

w interleaving

# Stability of Persistence

- ① Cont's filtrations, + 10 mins
  - ② P-modules up to interleaving, + 5 mins
  - ③  $d_H, d_{an}$
- Next: Bottleneck distance.

## ① Continuous filtrations

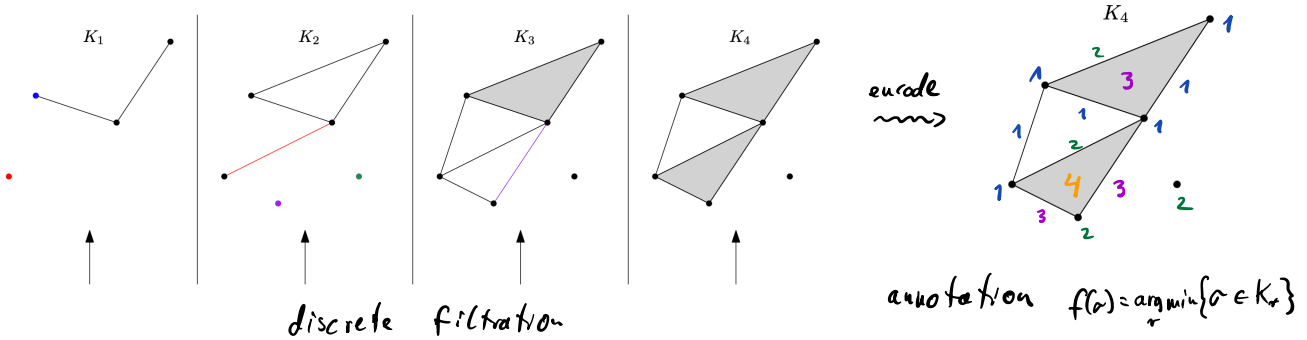
Def: A continuous filtration of a scc  $K$  is a collection of subcomplexes  $\{K_r\}_{r \geq 0}$  of  $K$ :

$$\forall r < s: K_r \subseteq K_s.$$

! To make life easier we assume:  $\forall \sigma \in K \arg \min \{\sigma \in K_r\}$  exists.

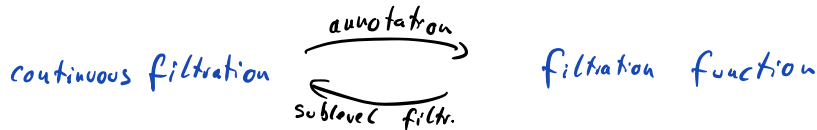
Examples: Rips, Čech, Alpha, ...

It appears AT some scale, not Beyond some scale.  
(closed notation @ defs of axes)



Def: Filtration function for scc  $K$ : map  $K \rightarrow [0, \infty)$  such that  $\tau \in \sigma \Rightarrow f(\tau) \leq f(\sigma)$   
 $\sigma \mapsto f(\sigma)$

Sublevel filtration of  $f$ :  $\{f^{-1}([0, r])\}_{r \geq 0} \leftarrow$  continuous filtration



Example: Filtration functions: Rips  $\rightsquigarrow$  diam

Čech  $\rightsquigarrow$  radius of smallest enclosing ball.

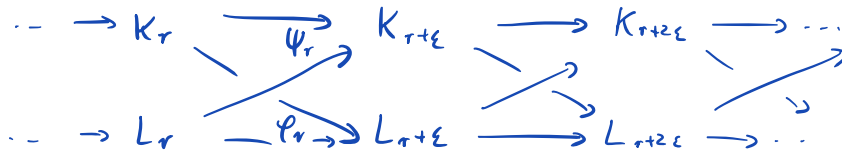
! Computation of Persistence is unhindered as it only requires the  
finitely many annotation values.

Example: [offsets]  $SCID^n$  finite

$\{\check{C}ech(S, r)\}_{r \geq 0} \simeq \{N(S, r)\}_{r \geq 0}$  models growth of offsets. The changes  
Nerve THM in homology can be computed & tracked by pers. hom.

