Persistent Homology: Interpretation and Stability

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- Notice that our ultimate goal is to use persistent homology to infer the "shape" of data (i.e., homology inference)
- To do this, we need to fully understand the meanings of PD or barcode, from different aspects

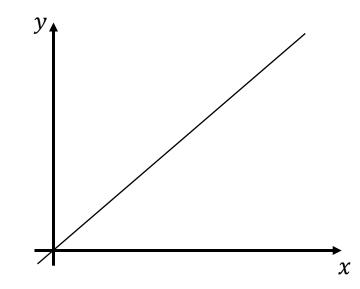
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- To do this, we need to fully understand the meanings of PD or barcode, from different aspects
- Moreover, we shall also see some properties that are critical to showing that persistent homology is a "reliable" way for inferring the shape

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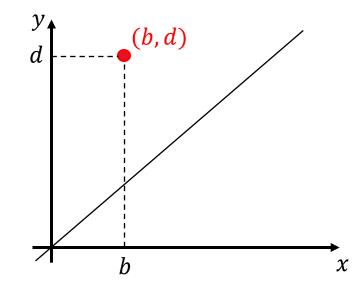
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- Why above the diagonal?

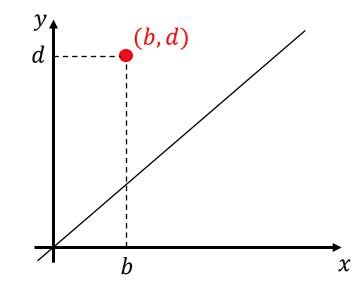
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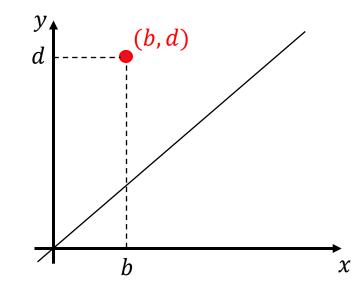
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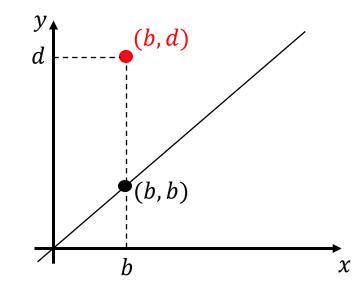
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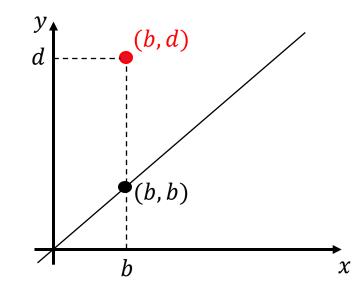
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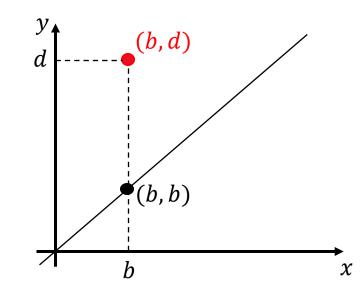
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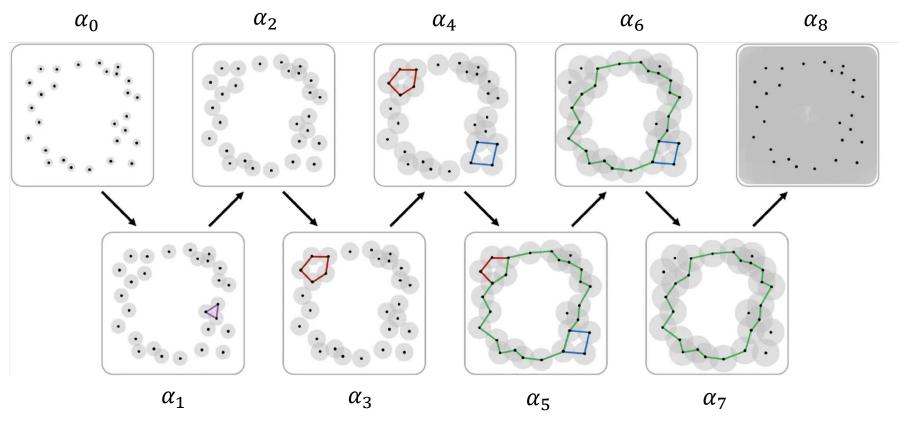
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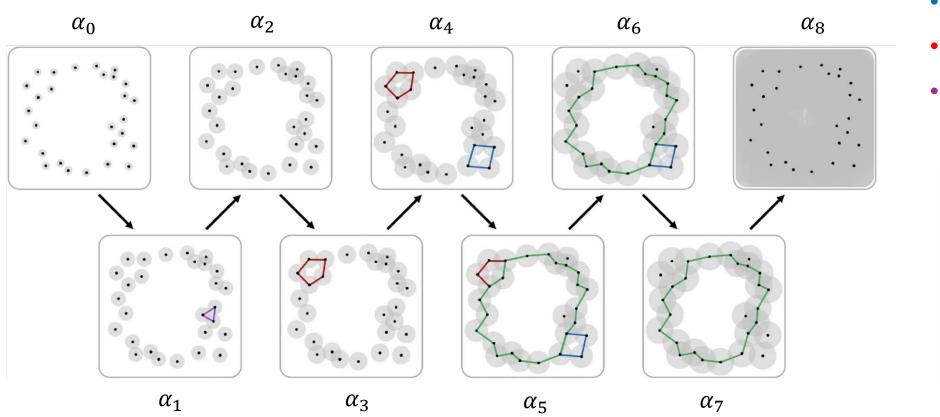
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- So (b, d) is above the diagonal



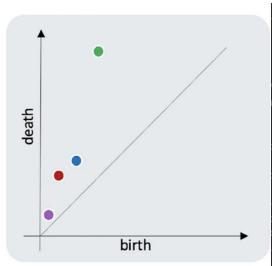
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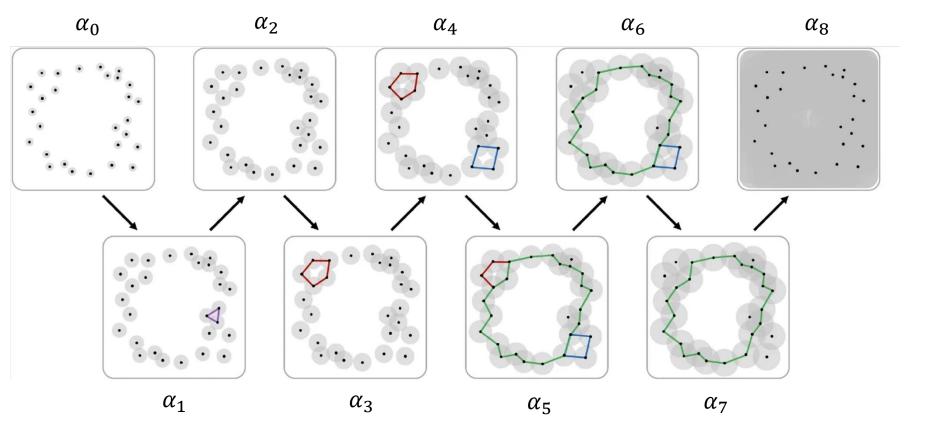


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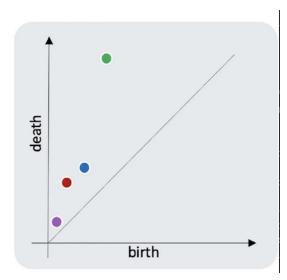


