

Statistical Inference for Topological Data Analysis

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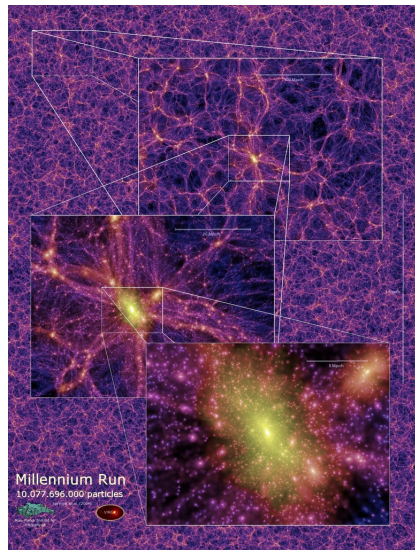
Introduction

Homology and Persistent Homology

Statistical Inference for Persistent Homology

Reference

Topological structures in the data provide information.



¹ http://www.mpa-garching.mpg.de/galform/virgo/millennium/poster_half.jpg

Persistent Homology: observe topological structure with multi resolutions.



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- ▶ Georges Seurat, A Sunday afternoon on the island of La Grande Jatte (Un dimanche après-midi à l'Île de la Grande Jatte)



Statistic Inference for Topological Data Analysis is explored.

- ▶ Introduction to Topological Data Analysis
 - ▶ Computational Topology: An Introduction (Edelsbrunner, Harer, 2010)
 - ▶ Topological Data Analysis (Wasserman, 2016)
 - ▶ An Introduction to Topological Data Analysis: Fundamental and Practical Aspects for Data Scientists (Chazal, Michel, 2021)
- ▶ Statistical Inference for Persistent Homology
 - ▶ Confidence sets for persistence diagrams (Fasy, Lecci, Rinaldo, Wasserman, Balakrishnan, Singh, 2014)




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The number of holes is used to summarize topological features.

- ▶ Geometrical objects:
 - ▶ A, B, C, D, E, F, G, H, I, J, K, L, M, O, P, Q, R, S, T, U, V, W, X, Y, Z, IJ
- ▶ The number of holes of different dimensions is considered.
 1. β_0 = # of connected components 
 2. β_1 = # of loops (holes inside 1-dim sphere) 
 3. β_2 = # of voids (holes inside 2-dim sphere) 

Example : Objects are classified by homologies.

1. $\beta_0 = \#$ of connected components



2. $\beta_1 = \#$ of loops (holes inside 1-dim sphere)

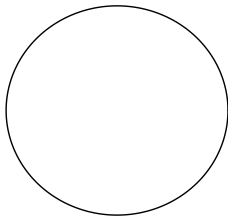


$\beta_0 \setminus \beta_1$	0	1	2
1	C, E, F, G, H, I, J, K, L, M, N, S, T, U, V, W, X, Y, Z	A, D, O, P, Q, R	B
2	IJ		

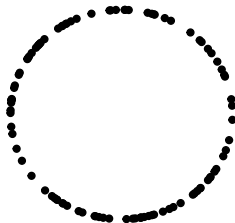
Homology of finite sample is different from homology of underlying manifold, hence it cannot be directly used for the inference.

- ▶ When analyzing data, we prefer robust features where features of the underlying manifold can be inferred from features of finite samples.
- ▶ Homology is not robust:

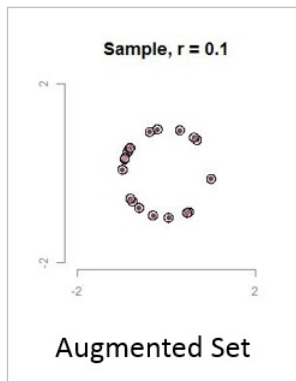
Underlying circle: $\beta_0 = 1, \beta_1 = 1$



