Persistent Homology: Filtration building techniques

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Outline for studying persistent homology

- 1. Intro to persistent homology
 - Build intuitions of persistent homology: what it does, what it produces
- 2. Formalizing persistent homology
 - Introduce its input (filtration) and study an algorithm for computation
- 3. Different ways for building filtrations
 - Vietoris-Rips filtration, sub-levelset filtration
 - Cubical complexes (for images)
- 4. Interpretation and stability of persistence diagram

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- So far, we have formally defined the PD for a (discrete) filtration, which is a finite sequence of simplicial complexes that are also nested
- But we haven't formally defined PD for a continuous filtration, where we have a space varying over $\alpha \in [0, \infty)$ (technically, there're infinitely many of them)



"Continuous" filtration

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- So to harness the power of persistence, you have to do this
- So we shall not only formally define PD on input data (which are typically continuous at least theoretically), but also learn ways to preprocess the data into filtrations to feed into the persistent homology pipeline



https://pixabay.com/photos/new-year-background-tree-sunset-736885/; Adler et al. Persistent homology for random fields and complexes.; https://builtin.com/artificial-intelligence/transformer-neural-network

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- We shall eventually show that continuous filtration is in some sense "equivalent" to the discrete one



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Image source: https://www.datacamp.com/tutorial/k-means-clustering-python

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 - Even for supervised learning (another type of more popular? machine learning), if you ignore the "labels" for the data, then the data become point clouds
 - After all, each element in your data is in some sense a "point"

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- Trying to infer the shape of point cloud is indeed a major motivation for topological data analysis





- Recall that each space in the "growing of balls" filtration is to take a ball of the same radius α centering in each point, and then take the union of the α -radius balls of all points
- To get the (continuous) filtration, we then let the radius α increase from 0 to ∞, and let the union of balls grow with it (see: https://gjkoplik.github.io/pers-hom-examples/0d_pers_2d_data_widget.html)
- Also recall our goal is to use a discrete filtration to "emulate" this continuous filtration
- Since a discrete filtration consists of simplicial complexes, what we really need to do is to find a way to use a simplicial complex to "emulate" the union of balls
- To construct a simplicial complex for the union of balls, we first let the point in the point cloud be all the vertices in the simplicial complex
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- a
- b
- C
- d



- a
- b
- C
- d



- a
- b
- C
- d



- a
- b
- C
- d



- a ab
- b
- C
- d



Simplices:

- a ab
- b
- C
- d

a

• C

d ullet







Simplices:

- a ab
- b
- C
- d

a

• C

d ullet







Simplices:

- a ab
- b cd
- C
- d

●b

a

• C

d ullet







- a ab
- b cd
- C
- d



- a ab
- b cd
- C
- d



- a ab
- b cd
- c ad
- d



- a ab
- *b cd*
- c ad
- d



- a ab
- *b cd*
- c ad
- d


Build discrete filtration for "growing balls"

- a ab
- b cd
- c ad
- *d bc*



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• To understand this definition, a brute force (inefficient) algorithm for computing $\mathbb{C}^{\alpha}(P)$ out of a point cloud P: enumerate each subset p_0, p_1, \dots, p_d of P, and check whether intersection of balls of the points is non-empty

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- Of course, in practice, the algorithms that people use to compute Čech Complexes are much more efficient ones

- a
- b
- C
- d



- a
- *b*
- C
- d



- a ab
- b
- C
- d



Simplices:

- a ab
- *b cd*
- C
- d

Ъ

a

• C

d∙







- a ab
- *b cd*
- c ad
- d



- a ab
- b cd
- c ad
- *d bc*



- a ab
- b cd
- c ad
- *d bc*



- *a ab abcd* with all its faces
- b cd (abc, abd, acd, and
- c ad bcd).
- *d bc*



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- E.g., if three balls intersect, then any two balls also intersect. So the edges of the corresponding triangle are also in the Cech complex.