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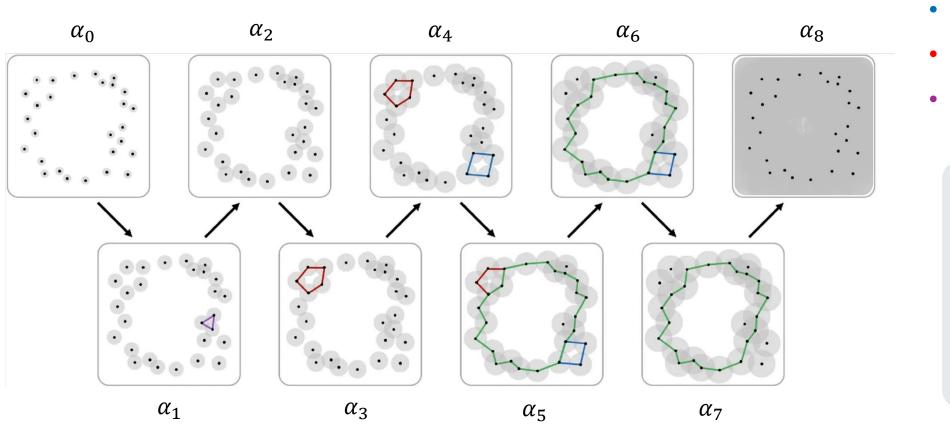


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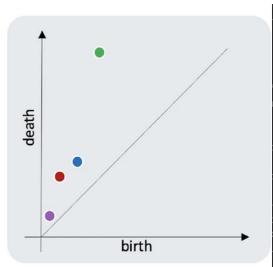


Meaning of "length" of a PD interval

- Define length of a PD interval (point), (b, d), as d b
- Observe the longer an interval is, the more significant its homology hole is

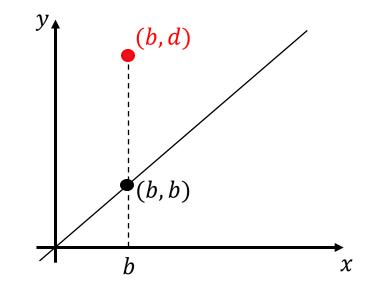


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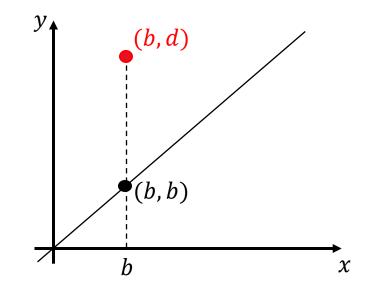
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- We observe that the length d b of a PD point (b, d) equals $1/\sqrt{2}$ times the distance of (b, d) to the diagonal
- This means that the distance of a point in PD to the diagonal indicates the length of the corresponding interval, and hence the significance of the homological feature



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- Mathematically, the tool to measure the difference of two objects is called a distance function (or metric)

Distance function (as on Wikipedia)

Definition [edit]

Formally, a metric space is an ordered pair (M, d) where M is a set and d is a metric on M, i.e., a function

 $d:M imes M o \mathbb{R}$

satisfying the following axioms for all points $x,y,z\in M$:^{[4][5]}

1. The distance from a point to itself is zero:

d(x,x)=0

2. (Positivity) The distance between two distinct points is always positive:

 $\text{If } x \neq y, \text{then } d(x,y) > 0$

3. (Symmetry) The distance from x to y is always the same as the distance from y to x:

d(x,y)=d(y,x)

4. The triangle inequality holds:

 $d(x,z) \leq d(x,y) + d(y,z)$

This is a natural property of both physical and metaphorical notions of distance: you can arrive at z from x by taking a detour through y, but this will not make your journey any shorter than the direct path.

If the metric d is unambiguous, one often refers by abuse of notation to "the metric space M".

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- L_{∞} -distance: $d_{\infty}(x, y) = \max\{|x_1 y_1|, |x_2 y_2|\}$

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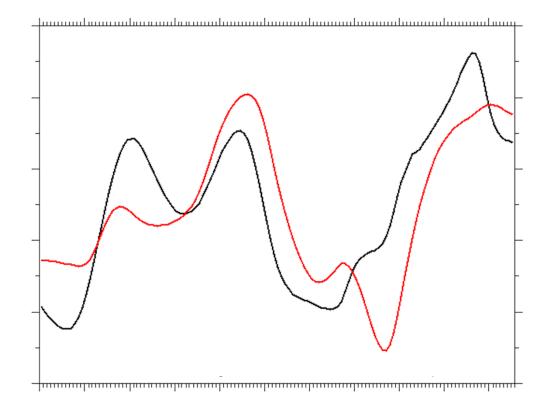
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- What is the data we try to measure here?
- It turns out that "functions" are a quite universal type of data (reason will be made clear later)
- Specifically, for measuring the difference of two functions, we assume the domain to be the same, aka. we measure two functions of the following form:
 - $f: X \to \mathbb{R}$
 - $g: X \to \mathbb{R}$

• The idea of our distance d(f, g) for the two functions f, g is to measure the maximum of difference of the function values at each point in the domain X

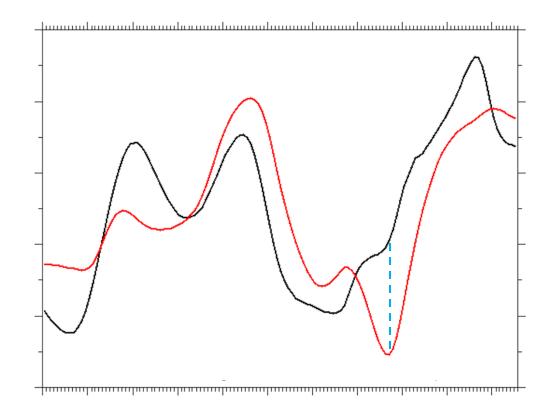
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- The distance is also denoted $\| f g \|_{\infty}$:

$$|| f - g ||_{\infty} = \max_{x \in X} \{|f(x) - g(x)|\}$$

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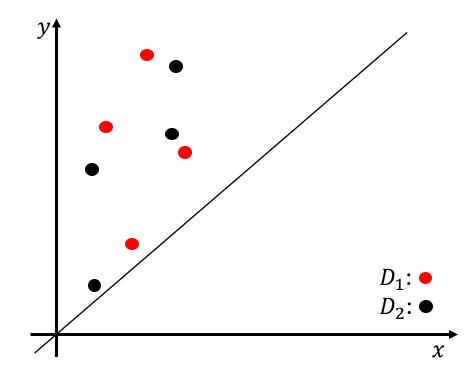


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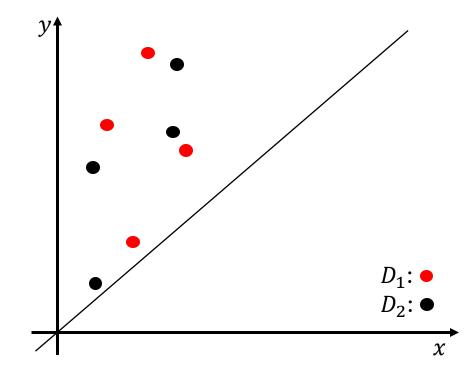


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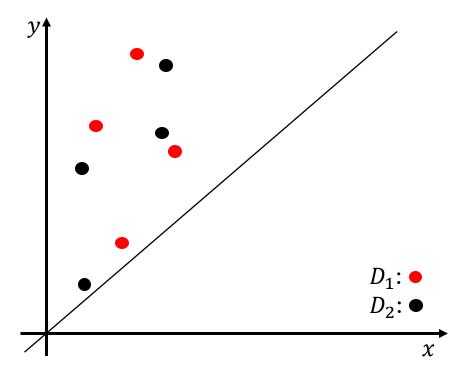
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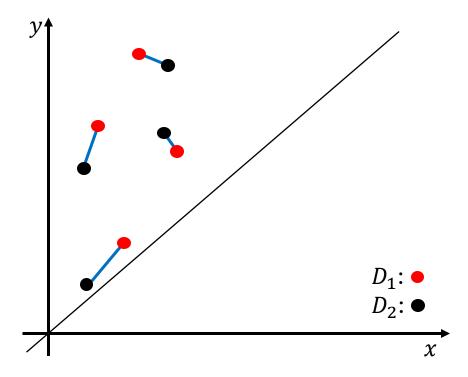
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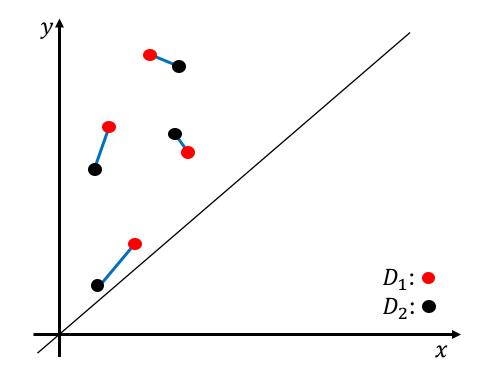
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- Furthermore, we want the matched points in the two PDs to be as close as possible, which indicates that overall the two PDs are "close" to each other



