Case study: Image segmentation for accurate topology inference

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Image segmentation for correct topology inference

This application is documented in the following paper

 Chung, Yu-Min, and Sarah Day. "Topological fidelity and image thresholding: A persistent homology approach." Journal of Mathematical Imaging and Vision 60 (2018): 1167-1179.

Persistent homology pipeline



homology inference

Some img from: AATRN; https://quantdare.com/understanding-the-shape-of-data-ii/

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 - β_1 , # of 1-dimensional holes: "tunnels"
 - β_2 , # of 2-dimensional holes: "cavities"



Image source: Eckert et al. The Topology of Pediatric Structural Asymmetries in Language-Related Cortex

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- After the binarization, we can then study the property of each region





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 - Tunnel can pinch off into cavities under the pressure of accumulated snow and thus the gas may get trapped
 - If tunnels are not correctly detected due to topological inaccuracies, then gas transfer simulations will underestimate the propagation of gases through the firn
 - Accurate estimates of gas transfer are of interest to climate scientists seeking to understand the age of atmospheric gases that have been trapped in air pockets within firn core samples

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 - This is also the method of comparison in the work

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Heidari et al. A new general model for quantum image histogram (QIH)

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- We will later see examples where the topology inferences based on Otsu's is far off

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- Given a digital image of two or more important subdomains, proposed an automated thresholding approach based on the PD
- It utilizes an optimization scheme so that important, robust topological features are present, but the number of features (holes) that are (most likely) due to noise is minimized
- As shown in the paper, the proposed method often leads to improved topological accuracy in the segmented image and in the resulting Betti numbers
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• For a threshold *t*, the work focuses on the black/dark region corresponding to this threshold, which is indeed the sublevel set of *t*:

$$f_t^- := \{ x \in P \mid f(x) \le t \}$$

• We then build the sublevelset filtration of *f*:

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- Recall: computing the number of intervals in the barcode \mathscr{P} intersecting a threshold *t* gives the Betti number
- which in turn is equivalent to counting the number of number of points in the upper-left quadrant of the point (t,t) in the PD \mathscr{P}





• So, as we move the lower right corner of the quadrant along the diagonal and count the number of points falling within it, we track the variation of the Betti number $\beta(\hat{f}_t^-)$ of the sublevel sets in the filtration

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

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• Now the problem boils down to finding a t such that $\beta(\hat{f}_t^-)$ is the "ground truth"



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- We take \hat{f} to be the "ground truth"



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- In either case, a threshold value between 50 (min value) and 200 (max value) would give you a correct binarization (in terms of Betti number)



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 - (Notice here since the domain of the function is only 1-dimensional, we could only count β_0 and all other Betti numbers are 0)
- So for this problem, the homology inference is a done deal once you have the barcode

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- Data often comes with noise
- Now, to make things closer to reality, we add some noise to the function \hat{f} , and get a more realistic function f that one could encounter





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- By looking at the intersection of the quadrant as we move the lower-right corner along the diagonal, we observe:
 - The range of values where the Betti number equals 2 is greatly reduced for the left function, now become [105, 140)
 - What's more, for the right function, there is now not a single value whose Betti number evaluates to 2 (we can never find the ground truth by simply using a threshold)
- BTW, to solve the problem of the right function, one could use a cut-off for the length of the intervals (distance to the diagonal) and keep only those points in PD above a 45° degree line (we could then count to 2 with remaining points)

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 - 1. In the middle of long features: This threshold tends to be in the middle of the life of long lifespan features so that these features tend not to have just appeared (been born) or be on the verge of disappearing (dying)
 - 2. Far from short features: It is also far enough from the short-lived features (most probably noise) from the both sides so that small perturbations of the threshold do not produce large changes in the noisy features included

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• To achieve the first goal (threshold sitting in middle of long features), they try to find a threshold *t* maximizing the following objective function:

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- We divide the whole function by $\# \mathscr{P}(t) + 1$ because we want to measure the average contribution of the term for the intervals (so for we do not penalize a t intersecting less long intervals)

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- The above objective function penalizes a *t* intersecting a lot of short intervals, and favors a *t* intersecting long intervals in the middle

