

VI - Statistical aspects of persistent homology

PSL Week - Topological Data Analysis

Abstract

We discuss how persistent homology behaves under random sampling. We highlight a notion of low intrinsic dimension called the (a, b) -standard assumption, and show how to leverage the stability of persistence, as well as elementary minimax theory, to study the problem of estimation of persistence diagrams.

Contents

1	Sampling assumptions and (a, b)-standard measures	1
1.1	Distance function and Hausdorff distance	1
1.2	(a, b) -standard measures	2
2	Hausdorff convergence of samples and consequences	3
2.1	A non-asymptotic bound	3
2.2	Plug-in estimation of persistence diagrams	4
3	Minimax risk and Le Cam's lemma	4
3.1	Risk and minimax risk	5
3.2	Total variation distance	5
3.3	Le Cam's two-point lemma	5
4	Minimax rates for persistence	6
4.1	Setup and upper bound	6
4.2	Lower bound via Le Cam	7

1 Sampling assumptions and (a, b) -standard measures

1.1 Distance function and Hausdorff distance

Let $K \subset \mathbb{R}^d$ be compact. We recall the *distance function* to K .

Definition 1.1 (Distance function). The distance to K is

$$\text{dist}(\cdot, \cdot) : \mathbb{R}^d \rightarrow [0, \infty), \quad \text{dist}(x, K) := \min_{p \in K} \|x - p\|.$$

Definition 1.2 (Offset). For $r > 0$, the r -offset (or *thickening*) of K is

$$K^r := \{x \in \mathbb{R}^d : \text{dist}(x, K) \leq r\} = \bigcup_{p \in K} \overline{B}_r(p).$$

Definition 1.3 (Hausdorff distance). Let A, B be compact subsets of \mathbb{R}^d . The *Hausdorff distance* between A and B is

$$d_H(A, B) := \min\{r \geq 0 : A \subset B^r \text{ and } B \subset A^r\}.$$

One can show the equivalent expression

$$d_H(A, B) = \sup_{x \in \mathbb{R}^d} |\text{dist}(x, A) - \text{dist}(x, B)|.$$

Intuitively, $d_H(A, B)$ is the smallest radius so that every point of A is within r of B and every point of B is within r of A (see Figure 1).

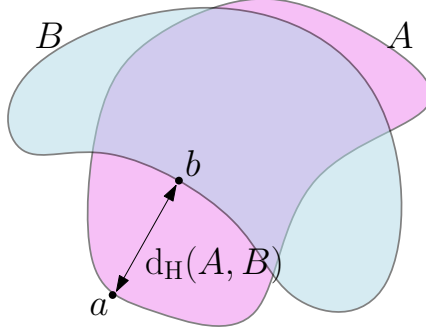


Figure 1: The Hausdorff distance between two subsets A and B of the plane. In this example, $d_H(A, B)$ is the distance between the point a in A which is the farthest from B and its nearest neighbor b on B .

1.2 (a, b) -standard measures

We now put a regularity condition on the sampling distribution.

Definition 1.4 ((a, b) -standard measure). Let P be a Borel probability measure on \mathbb{R}^d . We say that P is (a, b) -standard at scale r_0 if there exist constants $a > 0$, $b > 0$, $r_0 > 0$ such that for all $x \in \text{support}(P)$ and all $0 < r \leq r_0$,

$$P(B(x, r)) \geq ar^b.$$

Remark 1.5. Roughly speaking, a measure that is (a, b) -standard behaves on small scales like the b -dimensional Lebesgue measure:

- for $b = d$ and P having a density bounded below on its support, this condition is satisfied;
- b does not need to be an integer: this covers fractal-like supports (e.g. Cantor-type sets).

The exponent b plays the role of an *effective dimension*: balls of radius r carry at least a constant times r^b mass, so the support cannot be too thin.

To quantify how “massive” the support is, it is convenient to introduce covering and packing numbers.

Definition 1.6 (Covering and packing numbers). Let $K \subset \mathbb{R}^d$ be bounded and $r > 0$.

