagram w/ both ρ, η scales



top & bottom rows. Morozov zigzags
w/ different multipliers μ -ZZ

vertical maps induce maps, can consider images.

image-Reps zigzag. iR -ZZ

can also consider oscillation from bottom to top (\Rightarrow)

called "oscillating Repr zigzag" oR -ZZ

want to understand stability of bars in zigzag,
but interleaving techniques don't play well w/
reversed arrows.

Lemma: can reverse arrows & keep barcode.

let $v_1 \rightleftarrows \dots \rightleftarrows v_k \xrightarrow{A_k} v_{k+1} \rightleftarrows \dots \rightleftarrows v_n$ be quiver rep.
 then $\exists A'_k$ s.t.

$v_1 \rightleftarrows \dots \rightleftarrows v_k \xleftarrow{A'_k} v_{k+1} \rightleftarrows \dots \rightleftarrows v_n$ has same barcode.

Pf: can construct A'_k from barcode form.

$A_k = B_{k+1} E_k B_k^{-1}$. In barcode form, we can take transpose of E_k & reverse arrow
 $A'_k = B_k E_k^T B_{k+1}^{-1}$ (note: not A_k^T generally)

Lemma: we can replace $v_k \xrightarrow{A_k} v_{k+1} \xrightarrow{A_{k+1}} v_{k+2}$
 with $v_k \xrightarrow{A_{k+1} A_k} v_{k+2}$. Simply remove index $k+1$

$$I[k+1, k+1] \mapsto \emptyset$$

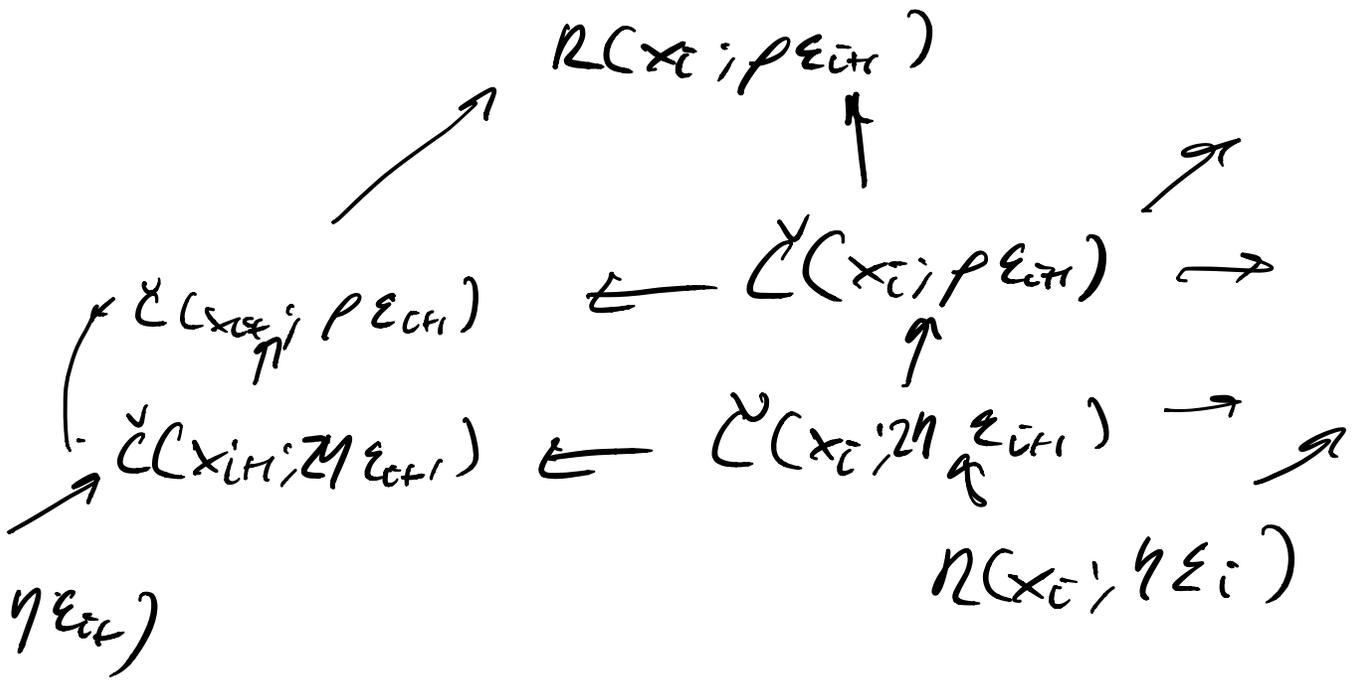
$$I[b, k+1] \mapsto I[b, k]$$

$$I[k+1, d] \mapsto I[k+2, d]$$

all other intervals unchanged.

