Topological Data Analysis

2022-2023

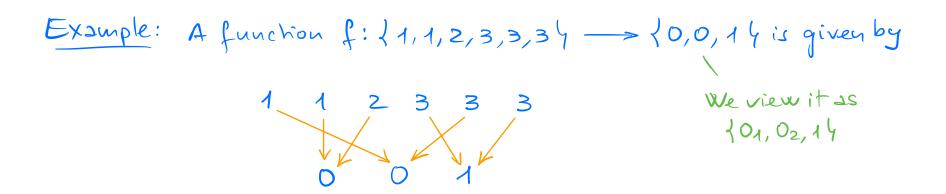
Lecture 8

Bottleneck Distance

28 November 2022

Suppose given persistence modules V and V' of finite type with respective persistence diagrams D and D'. Suppose, in addition, that $\dim V_\infty = \dim V_\infty'.$

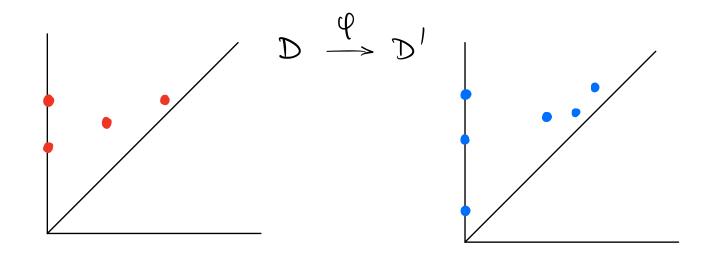
Recall that persistence diagrams are <u>multisets</u>, which means that their points may have a multiplicity. If we choose an <u>order</u> within each multiple point, then we can treat multisets as ordinary sets.

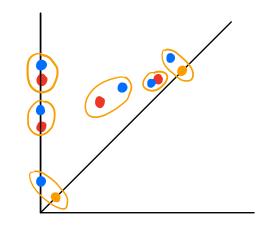


Convention: If Δ denotes the diagonal b=d, then points in Δ are members of every persistence diagram with countably infinite multiplicity.

A matching between D and D' is a bijective function $\varphi: D \to D'$ such that, for every $(x,x) \in \Delta$, either $\varphi(x,x) = (x,x)$ or $\varphi(x,x) = (b,d)$ with $b \neq d$.

Example:





A matching between D and D' For each matching $\varphi: D \to D'$, define $\|\varphi\| = \max \{ d_{\infty}((x,y), \varphi(x,y)) \mid (x,y) \in D \}$ where d_{∞} is the l_{∞} -distance on \mathbb{R}^2 , namely $d_{\infty}((x,y), (x',y'))) = \max \{ |x-x'|, |y-y'| \}'$

The bottleneck distance between two persistence diagrams is defined as $W_{\infty}(D,D') = \min \{\|\phi\| \mid \phi: D \to D' \text{ matching } \phi.$

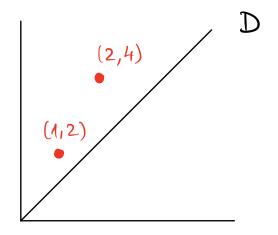
Hence $W_{\infty}(D,D')$ is the smallest $E\geqslant 0$ for which there exists a matching $\varphi:D\to D'$ for which $d_{\infty}((x,y),\varphi(x,y)) \leq E$ for all $(x,y)\in D$.

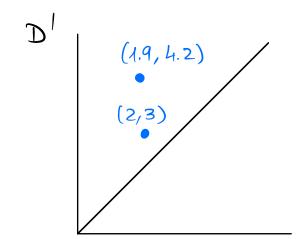
More generally, the Wasserstein distances are defined for P, 9>1 as

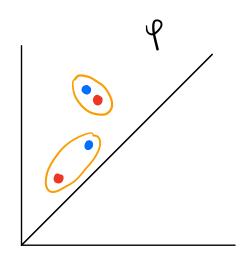
$$W_{P}[q](D,D') = \min_{\varphi:D\to D'} \left(\sum_{(x,y)\in D} d_{q}((x,y),\varphi(x,y))^{P} \right)^{1/P}$$

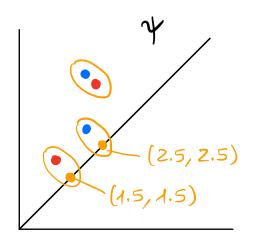
where $d_{q}((x,y),(x,y')) = (|x-x'|^{q} + |y-y'|^{q})^{1/q}$.

Example:









$$\| \varphi \| = \max \{1.0, 0.2 \} = 1$$

$$\|\gamma\| = \max\{0.5, 0.5, 0.2\} = 0.5$$

$$W_{\infty}(D,D')=0.5$$

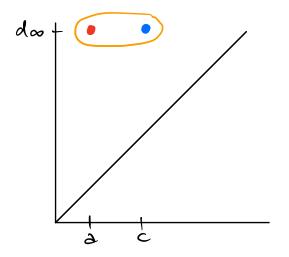
Isometry Theorem:

If V and V' are persistence modules of finite type with din $V_{\infty} = \dim V'_{\infty}$ and D(V), D(V') denote their respective persistence diagrams, then

$$W_{\infty}(D(V),D(V'))=d_{int}(V,V').$$

Proof:

 $\sqrt{1} = \mathbb{F}[a, \infty)$ $\sqrt{1} = \mathbb{F}[c, \infty)$



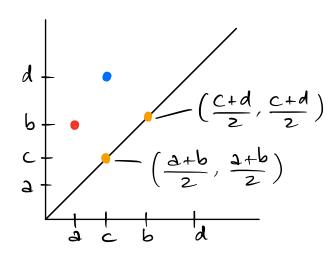
$$W_{\infty}(D(V),D(V')) = |a-c| =$$

$$= d_{int}(V,V').$$

$$V = \mathbb{F}[a,b)$$

$$V' = \mathbb{F}[c,d)$$

First case: a < c < b < d

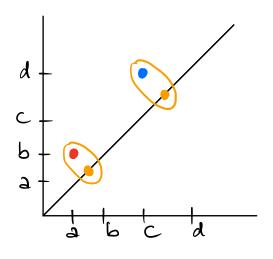


$$d_{\infty}((a,b),(c,d)) = \max \{c-a,d-b\}$$

$$d_{\infty}\left((a,b),\left(\frac{a+b}{2},\frac{a+b}{2}\right)\right) = \frac{a+b}{2} - a = \frac{b-a}{2}$$

$$W_{\infty}(D(V),D(V')) = \min \left\{ \max \left\{ c-a, d-b \right\}, \max \left\{ \frac{b-a}{2}, \frac{d-c}{2} \right\} \right\} = d_{int}(V,V')$$

Second case: a < b < c < d



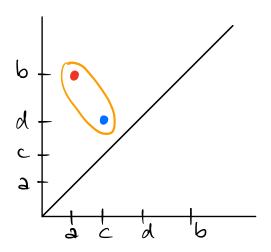
Note that $\int d-b \ge d-c > \frac{1}{2}(d-c)$ $|c-a \ge b-a > \frac{1}{2}(b-a)$

Hence $d_{\infty}((a,b),(c,d))$ is larger than $||\phi||$ if ϕ is the matching with diagonal points.

$$W_{\infty}(D(V), D(V') = \max \left\{ \frac{b-a}{2}, \frac{d-c}{2} \right\} = d_{int}(V, V').$$

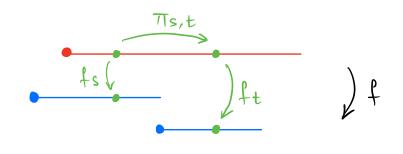
Third case:

 $a \le c < d \le b$



 $W_{\infty}(D(V),D(V')) = \max\{c-a,b-dy=dint(V,V').$

For the general case, note that V and V' are δ -interleaved if and only if their intervals IF [a,b) that are not δ -short can be pairwise matched in such a way that each pair of matched intervals are δ -interleaved.

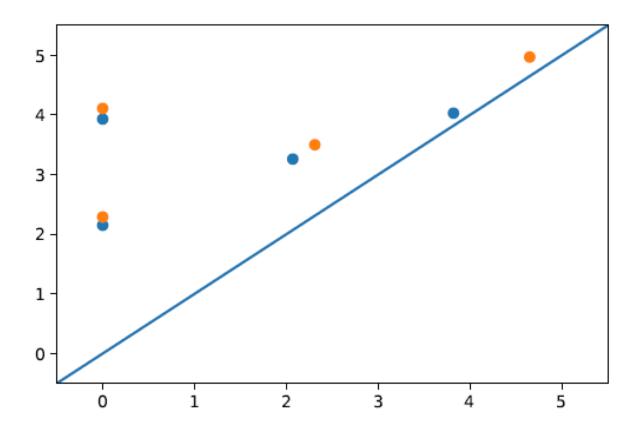


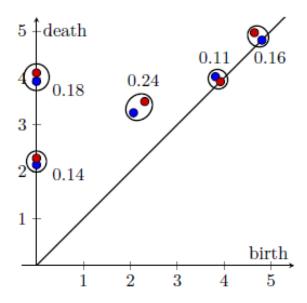
For this, we use the fact that every morphism between persistence modules must send intervals to intervals.

This cannot occur

 ${m V} = \mathbb{F}[0,2.15) \oplus \mathbb{F}[0,3.93) \oplus \mathbb{F}[2.07,3.26) \oplus \mathbb{F}[3.82,4.03)$

 $V' = \mathbb{F}[0, 2.29) \oplus \mathbb{F}[0, 4.11) \oplus \mathbb{F}[2.31, 3.50) \oplus \mathbb{F}[4.65, 4.97)$





$$d_{\text{int}}(V, V') = W_{\infty}(D(V), D(V')) = 0.24$$

```
R Console

> library("TDA")
> Diagl <- matrix(c(0, 0, 2.15, 0, 0, 3.93, 0, 2.07, 3.26, 0, 3.82, 4.03), ncol = 3, byrow = TRUE)
> Diag2 <- matrix(c(0, 0, 2.29, 0, 0, 4.11, 0, 2.31, 3.50, 0, 4.65, 4.97), ncol = 3, byrow = TRUE)
> par(mfrow = c(1,2))
> plot.diagram(Diag1, diagLim=c(0,6))
> plot.diagram(Diag2, diagLim=c(0,6))
> bottleneckDist <- bottleneck(Diag1, Diag2, dimension = c(0, 1))
> print(bottleneckDist)
[1] 0.24
> |
```