9					
Per:	sistence paths and signature				
	tures in topological data analysi				
(TDA)					
by	Chevyrer, Nanda and Oberhauser				
	vectorization				
	TDA ~ ML				

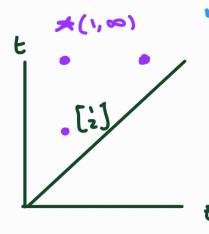
- Outline: 1 Persistent Homology
 - 2 Path embeddings
 - 3 Path signatures
 - 4 Results
- 1 Let X_{ξ} tell be a family of topsp. such that X_{i} C^{3} X_{ij} if i < j

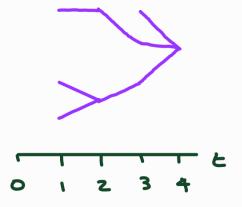
Let H(-) be your favorite topological invaviant (e.g. connected components)

Persistent homology consists on Heeping track of $H(X_t)$ as t varies.

Example:

Homm... the farther from diagonal the longer they persist.





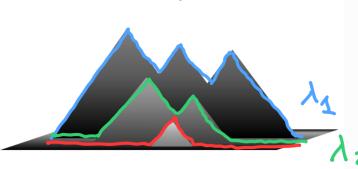
Persistence Diagram (PD)

Rmh: PD are bad for stats/ML!
integrated landscope emb.

2 From PD to Pather

* They describe 4 options, I will four on 1.

Pers landscape



 $\lambda = \begin{bmatrix} \lambda_1 \\ \lambda_2 \end{bmatrix} : [0, T] \longrightarrow \mathbb{R}^3$

aha! a (piece wise linear) path! $z_{iL}(t) = \int_{0}^{t} \lambda$

Met _ @ PD _ @ Banach

The path sig of
$$\lambda = \begin{bmatrix} \lambda_1 \\ \lambda_n \end{bmatrix}$$
 is

$$S(\lambda) = (S_0(\lambda), S_1(\lambda), \dots)$$

$$S_{\Lambda}(\lambda) = \int_{0}^{\text{end}} d\lambda_{\Lambda} e_{\Lambda} + \dots + \int_{0}^{\text{end}} d\lambda_{n} e_{n}$$

$$S_{2}(\lambda) = \underbrace{S}_{i_{1},i_{2} \in \{1,...,n\}} \underbrace{\int_{0}^{e_{1}} d\lambda_{i_{1}}(x) d\lambda_{i_{2}}(y)}_{i_{1}\otimes e_{i_{2}}} e_{i_{1}\otimes e_{i_{2}}}$$

Properties

I.
$$S(\lambda) = S(\lambda)$$
 iff λ and λ are tree-like equivalent

II. S(-) is reparametrization invariant

III. If
$$\lambda$$
 is 'a segment' then
$$\lambda = \xi \cdot \sigma$$

$$S(\lambda) = (1, \sigma, \frac{\sigma \sigma^2}{2!}, \frac{\sigma \sigma^3}{3!}, \dots)$$
[think 'moments']

Behaves well with concat. $S(\Gamma * \lambda) = S(\Gamma) \otimes S(\lambda)$.

AlgTop, SDE, Data (Chen 60's) (Lyons 90's) (mow)

Bar = PD

THEOREM 5. Define

$$\Phi: \mathbf{Bar}/\iota \to \mathbf{T}(V), \quad B \mapsto S \circ \iota(B).$$

On each compact subset $K \subset \mathbf{Bar}/\iota$, the map Φ has the following properties.

(1) (**Universal**) Let $f: K \to \mathbb{R}$ be continuous. For each $\epsilon > 0$, there exists ℓ in $\bigoplus_{m \geq 0} (V')^{\otimes m}$ (the dual space of the tensor algebra) such that

$$\sup_{B\in K}|f(B)-\langle\Phi(B),\ell\rangle|<\epsilon.$$

(2) (**Characteristic**) Denoting by M the set of Borel probability measures on K, the map

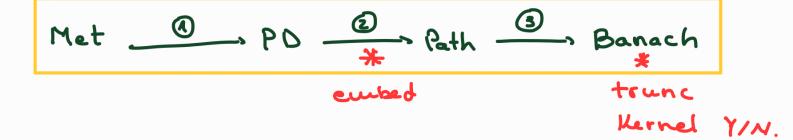
$$\mathcal{M} \to \mathbf{T}(V), \quad \mu \mapsto \mathbb{E}_{B \sim \mu} \left[\Phi(B) \right]$$

is injective.

(3) (**Kernelized**) Suppose further that V is a Hilbert space. Then the map

$$k: K \times K \to \mathbb{R}, \quad k(B, B') = \langle \Phi(B), \Phi(B') \rangle$$

defines a bounded, continuous kernel⁶ which is universal for the space of continuous *functions* $C(K,\mathbb{R})$ *and characteristic for Borel probability measures on* K.



 98.1 ± 0.7

Method **Textures Orbits** Shapes 96.8 ± 1.0 94.6 ± 1.3 95.8 ± 1.6 $k_{\rm SW}$ 93.7 ± 1.0 99.86 ± 0.21 90.3 ± 2.3 Φ_{PI} 90.4 ± 1.5 96.6 ± 0.9 92.7 ± 1.5 $k_{\rm E}$ k_{χ} 94.9 ± 0.6 NA 92.4 ± 3.0 k_{β} 97.8 ± 0.2 NA 93.0 ± 3.0 $\Phi_{\rm E}$ 88.1 ± 0.8 98.1 ± 1.0 95.0 ± 0.9 92.9 ± 0.7 98.8 ± 0.6 98.0 ± 1.1 Φ_{χ} 97.7 ± 0.8

 96.6 ± 0.6

 Φ_{β}

TABLE 1. Mean accuracy (\pm standard deviation).

TABLE 2. Best truncation level <i>M</i> .					
		Textures	Orbits	Shapes	
	k_{E}	2	2	2	
	k_{χ}	3	NA	2	
	k_{β}	3	NA	2	
	$\Phi_{\rm E}$	5	8	4	
	Φ_{χ}	8	7	7	
	k_{E} k_{χ} k_{β} Φ_{E} Φ_{χ}	6	8	5	

Shameless

Discrete signature tensors for persistence landscapes

VINCENZO GALGANO, HEATHER A. HARRINGTON, DANIEL TOLOSA

Abstract

Signature tensors of paths are a versatile tool for data analysis and machine learning. Recently, they have been applied to persistent homology, by embedding barcodes into spaces of paths. Among the different path embeddings, the persistence landscape embedding is injective and stable, however it loses injectivity when composed with the signature map. Here we address this by proposing a discrete alternative. The critical points of a persistence landscape form a time-series, of which we compute the discrete signature. We call this association the discrete landscape feature map (DLFM). We give results on the injectivity, stability and computability of the DLFM. We apply it to a knotted protein dataset, capturing sequence similarity and knot depth with statistical significance.

Keywords: persistent homology, barcodes, persistence landscapes, feature maps, time-series, time warping, path signatures, tensors, knotted proteins, vectorisation.

MSC2020 codes: 55N31, 68T09, 46B85.

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Introduction

As the scale and complexity of biological data increases, problems such as interpretation, classification and quantification require advanced mathematical methods for investigation. Applying persistent homology from topological data analysis to biological data provides a geometric interpretation of the shapes of data. Here, we propose an alternative approach combining persistent homology and nonlinear algebra.