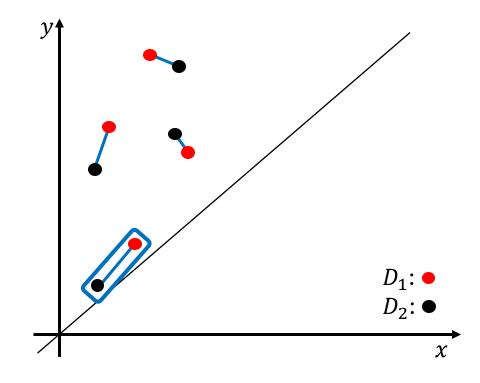
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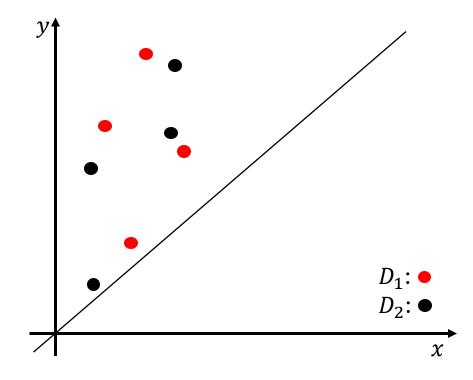
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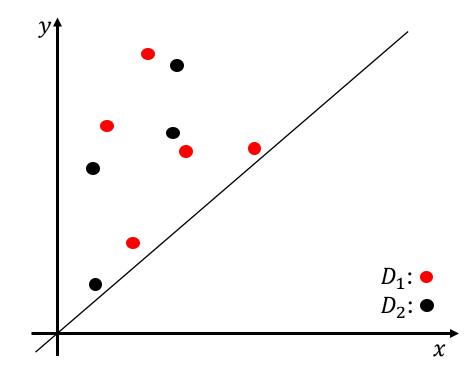
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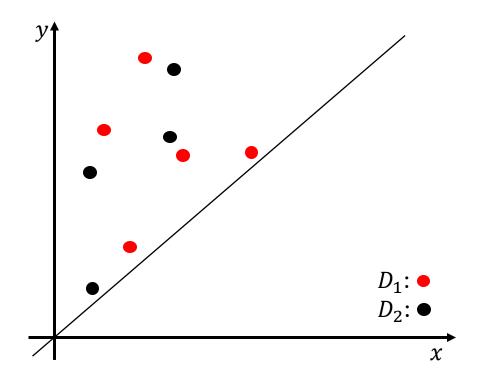
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 - However, we can only "ignore" then when they are not very "important", aka. being close to the diagonal

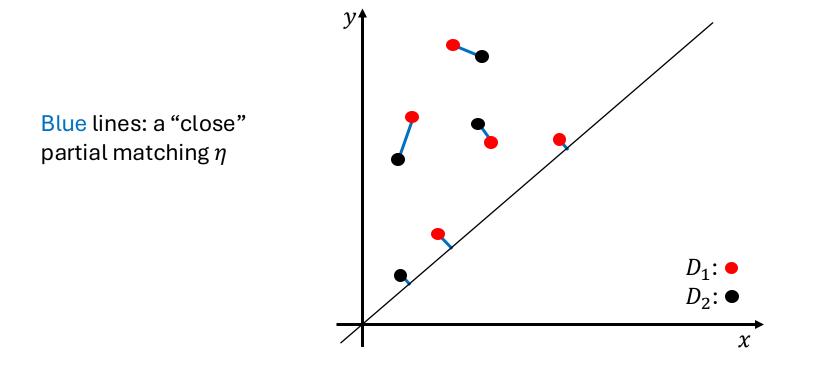
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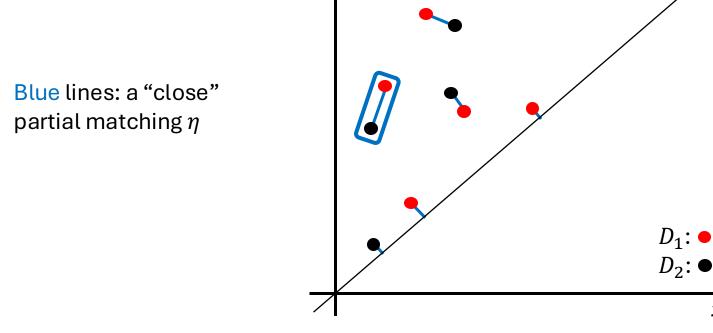


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The **bottleneck distance** is then defined as follows:

 $d_{\rm B}(D_1, D_2) = \min_{\eta \text{ over all partial matchings}} \{ cost(\eta) \}$

aka. the minimum cost of all partial matchings

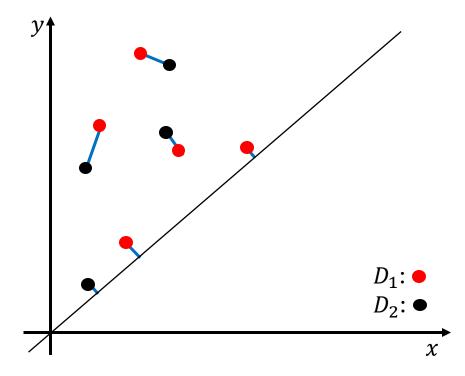
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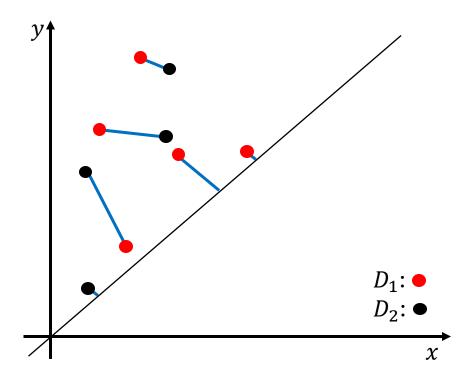
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 - Each unmatched point is "insignificant" (length is small) so we can "ignore" them
- So overall the two PDs are close

• The following is a "close" partial matching (with small cost) which achieves the bottleneck distance between D_1 , D_2 (aka. $d_B(D_1, D_2) = cost(\eta)$)



• The following is a partial matching where the max distance between matched points is high



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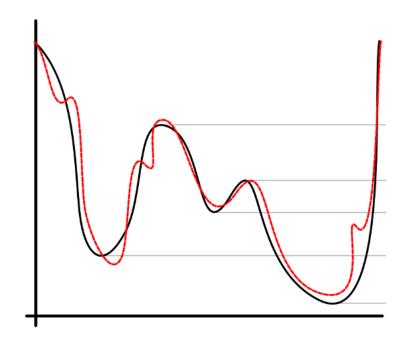
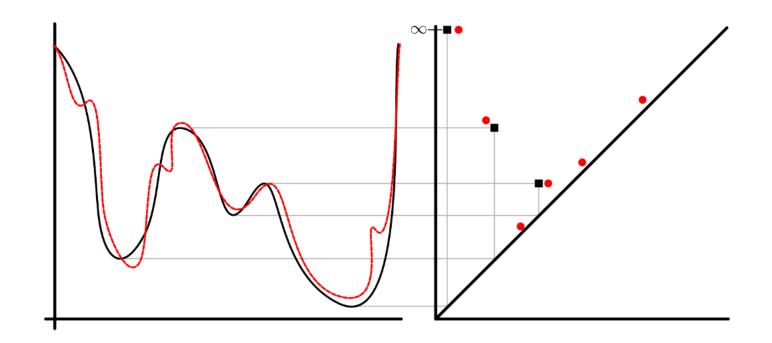