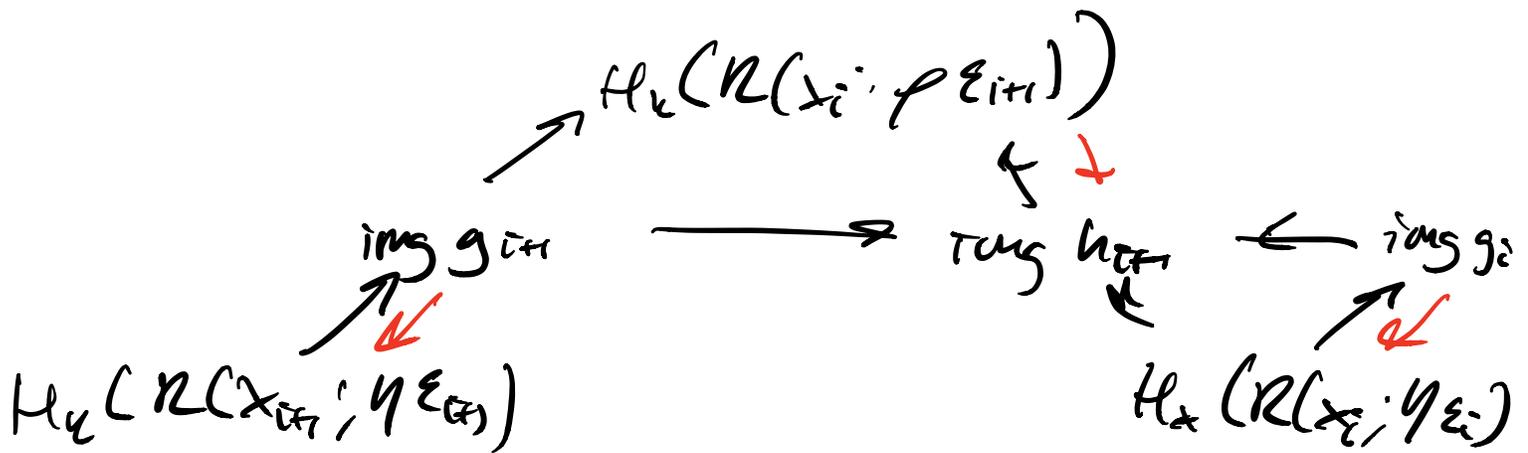
Pf: trivial.

Application to Rips zigzags. Assume $\eta < \frac{\epsilon}{2}$,
 then we can factor inclusions through $\mathcal{C}_{\epsilon, \eta}$

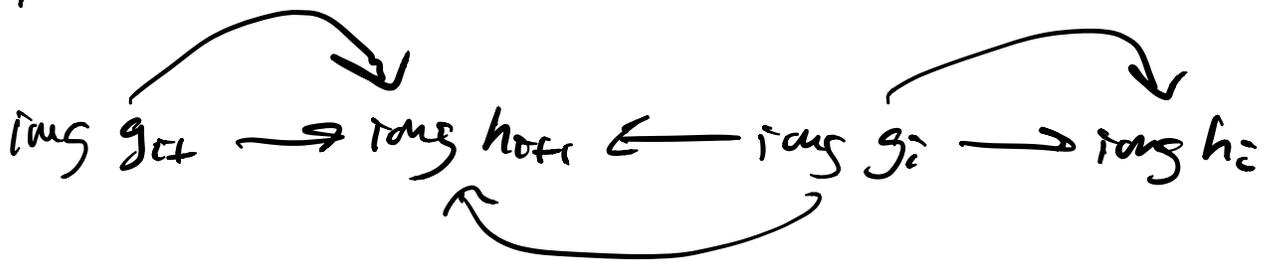


All arrows are inclusions, diagram commutes

Apply homology, take: $g_j = H_x(\check{C}(X_j; \eta \varepsilon_j) \rightarrow \check{C}(X_j; \rho \varepsilon_j))$
 $h_j = H_x(\check{C}(X_{jt}; \eta \varepsilon_{jt}) \rightarrow \check{C}(X_{jt}; \rho \varepsilon_{jt}))$