Def:  $\epsilon \geq 0$ . Filtrations  $\{K_r\}_{r \geq 0}$  and  $\{L_r\}_{r \geq 0}$  are  $\epsilon$ -interleaved if

$\exists$  simplicial maps  $\Psi_r: K_r \rightarrow L_{r+\epsilon}$   $\Psi_r: L_r \rightarrow K_{r+\epsilon}$  such that the following diagram commutes:



Isomorphism = 0-interleaving.

Interleaving distance:  $\arg \inf_{\epsilon} \{ \{K_r\}_{r \geq 0} \text{ and } \{L_r\}_{r \geq 0} \text{ } \epsilon\text{-interleaved} \}$  ← actually argmin in our case

max. distance between filtration functions  $f, g$  on  $K$ :

$$\|f-g\|_{\infty} = \max_{\sigma \in K} |f(\sigma) - g(\sigma)|$$

↑ metric on equiv. classes of filtrations

Proposition: Sublevel filtrations of  $f$  and  $g$  are  $\|f-g\|_{\infty}$ -interleaved.

← for example cubical or of images

Proof:  $\Psi_r$  &  $\Psi_r$  are identities on vertices.

observe:  $\forall \sigma \in K: f(\sigma) \leq r \Rightarrow g(\sigma) \leq r+\epsilon$  & vice versa

This concludes the proof. □

Also recall:  $X = \{x_1, \dots, x_n\}, Y = \{y_1, \dots, y_n\}, d(x_i, y_i) \leq \epsilon$ . Then:

- ① Rips filtrations of  $X$  &  $Y$  are  $2\epsilon$ -interleaved
- ② Čech filtrations of  $X$  &  $Y$  are  $\epsilon$ -interleaved.

② Persistence modules: fix  $f$  ← obtained from filtrations by applying homology

Def: A persistence module is a collection of finite-dimensional vector spaces  $\{V_r\}_{r \geq 0}$

along with linear maps  $h_{r,s}: V_r \rightarrow V_s, \forall r \leq s$ , such that

$$h_{r,r} = \text{id}_{V_r} \text{ and } h_{g,s} \circ h_{r,g} = h_{r,s} \quad \forall r \leq g \leq s.$$

Scale  $r$  is **regular** if  $\exists \epsilon > 0; h_{g,s}$  is an isomorphism,  $\forall g \in [r-\epsilon, r+\epsilon)$ .

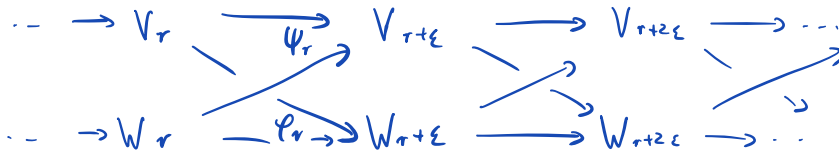
Scale  $r$  is **critical** if it is not regular.

Additional assumptions: • Eventually all  $h_{r,s}$  are isomorphisms:  $\exists R: \forall g \geq r \geq R, h_{r,g}$  is isomorphism } hold for homology of our filtrations

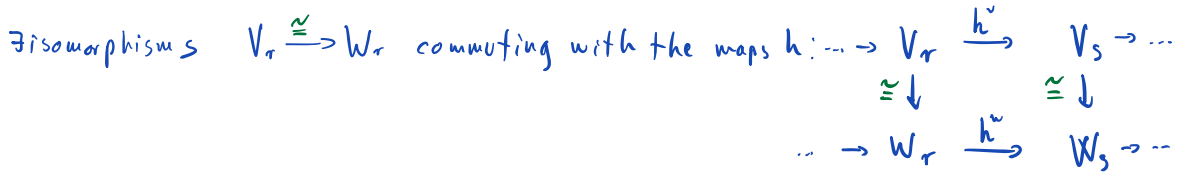
- $\forall r \geq 0 \exists r' > 0; \forall g \in [r, r') h_{r,g}$  is isomorphism
- There are only finitely many critical scales

Def:  $\epsilon > 0$ . Persis. modules  $\{V_r\}_{r \geq 0}$  and  $\{W_r\}_{r \geq 0}$  are  $\epsilon$ -interleaved if

$\exists$  linear maps  $\varphi_r: V_r \rightarrow W_{r+\epsilon}$   $\psi_r: W_r \rightarrow V_{r+\epsilon}$  such that the following diagram commutes



Def: Persistence modules  $\{V_r\}_{r \geq 0}$  and  $\{W_r\}_{r \geq 0}$  are isomorphic, if they are 0-interleaved!