Figure from: Bendich et al. Topological and statistical behavior classifiers for tracking applications

• Stability Theorem: For any two functions $f, g: X \to \mathbb{R}$, one has

 $d_{\mathrm{B}}(PD(f), PD(g)) \leq \| f - g \|_{\infty}$

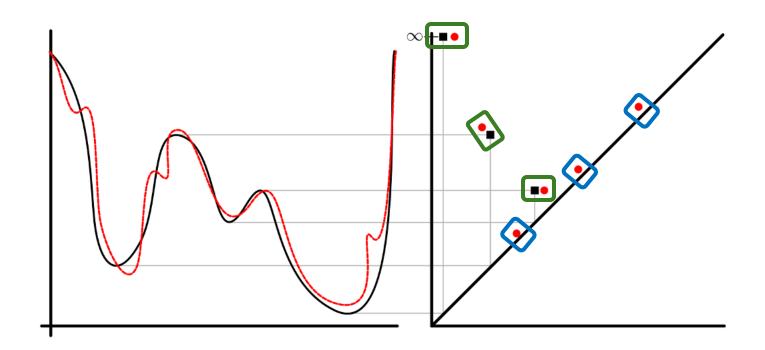
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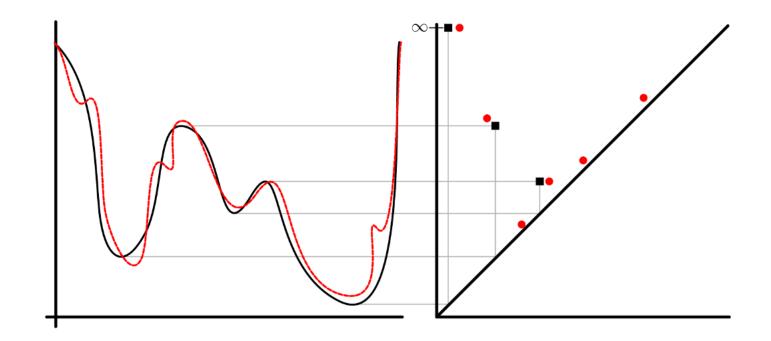
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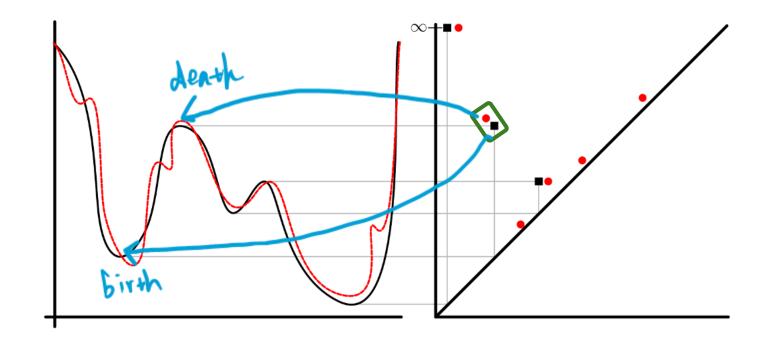
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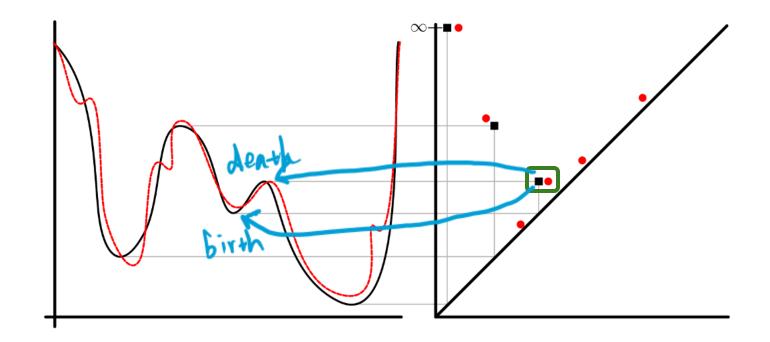
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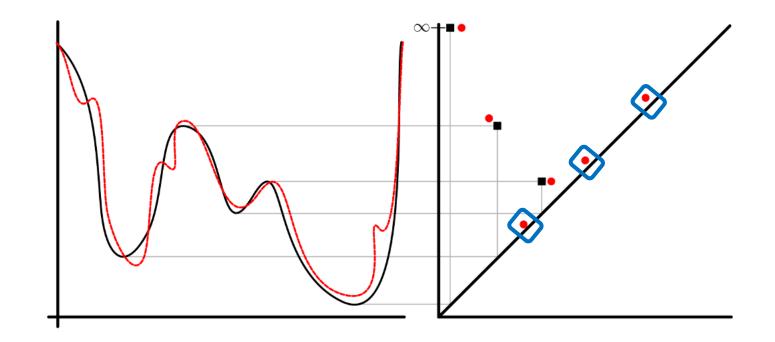
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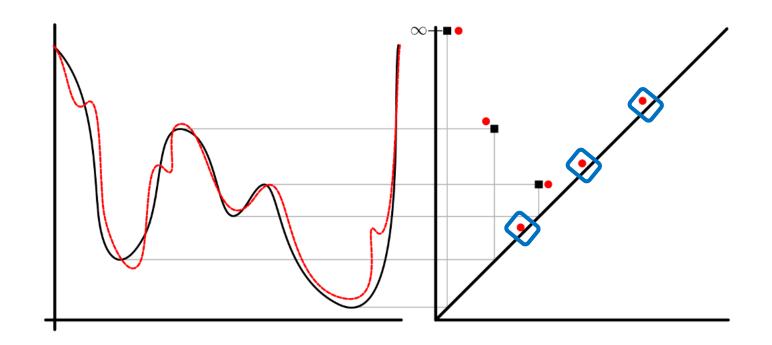
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- This also corresponds to the fact that the red curve is more noisy than the black one



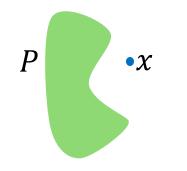
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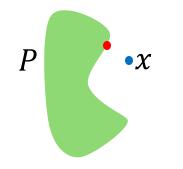
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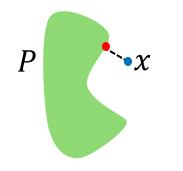
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- Corollary: The above observation also means that $PD(f_P)$ equals the PD of the Čech filtration of P