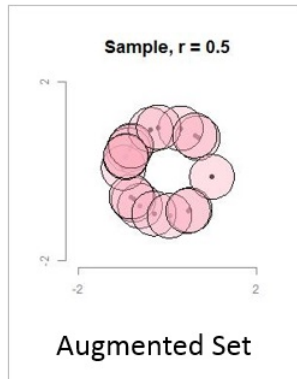
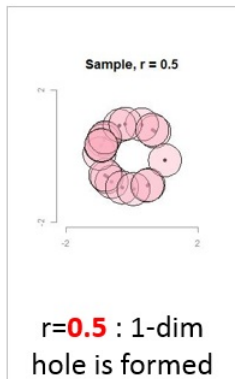
100 samples: $\beta_0 = 100, \beta_1 = 0$



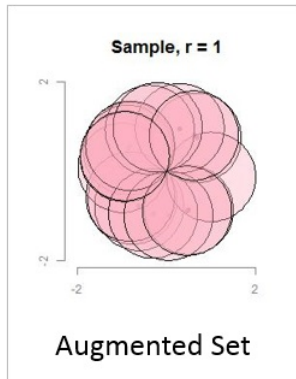
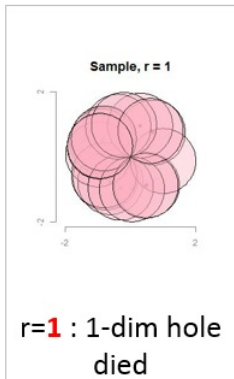
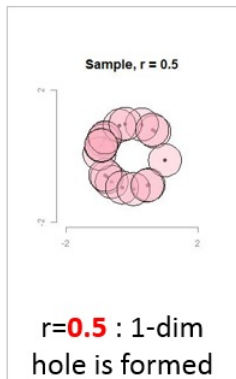
Persistent homology computes homologies on collection of sets, and tracks when topological features are born and when they die.



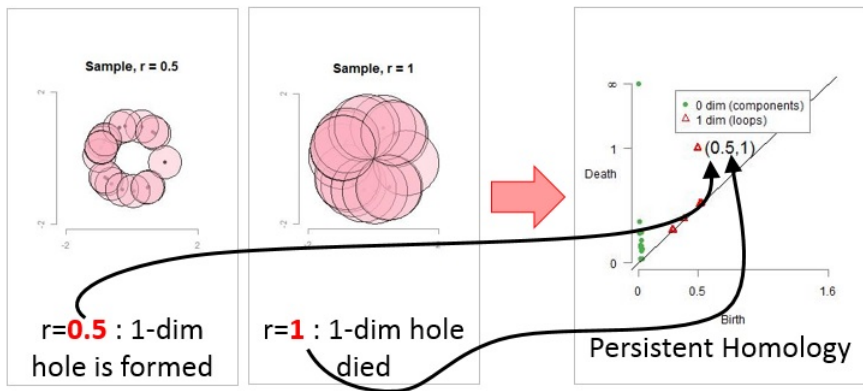
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We rely on the superlevel sets of the kernel density estimator to extract topological information of the underlying distribution.

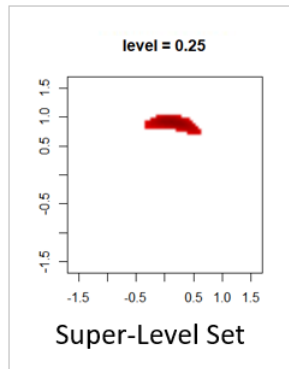
- ▶ The kernel density estimator is

$$\hat{p}_h(x) = \frac{1}{nh^d} \sum_{i=1}^n K\left(\frac{x - X_i}{h}\right).$$

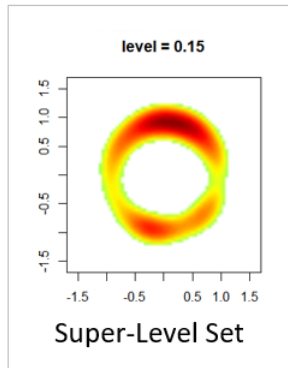
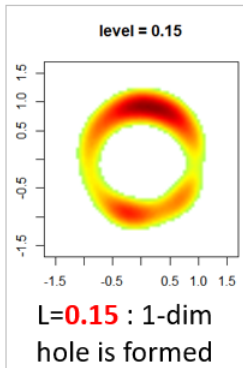
- ▶ We look at superlevel sets of the kernel density estimator as