Interleaving distance:  $\arg \inf_{\epsilon} \{ \{V_r\}_{r \geq 0} \text{ and } \{W_r\}_{r \geq 0} \text{ } \epsilon\text{-interleaved} \}$  ← actually argmin in our case

metric on isom. classes of pers. modules

Observations:  $\epsilon$ -interleaved filtrations  $\xrightarrow[\text{H}_2]{\text{induce}}$   $\epsilon$ -interleaved p. modules.

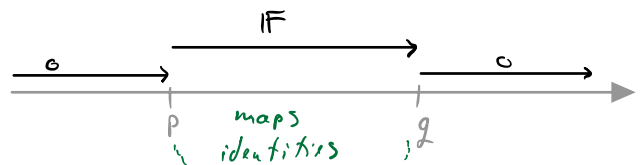
Barcodes are "horizontal decomposition" of pers. modules. Let us formalize it:

Def: The direct sum of persistence modules  $\{V_r\}_{r \geq 0}$  and  $\{W_r\}_{r \geq 0}$  is a persist. module  $\{V_r \oplus W_r\}_{r \geq 0}$  with bonding maps  $h^v \oplus h^w$ .

Def:  $0 \leq p < q$ . Interval module  $\mathbb{F}_{[p,q]}$  is a persist. module  $\{V_r\}_{r \geq 0}$  with

$$V_r = \mathbb{F} \text{ if } r \in [p, q]$$

$$V_r = 0 \text{ if } r \notin [p, q]$$



bonding maps are isomorphisms whenever possible

Theorem: [Structure theorem for persistent homology] Each persistence module is isomorphic to a (finite) direct sum of interval modules. The decomposition is unique up to permutation of the intervals.

Barcode decomposition formally

Example: interleaving distance between interval modules =  $d_B$  between pos.

Are there interleavings on p. modules not arising from interleave. of filtrations?  
Yes! By "similar spaces". Let's explain it.

Def:  $X$  a metric space,  $A, B \subseteq X$  finite subspaces.

Hausdorff distance  $d_X(A, B) = \max \left\{ \max_{a \in A} \min_{b \in B} d(a, b), \max_{b \in B} \min_{a \in A} d(a, b) \right\}$

is a metric on finite subsets of  $X$ .

Def:  $A, B$  finite metric spaces.

Gromov-Hausdorff distance

$$d_{GH}(A, B) = \inf_{\nu, \mu} d_H(\nu(A), \mu(B))$$

is a metric on isometry classes of finite metric spaces.

$\nu: A \hookrightarrow X, \mu: B \hookrightarrow X$  isometric embeddings.

inf always attained.

Proposition:  $d_{GH} \leq d_H$

Proposition:  $\epsilon > 0; A, B$  finite metric spaces,  $d_{GH}(A, B) \leq \epsilon, g \in \{0, 1, \dots\}$ . Then:

(a)  $\{H_g(\text{Rips}(A, r); \mathbb{F})\}_{r \geq 0}$  and  $\{H_g(\text{Rips}(B, r); \mathbb{F})\}_{r \geq 0}$  are  $2\epsilon$ -interleaved.

(b)  $\{H_g(\bar{\text{Cech}}(A, r); \mathbb{F})\}_{r \geq 0}$  and  $\{H_g(\bar{\text{Cech}}(B, r); \mathbb{F})\}_{r \geq 0}$  are  $\epsilon$ -interleaved.

Proof: [for Rips,  $g=1$ ]

$$\begin{array}{ccc} H_1(\text{Rips}(A, r); \mathbb{F}) & \xrightarrow{\varphi_r} & H_1(\text{Rips}(B, r+2\epsilon); \mathbb{F}) \\ & \searrow \psi_r & \\ H_1(\text{Rips}(B, r); \mathbb{F}) & \xrightarrow{\psi_r} & H_1(\text{Rips}(A, r+2\epsilon); \mathbb{F}) \end{array}$$

Assume  $A, B \subseteq X$  by def.