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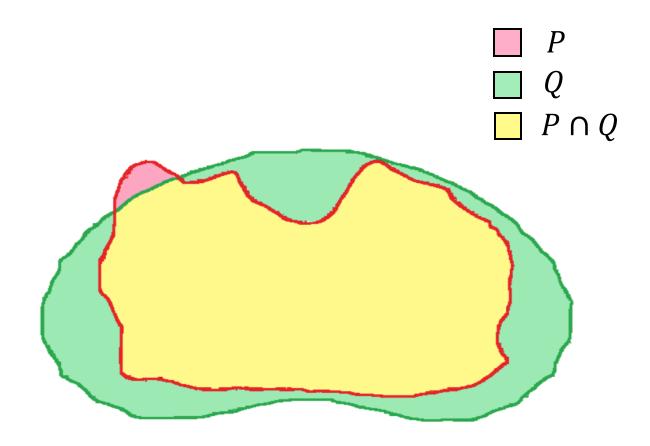
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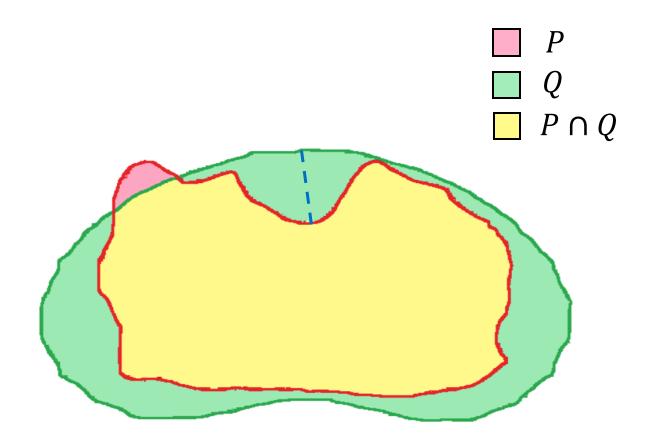
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• The **Hausdorff** distance is then the maximum of the two:

$$d_H(P,Q) = \max\{d_H^{unsym}(P,Q), d_H^{unsym}(Q,P)\}$$



Img source: Karimi & Salcudean. Reducing the Hausdorff Distance in Medical Image Segmentation With Convolutional Neural Networks



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- Let p be the point in P such that d(x, P) = d(x, p).
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- Based on the definition of Hausdorff distance, we have

$$d(p,q) = d(p,Q) \le d_H(P,Q)$$

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• We then have $d(x,Q) - d(x,P) \le d_H(P,Q)$

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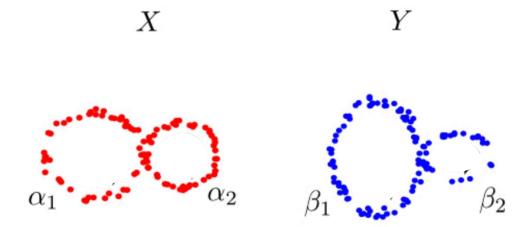
- Define $PD_{Rips}(P) \coloneqq PD(\mathcal{R}(P))$ for a point cloud P and recall that $\mathcal{R}(P)$ is the Rips filtration of P
- Stability Theorem for Point Clouds (Vietoris-Rips): One has

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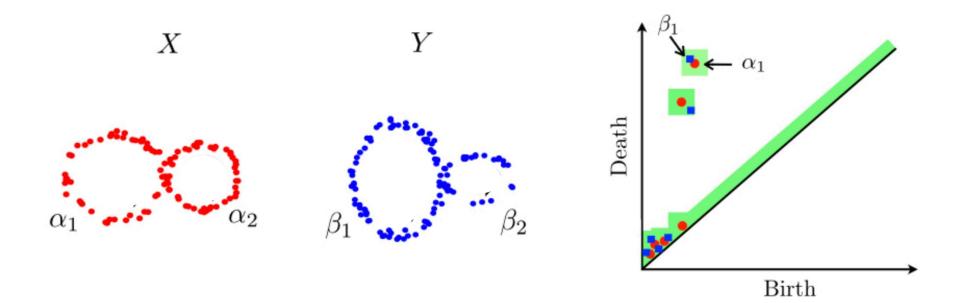
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 Proof of this needs the more advanced notion of "interleaving" stability and is beyond scope



Img source: Kusano, Fukumizu, Hiraoka. Kernel Method for Persistence Diagrams via Kernel Embedding and Weight Factor



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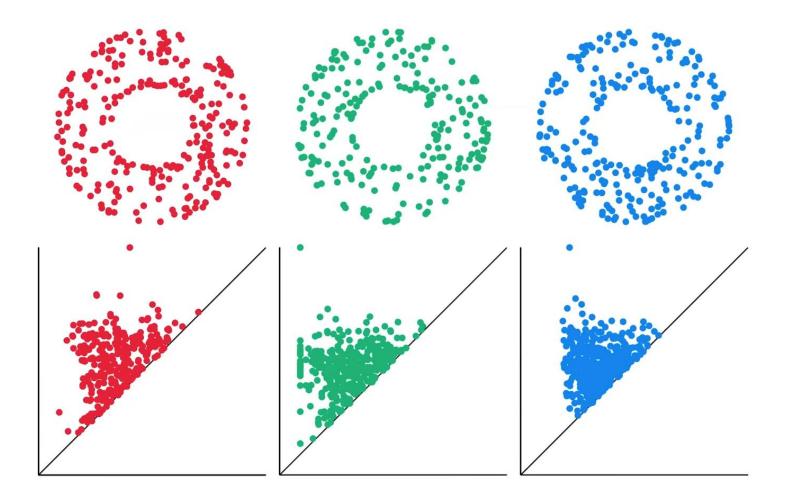


Figure from: Bastian Rieck: Topological Data Analysis for Machine Learning III: Topological Descriptors & How to Use Them (https://www.youtube.com/watch?app=desktop&v=7i1kabhl5IU)

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 The theorem follows from the previous "interleaving" between Čech and Rips complexes (but details omitted)

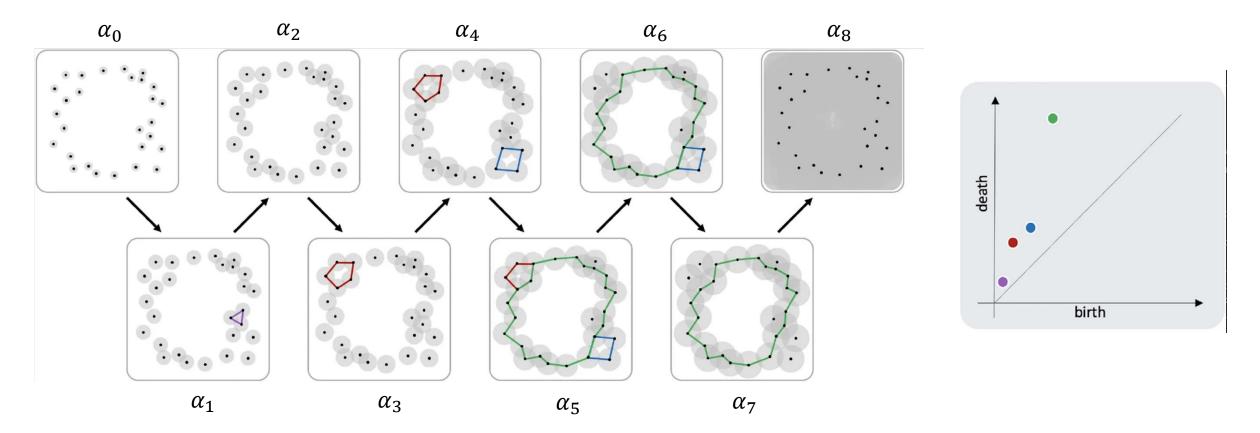
 $\mathbb{C}^{\alpha}(P) \subseteq \mathbb{V}\mathbb{R}^{\alpha}(P) \subseteq \mathbb{C}^{2\alpha}(P)$

Computing bottleneck distance

- <u>https://github.com/nihell/tutorialathon/blob/master/BottleneckTutorial.ipyn</u>
 <u>b</u>
- https://www.youtube.com/watch?v=4WswT9snTjc

• A practical use of drawing a PD as a barcode is that barcode provides a way to "visualize" the Betti number across the different range (value α)

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- For the previous filtration on a point cloud and its 1d PD