- An r -covering of K is a family of balls of radius r whose union contains K . The *covering number* is

$$\text{cov}(K, r) := \min \left\{ k : K \subset \bigcup_{i=1}^k B(x_i, r) \right\}.$$

- An r -packing of K is a family of disjoint balls of radius r with centres in K . The *packing number* is

$$\text{pack}(K, r) := \max \left\{ k : \exists x_1, \dots, x_k \in K, B(x_i, r) \text{ disjoint} \right\}.$$

Proposition 1.7 (Massiveness of (a, b) -standard measures). Let P be (a, b) -standard at scale r_0 and $K = \text{support}(P)$. Then there exists a constant $C_{a,b} > 0$ such that for all $r \leq r_0$,

$$\text{pack}(K, r) \leq \frac{1}{ar^b}, \quad \text{cov}(K, r) \leq \frac{C_{a,b}}{r^b}.$$

Idea. If $B(x_1, r), \dots, B(x_N, r)$ is a maximal packing, then these balls are disjoint and all contained in K^r , so

$$1 = P(\mathbb{R}^d) \geq \sum_{i=1}^N P(B(x_i, r)) \geq N ar^b,$$

which yields $N \leq 1/(ar^b)$. Using duality between coverings and packings (i.e. $\text{pack}(K, r) \leq \text{cov}(K, r) \leq \text{pack}(K, r/2)$) gives the covering bound. \square

Thus a (a, b) -standard measure has at most on the order of r^{-b} well-separated points at scale r , just like a b -dimensional cube $[0, 1]^b$.

2 Hausdorff convergence of samples and consequences

2.1 A non-asymptotic bound

Let P be (a, b) -standard at scale r_0 , and let X_1, \dots, X_n be i.i.d. with distribution P . We denote the sample by

$$\mathcal{X}_n := \{X_1, \dots, X_n\} \subset \mathbb{R}^d.$$

The following proposition shows that \mathcal{X}_n converges to the support of P in Hausdorff distance, with a rate governed by b .

Proposition 2.1 (Hausdorff convergence under (a, b) -standardness). *Let P be (a, b) -standard at scale r_0 , with compact support $K = \text{support}(P)$. Let \mathcal{X}_n be an i.i.d. sample from P . Then:*

(a) *There exist constants $C_{a,b,\alpha} > 0$ such that for any $\alpha > 0$ and all n large enough,*

$$\mathbb{P}\left(d_H(K, \mathcal{X}_n) > \left(C_{a,b,\alpha} \frac{\log n}{n}\right)^{1/b}\right) \leq n^{-\alpha}.$$

(b) *Equivalently, for any confidence level $\delta \in (0, 1)$ and any $r \leq 2r_0$, one has*

$$\mathbb{P}(d_H(K, \mathcal{X}_n) \leq r) \geq 1 - \delta$$

as soon as

$$n \geq \frac{C'_{a,b}}{r^b} \left(\log \frac{1}{r} + \log \frac{1}{\delta} \right).$$

Idea. Fix $r \leq 2r_0$ and consider a minimal $(r/2)$ -covering of K with $N = \text{cov}(K, r/2) \lesssim r^{-b}$ balls B_1, \dots, B_N of radius $r/2$. If $d_H(K, \mathcal{X}_n) > r$, one easily checks that at least one ball B_j contains no sample point. Since $P(B_j) \geq ar^b$, the probability that B_j is empty is at most $(1 - a(r/2)^b)^n \leq \exp(-an(r/2)^b)$. A union bound over all j then gives

$$\mathbb{P}(d_H(K, \mathcal{X}_n) > r) \leq N \exp(-an(r/2)^b) \lesssim r^{-b} \exp(-an(r/2)^b).$$

Optimizing in r yields the rate $r_n \asymp (\log n/n)^{1/b}$ and the stated bounds. \square

In words: for an (a, b) -standard measure, with n points we typically resolve the support down to a scale of order $(\log n/n)^{1/b}$ in Hausdorff distance.