$$\{x \in \mathbb{R}^d : \hat{p}_h(x) \geq L\}_{L>0}.$$

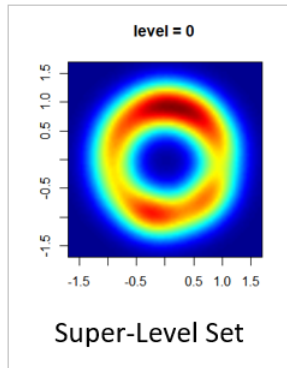
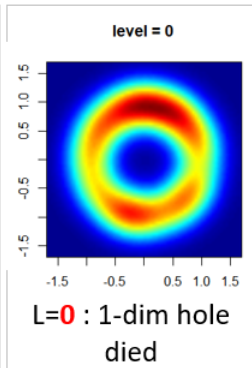
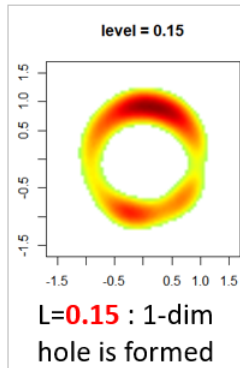
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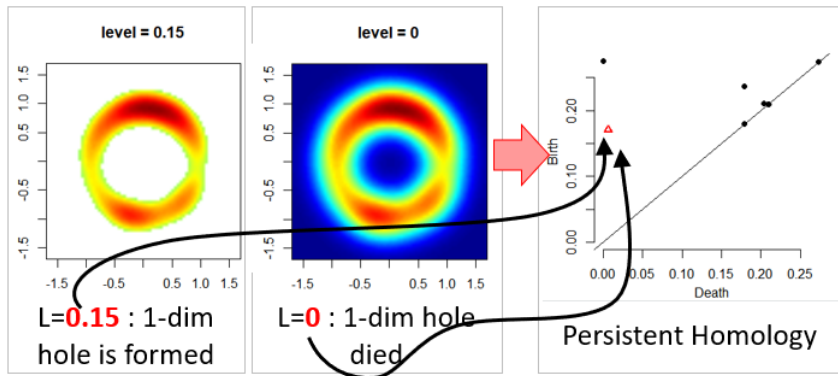
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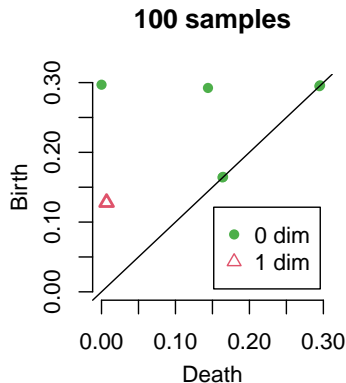
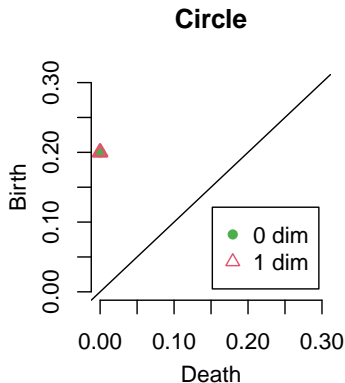
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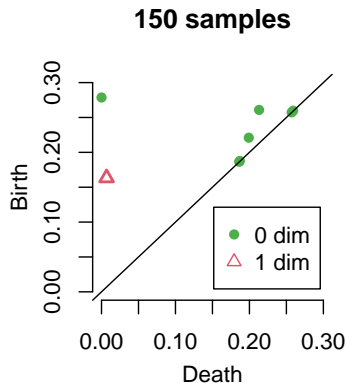
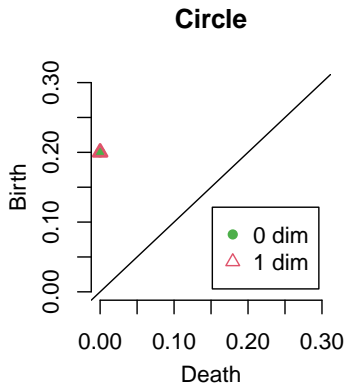
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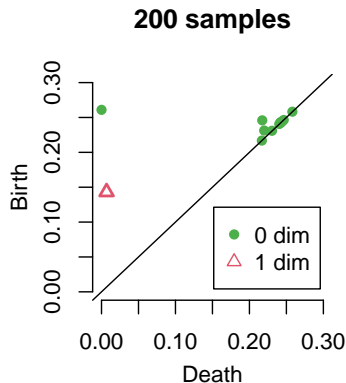
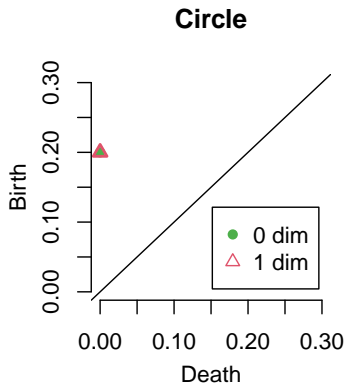
Persistent homology of the underlying manifold can be inferred from persistent homology of finite samples.