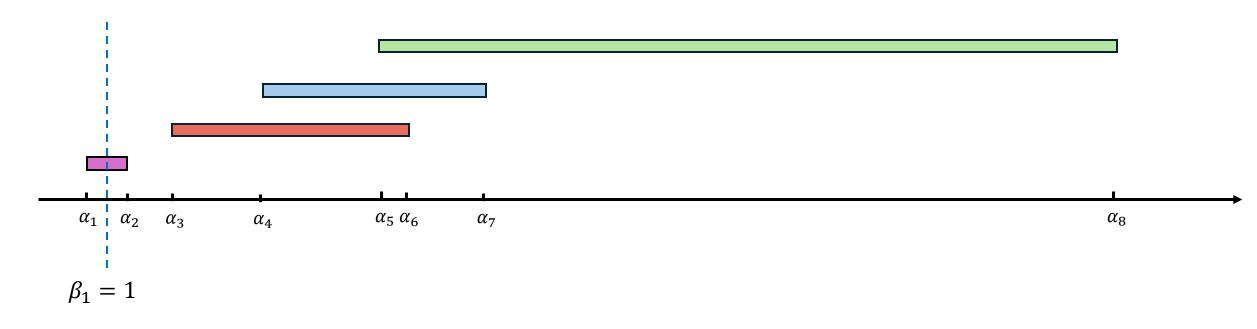
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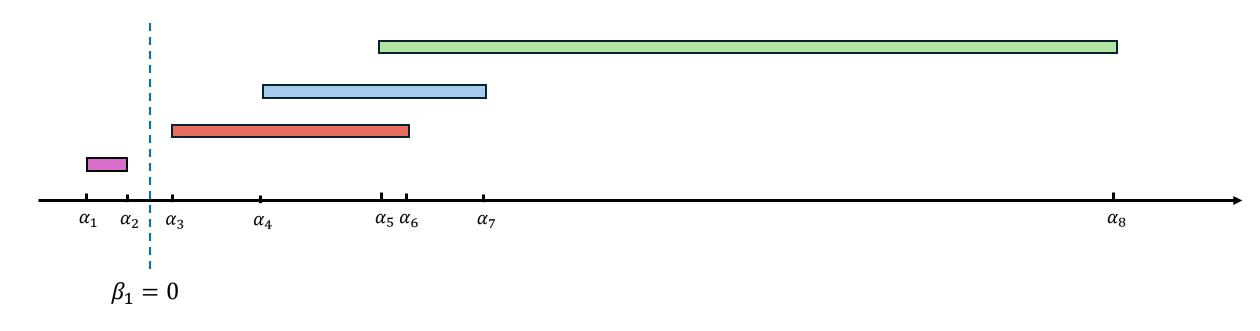
- The following is its 1d barcode
- Observe, if you count the number of intervals (bars) containing a certain value, then it gives you the 1st Betti for the complex corresponding to the range (value α) in the filtration