2.2 Plug-in estimation of persistence diagrams

We now want to transfer the Hausdorff convergence of \mathcal{X}_n to convergence of persistence diagrams. Let (M, ρ) be a compact metric space, and let Filt be a filtration functor that associates to each compact subset $A \subset M$ a filtration of simplicial complexes $\text{Filt}(A)$ (e.g. the Vietoris–Rips or Čech filtration). Under mild assumptions, recall from Chapter III that persistent homology is *stable* with respect to perturbations of A in the Hausdorff metric.

Theorem 2.2 (Stability for spaces (informal)). *Let (M, ρ) be a compact metric space and $A, B \subset M$ compact. For a fixed homological degree k , let $D_k(A)$ and $D_k(B)$ denote the persistence diagrams of $H_k(\text{Filt}(A))$ and $H_k(\text{Filt}(B))$ (with coefficients in a field). Then,*

$$d_B(D_k(A), D_k(B)) \leq d_H(A, B),$$

where d_B is the bottleneck distance.

We now combine Proposition 2.1 and Theorem 2.2. Let (M, ρ) be a compact metric space and let μ be a Borel probability measure on M with compact support $X_\mu := \text{support}(\mu) \subset M$. Let X_1, \dots, X_n be i.i.d. with distribution μ and $\mathcal{X}_n := \{X_1, \dots, X_n\}$.

Definition 2.3 (Statistical model). Fix $a, b > 0$. We denote by $\mathcal{P}_{M,a,b}$ the collection of Borel probability measures μ on M such that:

- the support X_μ is compact in M ;
- μ is (a, b) -standard (with respect to ρ).

Theorem 2.4 (Upper bounds for persistence diagrams). *Assume $\mu \in \mathcal{P}_{M,a,b}$ and let $D_k(\mu)$ denote the k -th persistence diagram of $\text{Filt}(X_\mu)$, and $D_k(\mathcal{X}_n)$ that of $\text{Filt}(\mathcal{X}_n)$. Then:*

(a) For all $\varepsilon > 0$,

$$\mathbb{P}(d_B(D_k(\mu), D_k(\mathcal{X}_n)) > C\varepsilon) \leq \min\left(\frac{C'}{\varepsilon^b} \exp(-cn\varepsilon^b), 1\right),$$

where $C, C', c > 0$ depend only on the filtration and on a, b .

(b) For n large enough,

$$\sup_{\mu \in \mathcal{P}_{M,a,b}} \mathbb{E}_{\mu^n} [d_B(D_k(\mu), D_k(\mathcal{X}_n))] \leq C_{a,b} \left(\frac{\log n}{n}\right)^{1/b},$$

where $C_{a,b}$ depends only on a, b and the filtration.

Idea. For each μ , by stability for spaces,

$$d_B(D_k(\mu), D_k(\mathcal{X}_n)) \leq C d_H(X_\mu, \mathcal{X}_n).$$

Apply Proposition 2.1 with $K = X_\mu$, then take the supremum over $\mu \in \mathcal{P}_{M,a,b}$ and integrate the tail bound to control the expectation by using $\mathbb{E}[Y] = \int_0^\infty \mathbb{P}(Y > y) dy$ for a random variable $Y \geq 0$. \square

Thus the usual estimator $D_k(\mathcal{X}_n)$ is consistent, and its accuracy improves at the rate $(\log n/n)^{1/b}$, up to constants.

3 Minimax risk and Le Cam's lemma

As standard in statistical decision theory, we now turn to the question of optimality: Does there exist any better estimator than mine? Said otherwise: How well can *any* estimator do, in the worst case, over a given statistical model?

3.1 Risk and minimax risk

Let $(\mathcal{X}, \mathcal{A})$ be a measurable space (the observation space). A *statistical model* is a collection \mathcal{Q} of probability measures on $(\mathcal{X}, \mathcal{A})$.

We are interested in a parameter

$$\theta : \mathcal{Q} \rightarrow \Theta,$$

where (Θ, ρ) is a metric space (for us, Θ will be a space of persistence diagrams, and ρ the bottleneck distance).

An *estimator* is a measurable map

$$\hat{\theta}_n : \mathcal{X}^n \rightarrow \Theta,$$

applied to n i.i.d. observations $X_1, \dots, X_n \sim Q$.