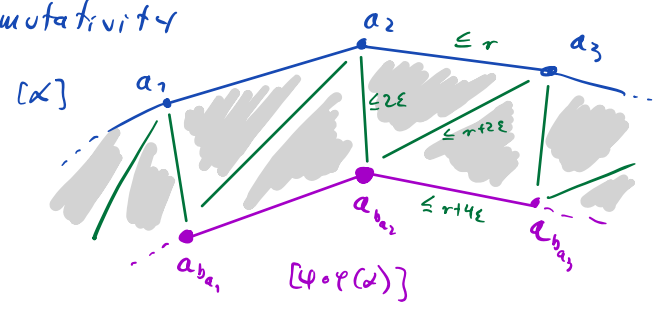
$\forall a \in A$  choose  $b_a \in B: d(a, b_a) \leq \epsilon. \forall b \in B$  choose  $a_b \in A: d(a, b) \leq \epsilon$

Take cycle  $\alpha = \langle a_1, a_2 \rangle + \langle a_2, a_3 \rangle + \dots + \langle a_k, a_1 \rangle$

Def  $\varphi_r(\alpha) = [\langle b_{a_1}, b_{a_2} \rangle + \dots + \langle b_{a_k}, b_{a_1} \rangle]$ .  $\leftarrow$  turns out to be a well-defined lin. map.

$$\varphi_r(\langle b_1, b_2 \rangle + \dots + \langle b_m, b_1 \rangle) = [\langle a_{b_1}, a_{b_2} \rangle + \dots + \langle a_{b_m}, a_{b_1} \rangle]$$

Commutativity



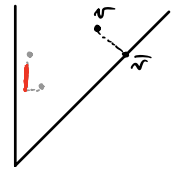
all sides  $\leq r+2\epsilon$   
 $\leftarrow$  homology between  $\alpha$  &  $\varphi_r(\alpha)$



③ Bottleneck distance: a metric on persistence diagrams.

For  $(x_1, y_1), (x_2, y_2) \in \mathbb{R}^2$  define:  $d_{\infty}((x_1, y_1), (x_2, y_2)) = \max\{|x_1 - x_2|, |y_1 - y_2|\}$

$$\overline{(x, y)} = \left(\frac{x+y}{2}, \frac{x+y}{2}\right)$$



Assume  $\mathcal{A} = (a_1, a_2, \dots, a_m)$  and  $\mathcal{B} = (b_1, b_2, \dots, b_n)$  are persistence diagrams.

$a_i, b_j$  pts in  $\{(x, y) \in \mathbb{R}^2; y > x \geq 0\}$ , appearing possibly with repetitions in  $\mathcal{A}$  &  $\mathcal{B}$ .

Def: A partial matching between  $\mathcal{A}$  and  $\mathcal{B}$  is a bijective map

$$\varphi: \mathcal{A}' \rightarrow \mathcal{B}' \quad \text{for some } \mathcal{A}' \subseteq \mathcal{A}, \mathcal{B}' \subseteq \mathcal{B}.$$

The matching distance of  $\varphi$  is

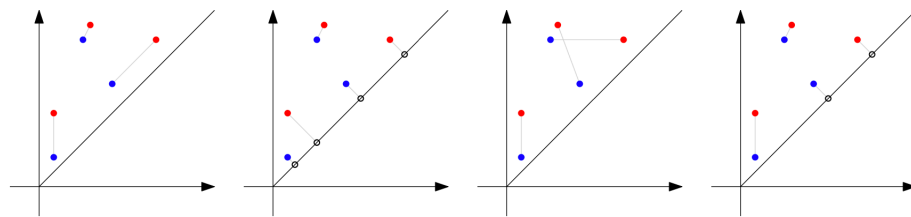
$$d_{\infty}(\varphi) = \max \left\{ \max_{v \in \mathcal{A}'} \{d_{\infty}(v, \varphi(v))\}, \max_{v \in \mathcal{A} \setminus \mathcal{A}'} \{d_{\infty}(v, \bar{v})\}, \max_{v \in \mathcal{B} \setminus \mathcal{B}'} \{d_{\infty}(v, \bar{v})\} \right\}$$

Let  $\mu(\mathcal{A}, \mathcal{B})$  be the collection of partial matchings  $\mathcal{A}$  to  $\mathcal{B}$ .