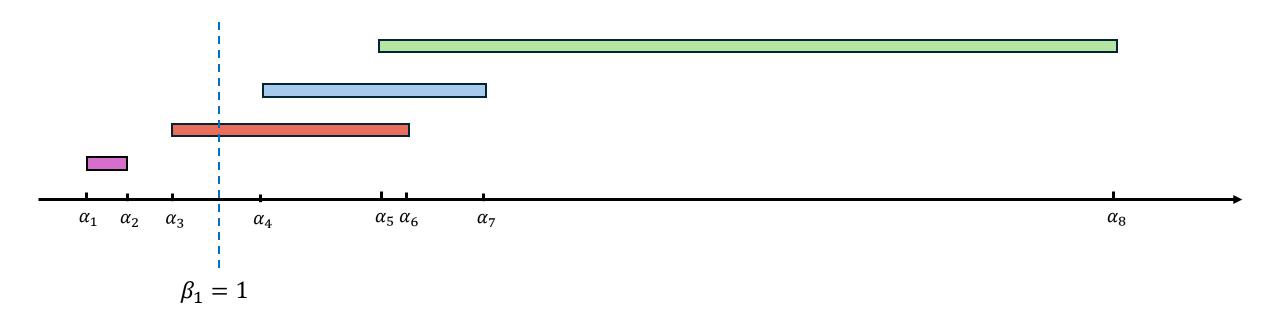
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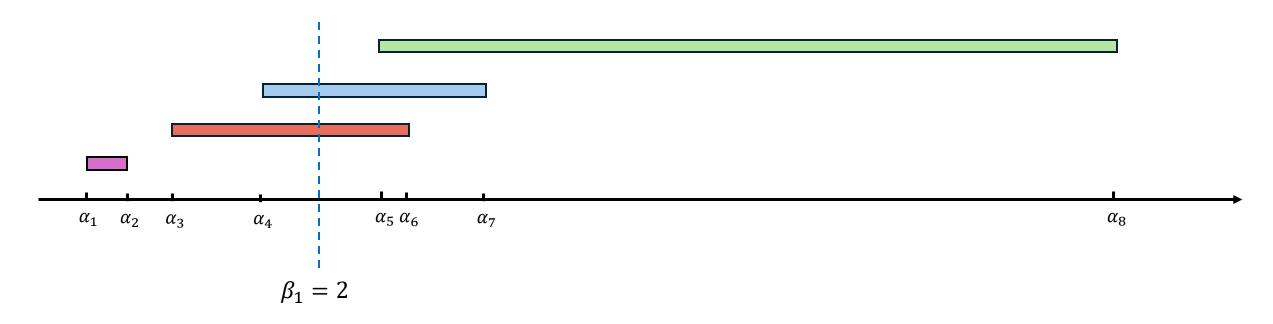
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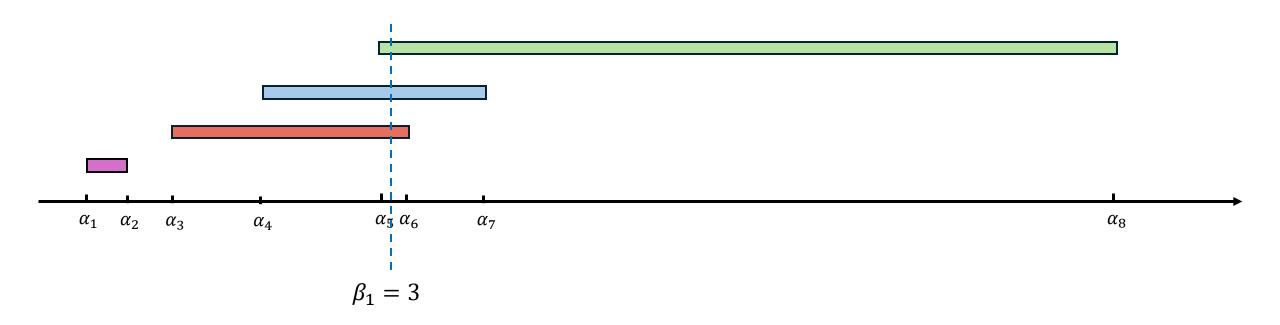
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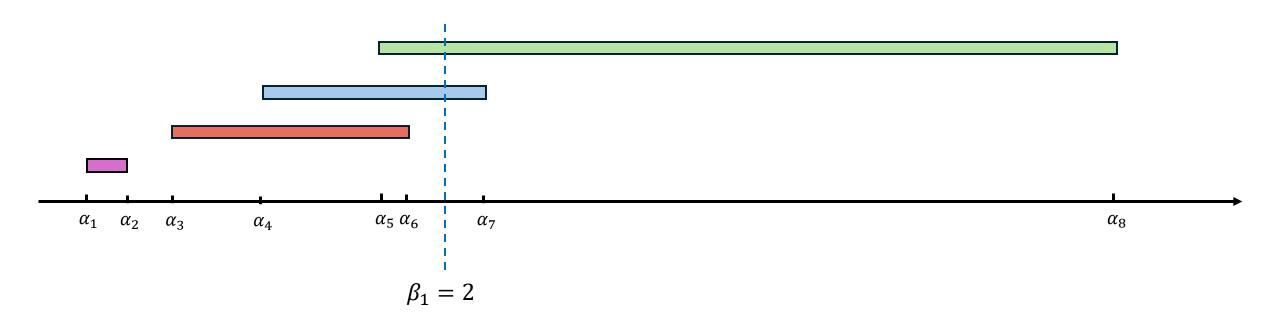
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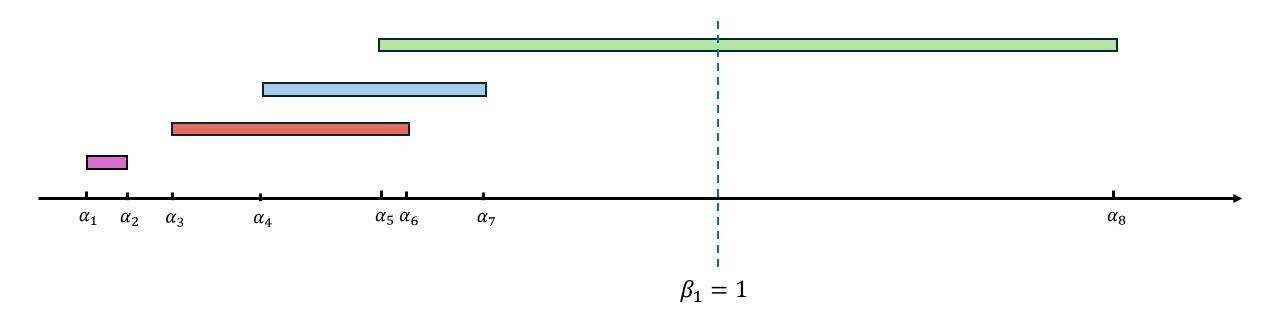
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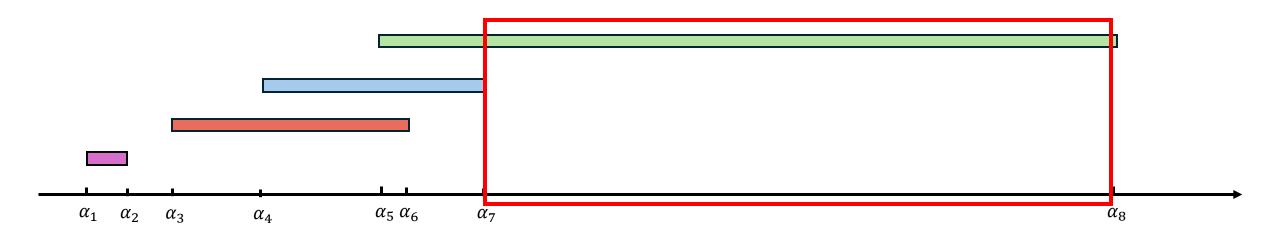
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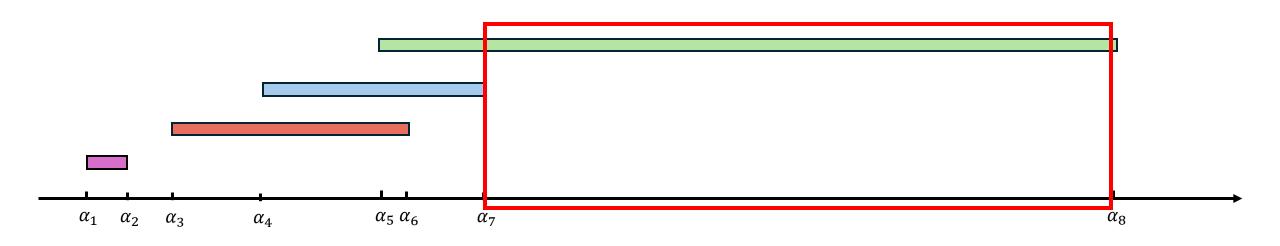
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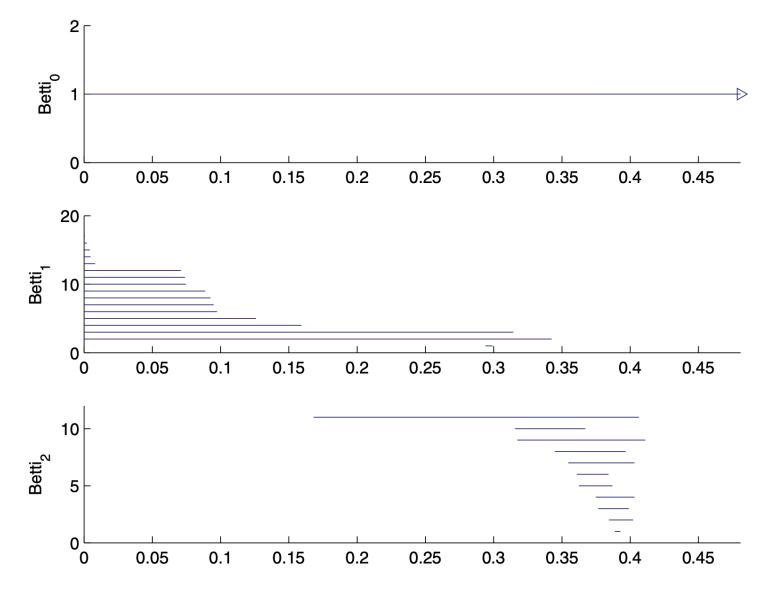
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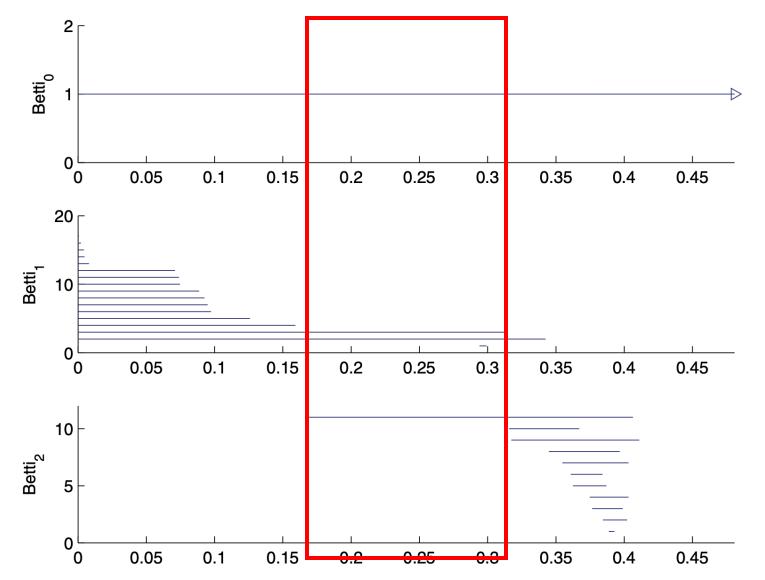
- More powerfully, if you look at the range of the values where the Betti number stays the same, and take the longest range, that would give you the most probably inference of the Betti number
- So most probably, for the previous point cloud, $\beta_1=1$ (complying out intuitions)



Another example



Another example

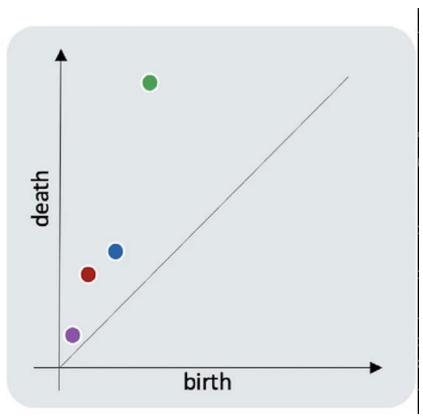


Counting Betti number using PD

• Now we know that taking the intersection of a vertical line with the intervals in the barcode gives you the Betti number at the value of the vertical line, what about PD? Can we do similar things in PD?