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- The above "equivalence" is called the "homotopy equivalence" in algebraic topology, whose definition is beyond the scope



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 - Notice that an interval of $PD(\mathcal{F}^c)$ is actually a continuous interval over the real line \mathbb{R} .


























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- After all, trade-offs were made everywhere in computer science between efficiency and quality

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- There are again finitely many α where the Rips complexes change, so the Rips filtration is a discrete finite filtration













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Image source: Reani, Bobrowski. Cycle Registration in Persistent Homology with Applications in Topological Bootstrap

Čech

Rips

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- But notice that other than the two triangles the remaining simplices in the two complexes are the same



Čech

Rips

- Following is an example where the Čech and Rips complexes are different
- But notice that other than the two triangles the remaining simplices in the two complexes are the same
- Furthermore, if we increase the radius for Čech complex, the two missing two triangles will come into picture ---- in some sense, the sequences of Čech and Rips complexes are "interleaved with each other"



Čech

Rips

• As for $\mathbb{C}^{\alpha}(P)$, we also need to show that $\mathbb{VR}^{\alpha}(P)$ is not only a set of simplices but also a simplicial complex, which is checking that the faces of each simplex in the set $\mathbb{VR}^{\alpha}(P)$ are also in $\mathbb{VR}^{\alpha}(P)$.

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- Verification: Take any $\sigma \in \mathbb{VR}^{\alpha}(P)$ and a face τ of σ (a subset of σ , i.e., $\tau \subseteq \sigma$).
- We have any pairs of the α -balls for the points in σ intersect. So any pairs of the α -balls for the points in τ intersect (because τ is a subset). So $\tau \in \mathbb{C}^{\alpha}(P)$

PD for Vietoris-Rips Filtration

- We now also define PD for a continuous filtration using the discrete Rips filtration
- Again, define the PD for the continuous filtration \mathcal{F}^c by "inducing" from the PD of the discrete Rips filtration $\mathcal{VR}(P)$:
 - Each complex in the discrete $\mathcal{VR}(P)$ corresponds to a range of α -values (and a range of spaces) in the continuous filtration
 - For each interval $[b, d) \in PD(\mathcal{VR}(P))$, consider the corresponding complexes in $\mathcal{VR}(P)$, which are $K_b, K_{b+1}, \dots, K_{d-1}$.
 - Then, the corresponding interval of $PD(\mathcal{F}^c)$ is the union of all the α -values corresponding to $K_b, K_{b+1}, \dots, K_{d-1}$.

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 - Then, the corresponding interval of $PD(\mathcal{F}^c)$ is the union of all the α -values corresponding to $K_b, K_{b+1}, \dots, K_{d-1}$.
- Notice now there is data loss introduced because $\mathcal{VR}(P)$ is not exactly the same as $\mathcal{C}(P)$

"Similarity" of Čech and Vietoris-Rips

 Interleaving Theorem of Čech and Rips Filtration : For any point set P and any radius α,

 $\mathbb{C}^{\alpha}(P) \subseteq \mathbb{V}\mathbb{R}^{\alpha}(P) \subseteq \mathbb{C}^{2\alpha}(P)$
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- **Claim**: Due to the previous interleaving of the two sequences, we have that $PD(\mathcal{VR}(P))$ "approximates" $PD(\mathcal{C}(P))$ well.
- Remark: This "approximation well" thing will be made more formal later.

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- Remark: This "approximation well" thing will be made more formal later.
- Another observation: The edges of the two Čech and Vietoris-Rips complexes for the same point set P over the same radius α are the same.
- Reason: Edges are formed by two points. If you check the definition of Čech and Vietoris-Rips ("all balls for a set of points intersect" and "each pair of balls for a set of points intersect"), when we only have two point, the two criteria become the same.