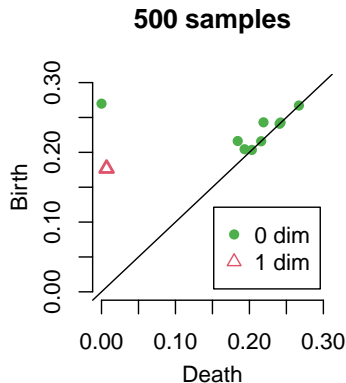
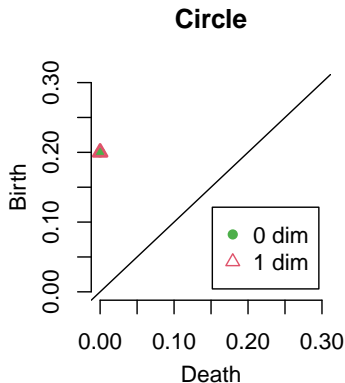
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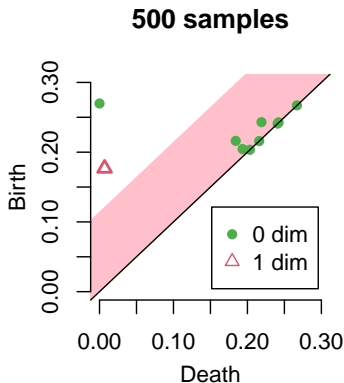
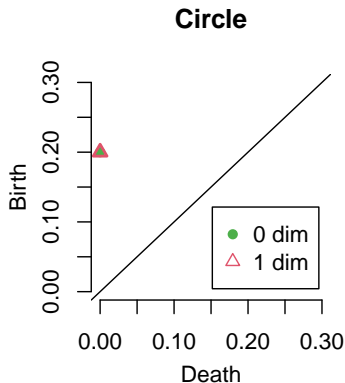
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Statistically significant homological features can be distinguished from statistically insignificant ones.

- Confidence sets for persistence diagrams (Fasy, Lecci, Rinaldo, Wasserman, Balakrishnan, Singh, 2014)



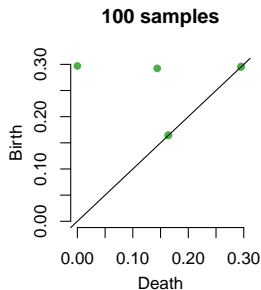
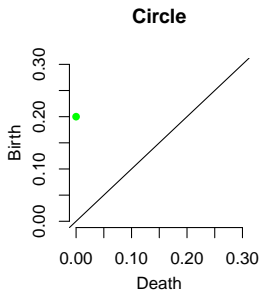
Bottleneck distance gives a metric on the space of persistent homology.