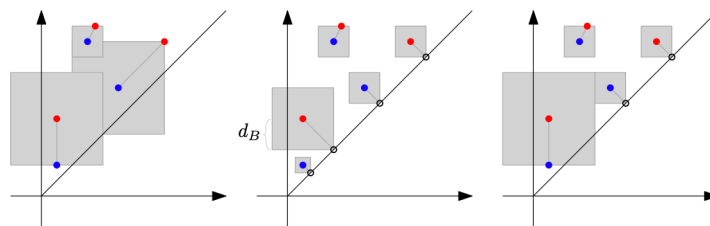
The Bottleneck distance is defined as

$$d_{\mathcal{B}}(\mathcal{A}, \mathcal{B}) = \min_{\varphi \in \mu(\mathcal{A}, \mathcal{B})} d_{\infty}(\varphi)$$

partial matchings



Bottleneck distance

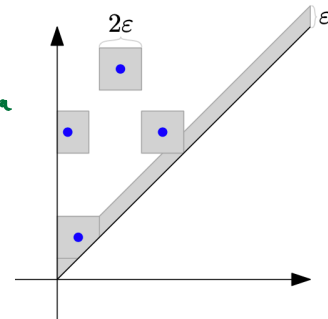


Isometry Theorem: The interleaving distance } algorithm  
between persistence modules

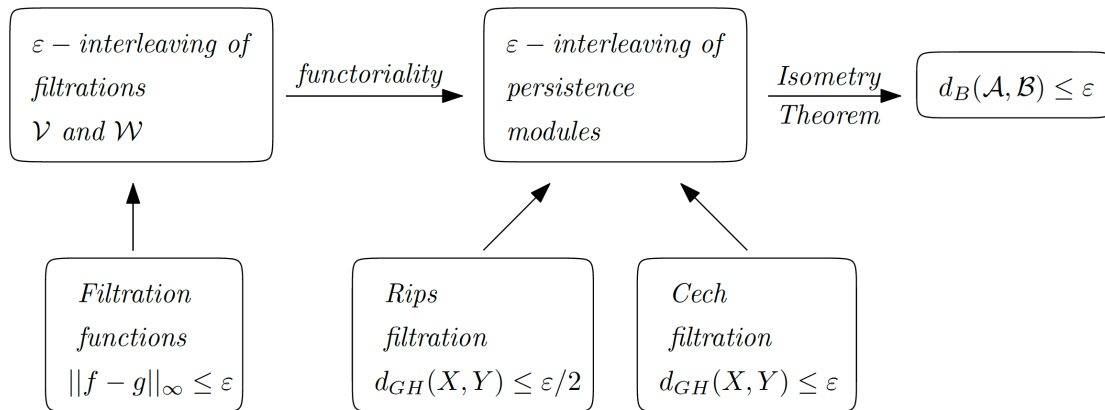
**EQUALS**

visualisation

the bottleneck distance between persistence diagrams.



# // Stability Theorem:



Alternative metric (sensitive to more changes):

Wasserstein distance:

let  $p > 1$

The  $p$ -matching distance of  $\varphi$  is

$$d_m^p(\varphi) = \sum_{v \in \mathcal{A}'} \{d_{\infty}(v, \varphi(v))\} + \sum_{v \in \mathcal{A} \setminus \mathcal{A}'} \{d_{\infty}(v, \bar{r})\} + \sum_{v \in \mathcal{B} \setminus \mathcal{B}'} d_{\infty}(v, \bar{r})$$

The  $p$ -Wasserstein distance is defined as

$$W_p(\mathcal{A}, \mathcal{B}) = \min_{\varphi \in \mu(\mathcal{A}, \mathcal{B})} d_m^p(\varphi)$$

$W_p$  is also stable under certain conditions.

!!critical axes not stable!!