Definition 3.1 (Risk and minimax risk). The *risk* of $\hat{\theta}_n$ at Q (for loss ρ) is

$$R(Q, \hat{\theta}_n) := \mathbb{E}_{Q^n} [\rho(\theta(Q), \hat{\theta}_n(X_1, \dots, X_n))].$$

The *minimax risk* over \mathcal{Q} is

$$R_n(\mathcal{Q}) := \inf_{\hat{\theta}_n} \sup_{Q \in \mathcal{Q}} R(Q, \hat{\theta}_n),$$

where the infimum is over all estimators $\hat{\theta}_n : \mathcal{X}^n \rightarrow \Theta$.

$R_n(\mathcal{Q})$ measures the best possible worst-case performance for the estimation problem (\mathcal{Q}, θ) at sample size n .

3.2 Total variation distance

We will use the *total variation* distance between probability measures.

Definition 3.2 (Total variation). Let Q, Q' be probability measures on $(\mathcal{X}, \mathcal{A})$. The *total variation* distance is

$$\text{TV}(Q, Q') := \sup_{A \in \mathcal{A}} |Q(A) - Q'(A)|.$$

If Q and Q' admit densities q, q' with respect to a reference measure ν , then

$$\text{TV}(Q, Q') = \frac{1}{2} \int_{\mathcal{X}} |q - q'| d\nu.$$

Remark 3.3. For product measures, one can show

$$\text{TV}(Q^n, Q'^n) \leq 1 - (1 - \text{TV}(Q, Q'))^n.$$

In particular, if $\text{TV}(Q, Q')$ is small, then $\text{TV}(Q^n, Q'^n)$ remains bounded away from 1 for moderate n .

3.3 Le Cam's two-point lemma

Le Cam's lemma is a simple but powerful way to get minimax lower bounds, by restricting attention to just two distributions $Q, Q' \in \mathcal{Q}$.

Lemma 3.4 (Le Cam). *Let \mathcal{Q} be a set of probability measures on $(\mathcal{X}, \mathcal{A})$, and $\theta : \mathcal{Q} \rightarrow \Theta$ a parameter, where (Θ, ρ) is a metric space. Then for any $Q, Q' \in \mathcal{Q}$,*

$$R_n(\mathcal{Q}) = \inf_{\hat{\theta}_n} \sup_{Q \in \mathcal{Q}} \mathbb{E}_{Q^n} \rho(\theta(Q), \hat{\theta}_n) \geq \frac{1}{2} \rho(\theta(Q), \theta(Q')) (1 - \text{TV}(Q, Q'))^n.$$

Proof. Fix any estimator $\hat{\theta}_n$, and any $Q, Q' \in \mathcal{Q}$. The key observation is to notice that

$$\sup_{Q \in \mathcal{Q}} \mathbb{E}_{Q^n} \rho(\theta(Q), \hat{\theta}_n) \geq \frac{1}{2} (\mathbb{E}_{Q^n} \rho(\theta(Q), \hat{\theta}_n) + \mathbb{E}_{Q'^n} \rho(\theta(Q'), \hat{\theta}_n)).$$

Then, changing the measure under which these two integral are taken and using the triangle inequality,

$$\rho(\theta(Q), \hat{\theta}_n) + \rho(\hat{\theta}_n, \theta(Q')) \geq \rho(\theta(Q), \theta(Q')).$$

This means that the estimator cannot simultaneously do very well under Q^n and under Q'^n if these two distributions are difficult to distinguish statistically. After that, we obtain

$$\sup_{Q \in \mathcal{Q}} \mathbb{E}_{Q^n} \rho(\theta(Q), \hat{\theta}_n) \geq \frac{1}{2} \rho(\theta(Q), \theta(Q')) (1 - \text{TV}(Q^n, Q'^n)),$$

and the proof is completed using a data-processing inequality for testing. \square

Informally: if two distributions in the model have parameters that are far apart, but are close in total variation, then any estimator must incur a nontrivial error on at least one of them.

4 Minimax rates for persistence

We now apply this framework to persistence diagrams of random point clouds sampled from (a, b) -standard measures.