Definition

Let D_1, D_2 be multiset of points. Bottleneck distance is defined as

$$W_\infty(D_1, D_2) = \inf_{\gamma} \sup_{x \in D_1} \|x - \gamma(x)\|_\infty,$$

where γ ranges over all bijections from D_1 to D_2 .



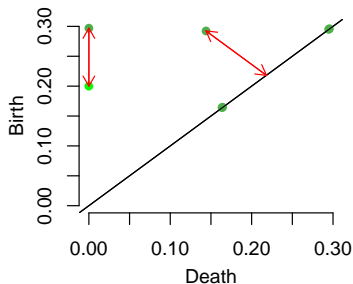
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$$\sup_{x \in D_1} \|x - \gamma_1(x)\|_\infty = 0.1$$

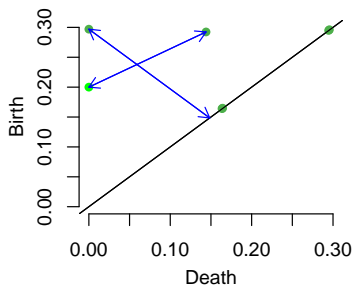
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$$\sup_{x \in D_1} \|x - \gamma_2(x)\|_\infty = 0.15$$

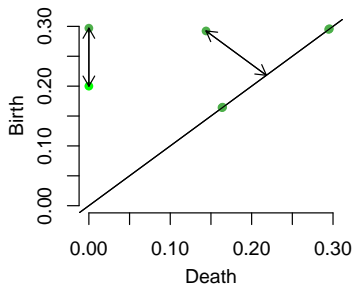
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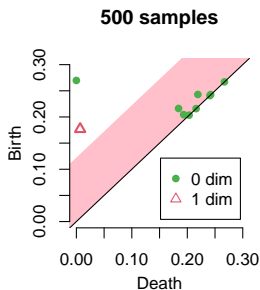
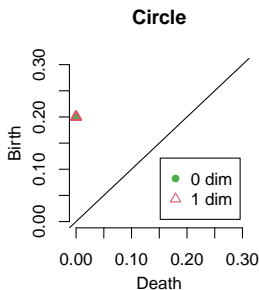


$$\inf_{\gamma} \sup_{x \in D_1} \|x - \gamma(x)\|_\infty = 0.1$$

Confidence band for persistent homology separates homological signal from homological noise.

Let $Dgm(M)$ and $Dgm(X)$ be persistent homologies of the manifold M and the data X , respectively. Given the significance level $\alpha \in (0, 1)$, $(1 - \alpha)$ confidence band $c_n = c_n(X)$ is a random variable satisfying

$$\mathbb{P}(W_\infty(Dgm(M), Dgm(X)) \leq c_n) \geq 1 - \alpha.$$



Confidence band for the persistent homology can be computed using the bootstrap algorithm.

1. Given a sample $X = \{x_1, \dots, x_n\}$, compute the kernel density estimator \hat{p}_h .
2. Draw $X^* = \{x_1^*, \dots, x_n^*\}$ from $X = \{x_1, \dots, x_n\}$ (with replacement), and compute $\theta^* = \sqrt{nh^d} \|\hat{p}_h^*(x) - \hat{p}_h(x)\|_\infty$, where \hat{p}_h^* is the density estimator computed using X^* .
3. Repeat the previous step B times to obtain $\theta_1^*, \dots, \theta_B^*$
4. Compute $\hat{z}_\alpha = \inf \left\{ q : \frac{1}{B} \sum_{j=1}^B I(\theta_j^* \geq q) \leq \alpha \right\}$
5. The $(1 - \alpha)$ confidence band for $\mathbb{E}[\hat{p}_h]$ is $\left[\hat{p}_h - \frac{\hat{z}_\alpha}{\sqrt{nh^d}}, \hat{p}_h + \frac{\hat{z}_\alpha}{\sqrt{nh^d}} \right]$.