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• $\mathscr{P}^{-}(t) = \{(b, d) \in \mathscr{P} \mid d \leq t\}$: set of intervals to left of t

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- For each term in $\Psi^{-}(t)$, if [b, d) is long, then the denominator is large making the whole term small; also, if t is far from [b, d), t d (distance of t to the interval) is large
- This means that it would penalize the situation where *t* is very close to some long intervals

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• The threshold T they look for is then the following:

$$T = \underset{t \in [0,I]}{\operatorname{arg\,max}} \Phi(t).$$

A side note

- One interesting thing to notice: for a threshold *t*, the quadrant with corner on (t,t) separates a PD into the three parts we were considering previously:
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- They performed 1000 trials applying thresholding to previous 1D function each time with an additive noise generated from a Gaussian distribution with mean 0 and standard deviation 2
- They then use Otsu's method and their method to generate threshold and compute the Betti number





• Their proposed method constantly outperforms Otsu's in terms of the inference of Betti numbers







 $\hat{\beta} = (\beta_0, \beta_1)$









 Although lifespan cutoff gives accurate Betti number, binary image in (c) produced by their method does not represent the original image well, in particular, in the lower region. This appears to be due to the inconsistent illumination in this image (difference in brightness)



 By manually partitioning the image into a 4×4 grid of subimages and segmenting each separately, both results are good. But Otsu's still produces a topological error (red)

Another synthetic dataset



- 1. Similarly to the 1D function, they start with a black / white image with a white square on the upper-left corner.
- 2. They then add some noise within a (slightly larger) square around the white square
- 3. Finally, the add noise on the whole image

Another synthetic dataset



- Histogram-base segmentation methods (e.g., Otsu) are strongly influenced by the large dark region
- This may result in losing the essential topological features of one connected component and a single hole ($\hat{\beta} = (1,1)$)

Another synthetic dataset



• And of course, with lifespan cut-off, PD thresholding produces correct result

An example with disparate brightness



(a) $\hat{\beta} = (5,4)$ (b) $\beta(f_{185}^-) = (2,1)$ (c) $\beta(f_{66}^-) = (5,4)$

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An example that PD thresholding fails



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An example that PD thresholding fails



- Add "salt and pepper" noise which randomly resets some pixels to white or black
- The produced PD contain a lot of long-lived intervals due to the small white or black pixels



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- The bound $\parallel f g \parallel_\infty$ is determined by the maximum value difference on a single pixel
- With the "salt-and-pepper" noise, distance of the two functions (before and after adding noise) can be arbitrarily large
- So the two PDs are not stable anymore

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- Otsu : $\beta(f^-) = (206, 432)$ with threshold 134
- PD-based: $\beta(f^{-}) = (194, 143)$ with threshold 151
- With length cut-off, $\beta(f^-) = (149, 33)$

- One variation of the objective function is to replace $\Phi(t)$ with $\Phi_i(t)$ which is the same as $\Phi(t)$ by considering only the *i*-th PD
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- Then, we can assign a weight w_i to each dimension p indicating how one wants to stress this dimension

- One variation of the objective function is to replace $\Phi(t)$ with $\Phi_i(t)$ which is the same as $\Phi(t)$ by considering only the *i*-th PD
 - So $\Phi_i(t)$ only tries to optimize over the *i*-dimensional homology features
- Then, we can assign a weight w_i to each dimension p indicating how one wants to stress this dimension
- Then final objective function is a weighted sum of $\Phi_i(t)$:

$$T = \arg \max \sum_{i=0}^{\dim -1} w_i \Phi_i(t),$$



$$\Psi(t) = \frac{1}{\#\mathscr{P}(t) + 1} \sum_{(b,d)\in\mathscr{P}(t)} [(d-t)(t-b)]^p$$

• Take the p-th power for each term in $\Psi(t)$, to put greater emphasis on long intervals

$$\Psi(t) = \frac{1}{\#\mathscr{P}(t) + 1} \sum_{(b,d)\in\mathscr{P}(t)} [(d-t)(t-b)]^p$$