4.1 Setup and upper bound

We consider the model $\mathcal{P}_{M,a,b}$ defined above, with parameter of interest

$$\theta(\mu) := D_k(\mu),$$

the k -th persistence diagram of $\text{Filt}(X_\mu)$, and loss

$$\rho(D, D') := d_B(D, D').$$

A natural estimator is

$$\hat{\theta}_n := D_k(\mathcal{X}_n),$$

the diagram built on the empirical point cloud.

Theorem 2.4 shows that this estimator satisfies

$$\sup_{\mu \in \mathcal{P}_{M,a,b}} \mathbb{E}_{\mu^n} d_B(D_k(\mu), D_k(\mathcal{X}_n)) \leq C_{a,b} \left(\frac{\log n}{n} \right)^{1/b}.$$

In particular,

$$R_n(\mathcal{P}_{M,a,b}) \leq C_{a,b} \left(\frac{\log n}{n} \right)^{1/b},$$

so the minimax risk cannot be worse than this rate.

4.2 Lower bound via Le Cam

We now sketch a lower bound, following a two-point argument. Assume that (M, ρ) is a metric space in which we can find a point $x \in M$ and a sequence $(x_n)_{n \geq 1} \subset M$ such that

$$\rho(x, x_n) \asymp (an)^{-1/b}.$$

(This holds in particular in \mathbb{R}^d with $b \leq d$, by choosing a grid of points with spacing of order $n^{-1/b}$.)

For each n , consider the two measures

$$\mu_0 = \delta_x, \quad \mu_{1,n} := \left(1 - \frac{1}{n}\right) \delta_x + \frac{1}{n} \delta_{x_n}.$$

One can check that both μ_0 and $\mu_{1,n}$ belong to $\mathcal{P}_{M,a,b}$ for suitable constants a, b (they are (a, b) -standard, since balls around x or x_n quickly contain all mass).

Let $D_k(\mu_0)$ and $D_k(\mu_{1,n})$ be the corresponding persistence diagrams (for a fixed filtration and degree k). Geometrically, μ_0 has a single-point support $\{x\}$, while $\mu_{1,n}$ has two points $\{x, x_n\}$ at distance $\rho(x, x_n) \asymp n^{-1/b}$.

- The bottleneck distance between $D_k(\mu_0)$ and $D_k(\mu_{1,n})$ is of order $\rho(x, x_n)$: the presence of the extra point x_n creates additional small features in the filtration at scale $\rho(x, x_n)$, so that

$$d_B(D_k(\mu_0), D_k(\mu_{1,n})) \gtrsim \rho(x, x_n) \asymp n^{-1/b}.$$

- The total variation distance between μ_0 and $\mu_{1,n}$ is exactly

$$\text{TV}(\mu_0, \mu_{1,n}) = \frac{1}{n},$$

so

$$\left(1 - \text{TV}(\mu_0, \mu_{1,n})\right)^n = \left(1 - \frac{1}{n}\right)^n \longrightarrow e^{-1}.$$

Applying Le Cam's lemma

$$R_n(\mathcal{P}_{M,a,b}) \geq \frac{1}{2} d_B(D_k(\mu_0), D_k(\mu_{1,n})) \left(1 - \text{TV}(\mu_0, \mu_{1,n})\right)^n$$

and using the two bullets above, we obtain

$$R_n(\mathcal{P}_{M,a,b}) \gtrsim n^{-1/b}.$$

Theorem 4.1 (Minimax lower bound for persistence). *Under the assumptions above, there exists a constant $c > 0$ such that for all n large enough,*

$$R_n(\mathcal{P}_{M,a,b}) \geq c n^{-1/b}.$$

Combining this with the upper bound from Theorem 2.4, we obtain that the estimator $D_k(\mathcal{X}_n)$ is *minimax optimal up to logarithmic factors*:

$$c n^{-1/b} \lesssim R_n(\mathcal{P}_{M,a,b}) \leq C_{a,b} \left(\frac{\log n}{n}\right)^{1/b}.$$

Remark 4.2. The log factor in the upper bound comes from the union bound over an r -covering of the support and is typical in nonparametric estimation with (a, b) -standard assumptions. For $b = 1$, it cannot actually be removed.