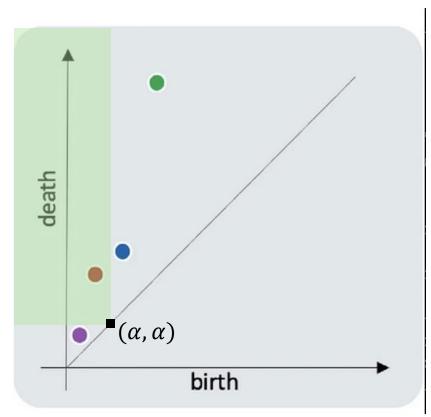
Counting Betti number using PD

• It turns out that, the number of intervals intersecting a value α as manifested on the PD is the number of pointing in the upper-left quadrant of the point (α, α) on the diagonal

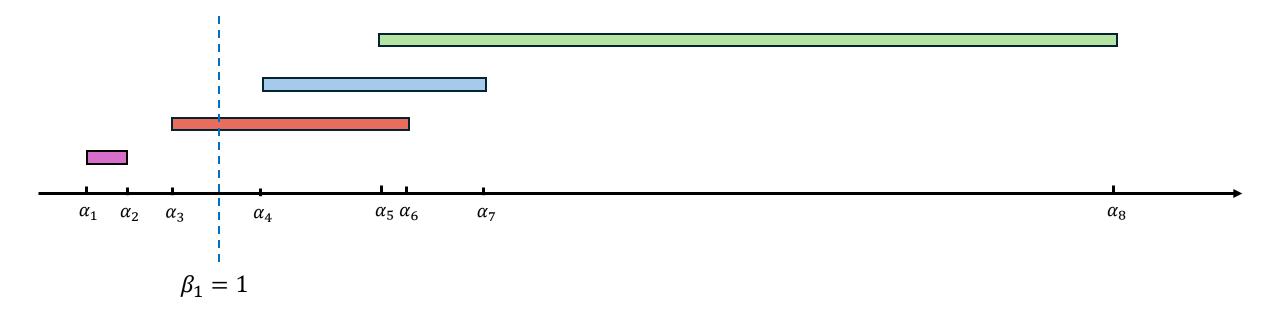


Counting Betti number using PD

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On barcode



On PD

• $\alpha_3 < \alpha < \alpha_4$

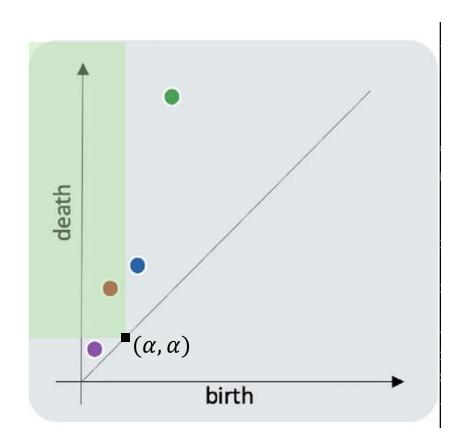
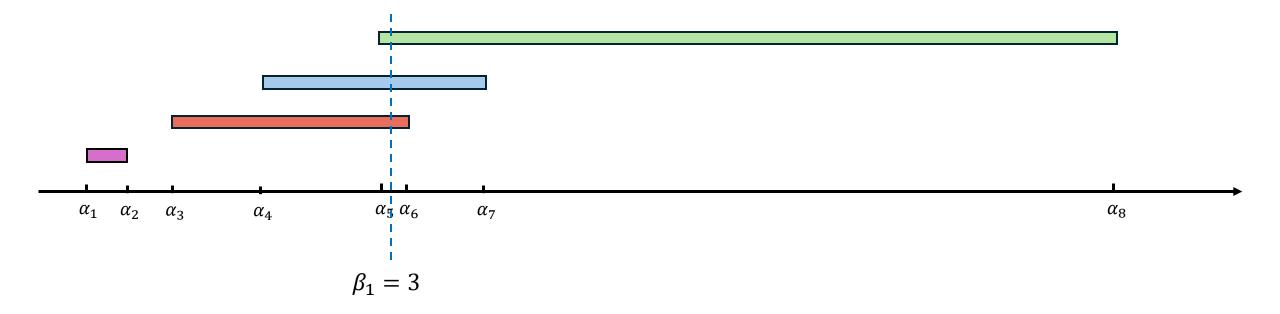


Image source: Bobrowski O, Skraba P. A universal null-distribution for topological data analysis.

On Barcode



On PD

• $\alpha_5 < \alpha < \alpha_6$

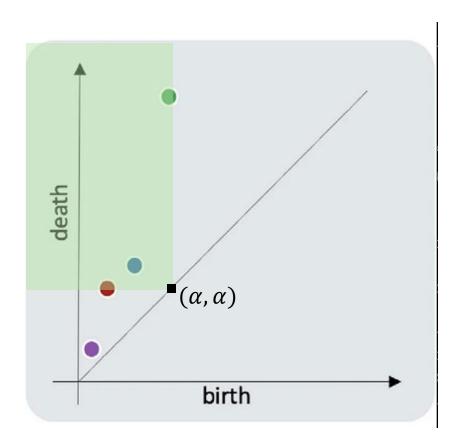
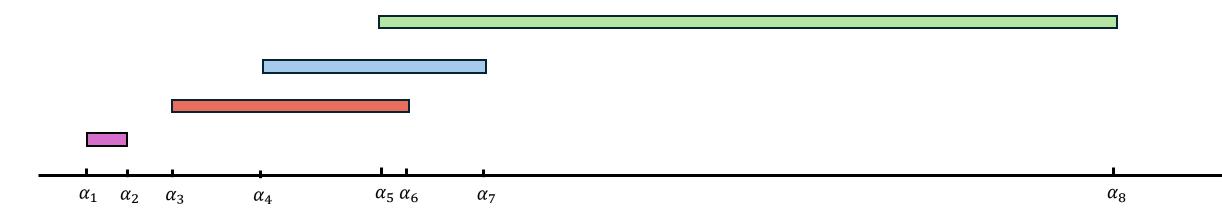


Image source: Bobrowski O, Skraba P. A universal null-distribution for topological data analysis.

- A typically belief in TDA is that people often think of features (intervals) with long lifespans as robust, important features, whereas a short lifespan may be an indication that the feature is less essential and may in fact be due to noise in data
- A commonly adopted approach in using PD/barcode for inference is to find a "cut-off" for the short intervals (noise), and focus on the long intervals (actual features) only

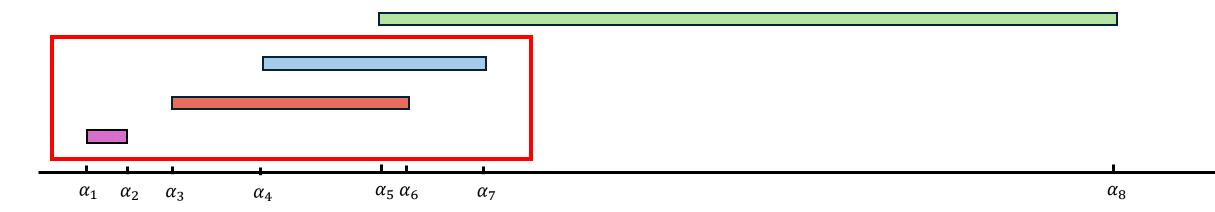
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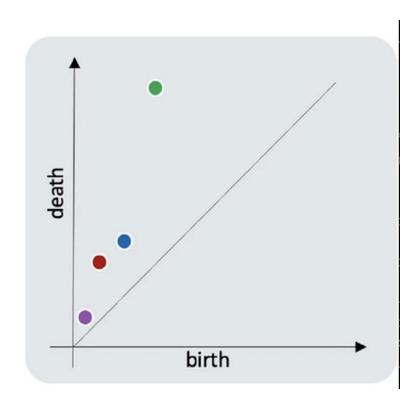
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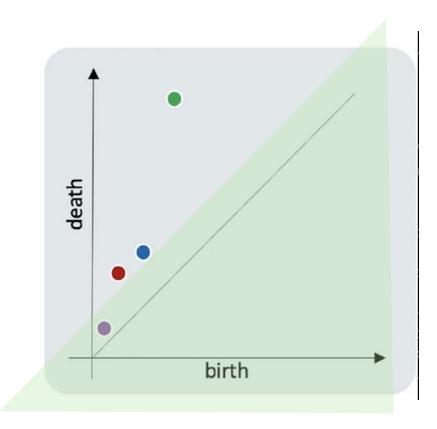
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