- **Definition**: Given a point set P and a distance r, the Vietoris-Rips complex of P corresponding to the distance r, denoted $\mathbb{VR}^r(P)$, is a simplicial complex whose vertices are points in P such that
 - A subset of points p_0, p_1, \dots, p_d of P form a d-simplex if and only if for each pair of points p_i, p_j in the subset, their distance is no more than r, i.e.,

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- The Rips complex in the above definition is the same as the previous Rips complex by taking r/2 as radius (if two points p_i , p_j have distance no more than r, then their r/2-balls intersect)
- The benefit of the above alternative definition is that we can completely eliminate balls and define Rips complexes / filtration by only considering the pair-wise distances between points

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- They are abstract objects but we have some pair-wise "distances" between these objects.

• In this example, the data points are "regions of the brain", and we can calculate their "similarity" by measuring the correlation of their blood oxygen fluctuation over time (a time series data).



Figure courtesy of: Duy Duong-Tran

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• These regions are not really technically having a position (each region is represented by a blood oxygen level function over time), but we have distances between the regions



• For this data, we still can build Rips filtration on these brain regions



Figure courtesy of: Duy Duong-Tran

- We shall briefly look at some facts concerning computing Rips Filtration.
- Recall:
- **Definition**: Given a point set P and a distance r, the **Vietoris-Rips** complex of P corresponding to the distance r, denoted $\mathbb{VR}^r(P)$, is a simplicial complex whose vertices are points in P such that
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- This means that a Rips complex over a certain distance (radius) *r* is completely determined by the distances of each pair of points in *P*
- Since a pair of points form an edge, this also means that a VR^r(P) can be completely determined once we have figured out the edges (1-simplices) for VR^r(P)

- So, for a certain r, to compute $\mathbb{VR}^r(P)$, our first thing to do: Enumerate each pair of points in P and check whether their distance is no more than r.
 - If this is true, we let the pair form an edge in $\mathbb{VR}^r(P)$
- By doing this, we have all the edges in $\mathbb{VR}^r(P)$.
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- $\mathbb{VR}^{r}(P)$ is then the **Clique complex** of the graph $\mathbb{G}^{r}(P)$.

Clique

- **Definition**: A clique of a graph G = (V(G), E(G)) is a subset S of V(G) such that each pair of vertices of S form an edge in G.
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Image source: Santamaría, Therón. Overlapping Clustered Graphs: Co-authorship Networks Visualization

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- E.g, the following graph has three maximal cliques (a clique not contained in another clique)



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- E.g, the clique complex of below graph has three maximal simplices G_1 , G_2 , and G_3 (a simplex not being a face of another simplex) and all their faces





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- This means that only crossing the values $d(e_1) < d(e_2) < \cdots < d(e_m)$, $\mathbb{G}^r(P)$ changes
- Finding these values takes $O(n^2 \log n)$ time dominated by the sorting, where n is the number of points in P

Another type of filtration

- **Delauney** complexes / filtrations: growing the balls around points, construct a simplex whenever their set of balls intersect (the same as Cech complexes)
- Difference: the part of a ball stops growing when touching another ball.



Image source: Li & Liang. Knowledge-Based Energy Functions for Computational Studies of Proteins



Image source: Mishra, Motta. Stability and machine learning applications of persistent homology using the Delaunay-Rips complex

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- Disadvantage of Delauney complexes: costly to compute, especially when the dimension of the points in the point cloud is high
- The go-to filtration for point cloud is **Rips filtration** because of (1) its computational efficiency and (2) the fact that it still faithfully recover the shape of the data (despite data loss)

Other types of complexes

- There are other types of complexes:
 - Witness complex
 - Graph-induced complex
 - Tangential complex
 - ...
- Will not cover them at least for the time being

Data as a function

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• E.g., all pixels in an image form the domain of a function and the color value on each pixel is basically the function value on a point of the domain

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 - Even if the range of the function is more than a single real value, say again, a colored image, we can take each channel (RGB), this will give you three individual real-valued functions. We can analyze each individually using persistence

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- Doing this for 3d images or arbitrary simplicial complexes can be generalized

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 $\implies f: X \to \mathbb{R} \implies \text{Filtration} \implies \text{PD}$