apply arrow reversal to get red arrows,



Thm: Oudst/Sheehy 2015:

let $p > 0$, $3 \leq \eta = \frac{p-4}{2}$, $K \subseteq \mathbb{R}^d$ compact

$$d_K(x, K) \leq \varepsilon,$$

$$\varepsilon \leq \min \left(f(\eta, p) \right) \text{ wfs}(K)$$

\Rightarrow for any $l > k$ w/

$$\varepsilon_l > \max \left\{ \frac{3\varepsilon}{\eta-3}, \frac{4\varepsilon}{p-2\eta-1} \right\}$$

$$\varepsilon_k < \min \left\{ \frac{1}{6} \text{wfs}(K) - \varepsilon, \frac{1}{p-1} (\text{wfs}(K) - \varepsilon) \right\}$$

The barcode of $\mathbb{R}-\mathbb{Z}\mathbb{Z}$ restricted $[k, l]$
has 2 types of intervals:

- 1) those that span $[k, l]$, encode $H_*(K)$
- 2) remaining intervals have length 0

\Rightarrow barcode has same range that agrees with
 $H_*(K)$

this holds for $\mathbb{R}-\mathbb{Z}\mathbb{Z}$, $i\mathbb{R}-\mathbb{Z}\mathbb{Z}$, but no
guarantees for $M-\mathbb{Z}\mathbb{Z}$, need stack in p. 4
to kill spurious homology

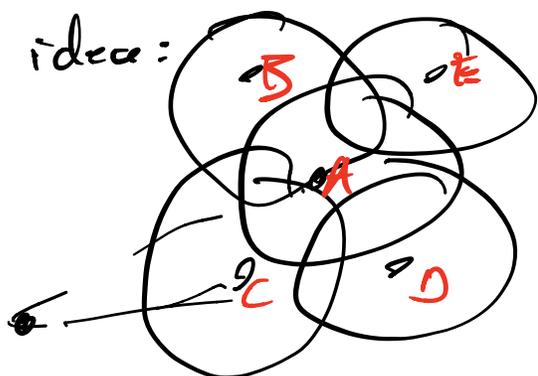
can drastically cut down on # simplices.

$\sum \binom{d(n_i)}{2}$ simplices in each space, n spaces

(x_1, \dots, x_n)
 \Rightarrow linear # simplices in n .

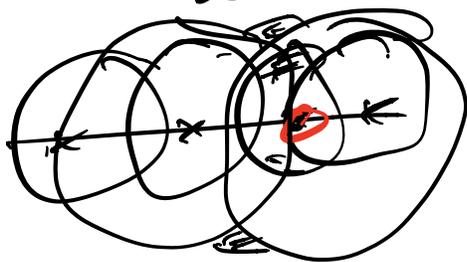
Part 2: Linear-size Rips approximations introduced by Sheehy 2013.

want something that has $O(n)$ simplices, but is filtration, not zigzag.



at some parameter, ball around A will be covered by other balls.
 \rightarrow safely remove it from filtration.

Problem: sometimes balls are not entirely covered



so we can perturb metric slightly to remove ball.

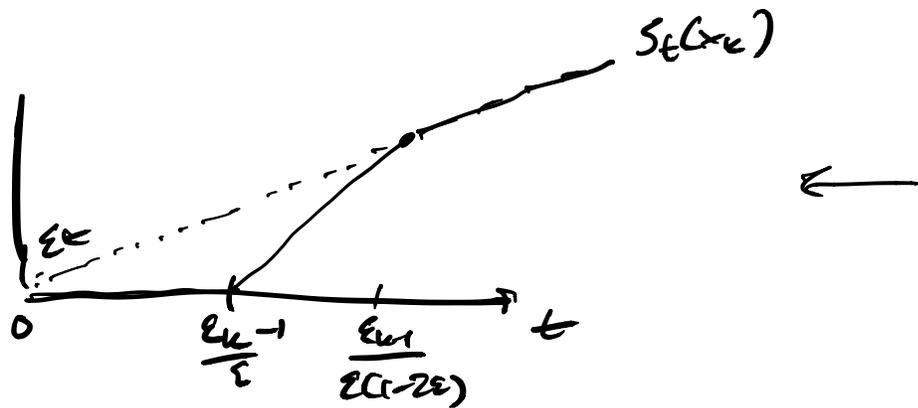
Again, let $X_i = \{x_1, \dots, x_i\}$, $\epsilon_i = d_H(X_i, X)$