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- Frédéric Chazal and Bertrand Michel. An introduction to topological data analysis: Fundamental and practical aspects for data scientists. *Frontiers Artif. Intell.*, 4:667963, 2021. doi: 10.3389/frai.2021.667963. URL <https://doi.org/10.3389/frai.2021.667963>.
- Frédéric Chazal, Vin de Silva, Marc Glisse, and Steve Oudot. The structure and stability of persistence modules. *arXiv preprint arXiv:1207.3674*, 2012.
- Frédéric Chazal, Brittany Terese Fasy, Fabrizio Lecci, Bertrand Michel, Alessandro Rinaldo, and Larry Wasserman. Robust topological inference: Distance-to-a-measure and kernel distance. *Technical Report*, 2014.
- Herbert Edelsbrunner and John L. Harer. *Computational topology*. American Mathematical Society, Providence, RI, 2010. ISBN 978-0-8218-4925-5. doi: 10.1090/mbk/069. URL <https://doi.org/10.1090/mbk/069>. An introduction.

Reference II

Brittany Terese Fasy, Fabrizio Lecci, Alessandro Rinaldo, Larry Wasserman, Sivaraman Balakrishnan, and Aarti Singh. Confidence sets for persistence diagrams. *Ann. Statist.*, 42(6):2301–2339, 2014. ISSN 0090-5364. doi: 10.1214/14-AOS1252. URL <https://doi.org/10.1214/14-AOS1252>.

Larry Wasserman. Topological data analysis, 2016.

Thank you!

Persistent Homology

Bottleneck distance can be controlled by the corresponding distance on functions: Stability Theorem.

Theorem

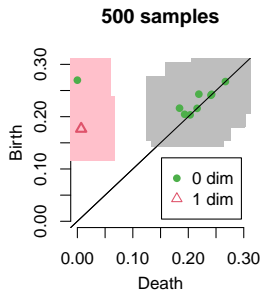
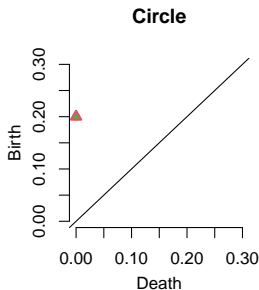
[Edelsbrunner and Harer, 2010][Chazal, de Silva, Glisse, and Oudot, 2012] Let \mathbb{X} be finitely triangulable space and $f, g : \mathbb{X} \rightarrow \mathbb{R}$ be two continuous functions. Let $Dgm(f)$ and $Dgm(g)$ be corresponding persistence diagrams. Then

$$W_{\infty}(Dgm(f), Dgm(g)) \leq \|f - g\|_{\infty}.$$

Confidence set for the persistent homology is a random set containing the persistent homology with high probability.

Let $Dgm(M)$ and $Dgm(X)$ be persistent homologies of the manifold M and the data X , respectively. Given the significance level $\alpha \in (0, 1)$, $(1 - \alpha)$ confidence set $\{D \in Dgm : W_\infty(Dgm(X), D) \leq c_n\}$ is a random set satisfying

$$\mathbb{P}(Dgm(M) \in \{D \in Dgm : W_\infty(Dgm(X), D) \leq c_n\}) \geq 1 - \alpha.$$



Confidence band for the persistent homology can be obtained by the corresponding confidence band for functions.

From Stability Theorem, $\mathbb{P}(\|f_M - f_X\| \leq c_n) \geq 1 - \alpha$ implies

$$\mathbb{P}(d_B(Dgm(f_M), Dgm(f_X)) \leq c_n) \geq \mathbb{P}(\|f_M - f_X\|_\infty \leq c_n) \geq 1 - \alpha,$$

so the confidence band of corresponding functions f_M can be used for confidence band of persistent homologies $Dgm(f_M)$.

Confidence band for the persistent homology can be computed using the bootstrap algorithm.

Bootstrap algorithm can be applied to persistent homology.

- ▶ for the case of kernel density estimator in Fasy et al. [2014],
- ▶ for the case of distance to measure and kernel distance in Chazal et al. [2014].