Image

• We visualize the domain X of a 2d image as a regular grid, where pixels are grid points (below an example of 4x4 image)

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- So we subdivide the grid to be consisting of triangles, so X becomes a simplicial complex



- Another problem: the function values are only given on the vertices (which are gray-scale values on the pixels from the given image)
- We need function values on the edges and triangles: for this we take the "maximum" value of the vertices that an edge or triangle contains



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Image function

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- We then take all possible functions values (there are finitely many of them) and sort them (i.e, start with the lowest value):

 $\alpha_0 < \alpha_1 < \dots < \alpha_m$

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• It should be esay to verify that $f^{-1}(-\infty, \alpha_i] \subseteq f^{-1}(-\infty, \alpha_{i+1}]$ for any i

• $f^{-1}(9] \subseteq f^{-1}(10] \subseteq f^{-1}(11] \subseteq f^{-1}(12] \subseteq f^{-1}(13] \subseteq f^{-1}(14] \subseteq f^{-1}(15]$







• $f^{-1}(9] \subseteq f^{-1}(10]$



• $f^{-1}(9] \subseteq f^{-1}(10] \subseteq f^{-1}(11]$



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PD for the sublevelset filtration

PD for the sublevelset filtration

• There is a 1-dimensional bar [14,15) in the PD



PD for the sublevelset filtration

• There is a 0-dimensional bar [10,12) in the PD



3D image

• We view the domain X for a 3D image as a 3D grid, and we have a function value on each grid point



3D image

• We also need to subdivide the cube into (six) tetrahedra to make the domain a simplicial complex



3D image

- And then we only need to assign value to each edge, triangle, tetrahedron based on the maximum values of their vertices
- The sublevelset filtration can then be defined similarly



More about 3D images

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- 3D images can be considered as a stacking of several 2D images, and are commonly used in medical imaging (e.g., CT-scans, MRI)
- Analyzing medical images is a hot and important in image processing



Img source: Jackowski, Papademetris, Dobrucki, Staib. Characterizing Vascular Connectivity from microCT Images

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- Naturally, we could also define sublevelset filtrations on triangular meshes by assigning function values to the vertices (edges / triangles are then induced)
- There is a natural way to assign values to the vertices which is to use the "height function"

















More sublevelset filtrations

- Indeed we have also seen sublevelset filtrations in previous slides
- An interactive example: <u>https://iuricichf.github.io/ICT/filtration.html</u>

- There is a counterpart of sublevelset filtration called **super**levelset filtration
- A superlevelset of is the subset of X whose function values are greater than or equal to a value α , and we denote it as $f^{-1}[\alpha, \infty)$
- We then take all possible functions values and **descreasingly** sort them (i.e, start with the height value):

$$\alpha_0 > \alpha_1 > \dots > \alpha_m$$

• The superlevelset filtration is then the superlevelsets over the above values:

$$f^{-1}[\alpha_0,\infty) \subseteq f^{-1}[\alpha_1,\infty) \subseteq \cdots \subseteq f^{-1}[\alpha_m,\infty)$$

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$$\mathcal{F}\colon K_0\subseteq K_1\subseteq\cdots\subseteq K_m=K$$

intervals in the PD are "integer intervals" (e.g., $[3, 6) = \{3, 4, 5\}$).

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- This applies when it is not clear where the discrete filtration is built from
- In practice, filtrations are built from different types of data. Each complex in the discrete filtration is associate with a real value (or a bunch of them)
- Intervals in the PD for such a filtration (when we know source data) is then continuous intervals of real values (e.g., [3.52, 6.37))

• For the previous sublevelset filtration for image, we can also number each complex in the filtration from 0 to 6



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- For the previous sublevelset filtration for image, we can also number each complex in the filtration from 0 to 6
- But we use the pixel values (e.g., 9, 10, ...) instead of the integer indices (e.g., 0, 1, ...) for the PD



• E.g., the below 1d interval is [14,15) rather than [5,6)



• The below 0d interval is [10,12) rather than [1,3)