given target approx. error $\epsilon \in (0, \frac{1}{2})$

perturb metric d_X by incorporating weights

$$d_X^t(x_i, x_j) = d_X(x_i, x_j) + S_t(x_i) + S_t(x_j)$$

$$S_t(x_i) = \begin{cases} 0 & \text{if } t \leq \frac{\epsilon_{i-1}}{2} \\ \frac{1}{2} \left(t - \frac{\epsilon_{i-1}}{2} \right) & \text{if } \frac{\epsilon_{i-1}}{2} \leq t \leq \frac{\epsilon_{i-1}}{2(1-2\epsilon)} \\ \epsilon & \text{if } t \geq \frac{\epsilon_{i-1}}{2(1-2\epsilon)} \end{cases}$$



metric balls are smaller in perturbed metric.

to simulate ball growth & removal, rely on sparse Rips complexes.

$$S(X, d_X, \epsilon; t) = \mathcal{R}(X_i, d_X) \cup \bigcup_{j=0}^n (X_j, d_X, \rho_{j-1}, \rho_j)$$

for $t \in (\rho_{\epsilon_i}, \rho_{\epsilon_{i-1}}]$, $\rho = \frac{t}{\epsilon(1-2\epsilon)}$

stops growth of balls at time they should be removed from union instead of removing them

Thus suppose (X, d_X) has doubling dimension m , and order on X obtained by furthest pt. sampling. Then $\forall k \geq 0$ # of k -simplices in $S(X, d_X, \epsilon)$ is at most $\left(\frac{1}{\epsilon}\right)^{O(km)}$ n .

linear in n .

There is an $O(\epsilon)$ log-scale interleaving btw $S(X, d_X, \epsilon)$ and $\mathcal{R}(X, d_X)$

$$\begin{array}{ccc}
 R(x, d_x; t) & \longrightarrow & R(x, d_x; t(1+2\varepsilon)) \\
 \uparrow & \searrow & \uparrow \\
 S(x, d_x, \varepsilon; t) & \longrightarrow & S(x, d_x, \varepsilon; t(1+2\varepsilon))
 \end{array}$$

Black arrows are all inclusions

red arrow is simplicial map extended from projection onto x_c .

$$\pi_t(x_k) = \begin{cases} x_k & \text{if } k \in i \text{ (} x_k \in x_c \text{)} \\ \text{argmin}_{l \in i} d_x^t(x_l, x_k) & \text{otherwise} \end{cases}$$

$$d(x_i, x_k) = t \Rightarrow d(x_i, x_j) \leq t \quad \forall i, j \in 0..k$$

$$\Leftrightarrow d^t(\pi_t x_i, \pi_t x_j) \leq t(1+2\varepsilon)$$

red arrow exist.

Composition of maps commutes up to homotopy.

$$\begin{array}{ccc}
 R(x, t) & \longrightarrow & R(x; t(1+2\varepsilon)) \\
 & \searrow & \uparrow \\
 & & S(x, \varepsilon; t(1+2\varepsilon))
 \end{array}$$

want to show $(x_0 \dots x_k) \mapsto (\pi_\epsilon x_0 \dots \pi_\epsilon x_k) \cong \text{id}$

note that simplex $(x_0 \dots x_k, \pi_\epsilon x_0 \dots \pi_\epsilon x_k)$

in $\mathbb{R}^n(x; \epsilon(1+2\epsilon))$

b/c projection map distorts by $\leq \epsilon$

\Rightarrow get maximum distance of $d(x_0 \dots x_k) + 2\epsilon$
by δ -ineq.

have k path through simplex $(x_0 \dots x_k, \pi_\epsilon x_0 \dots \pi_\epsilon x_k)$

Thm: bottleneck distance btw. log scale
persistence diagrams is $\leq \log_2(1+2\epsilon) = O(\epsilon)$

□

Discussion:

- Iterative subsampling procedure is $O(n^2)$
can approximate in $O(n)$ time w/ net trees.
 - use \mathbb{Z}_2 approach for inference.
 - if we care abt. Rips signature, then
sparse filtrations are better
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