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# **Topological Data Analysis** with the Gudhi library

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Topological Data Analysis is:

a mathematically grounded framework...

 $H_k = Z_k / B_k$ 

...that applies to a wide variety of data sets...



... for a wide variety of tasks.



Mapper: exploratory data analysis









Persistence diagrams: machine learning

Its main goal is to compute the topology of spaces (number and sizes of connected components, loops, cavities, etc) from samplings...



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...and to summarize this info into descriptors suitable for data science

#### The Gudhi library





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conda:  ${\sim}250~000$  total downloads pip:  ${\sim}34~000$  downloads over the last 6 months



I. Turn datasets into simplicial complexes II. Compute and compare persistence diagrams III. Feed / regularize ML models w/ topology I. Turn datasets into simplicial complexes II. Compute and compare persistence diagrams III. Feed / regularize ML models w/ topology

**Def:** Given a point cloud  $P = \{P_1, \ldots, P_n\} \subset \mathbb{R}^d$ , its Čech complex of radius r > 0 is the abstract simplicial complex C(P, r) s.t. vert(C(P, r)) = P and

 $\sigma = [P_{i_0}, P_{i_1}, \dots, P_{i_k}] \in C(P, r) \quad \text{iif} \quad \cap_{j=0}^k B(P_{i_j}, r) \neq \emptyset.$ 

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In [ ]: torus = gudhi.read\_points\_from\_off\_file(off\_file='datasets/tore3D\_1307.off')
ac = gudhi.AlphaComplex(points=torus)
st = ac.create\_simplex\_tree()

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#### Input:

- topological space  $\boldsymbol{X}$
- continuous function  $f: X \to Y$  (99% of the time  $Y = \mathbb{R}^D$ )
- cover  ${\mathcal I}$  of  $\operatorname{im}(f)$  by open intervals:  $\operatorname{im}(f) \subseteq \bigcup_{I \in {\mathcal I}} I$

Method:

- Compute *pullback cover*  $\mathcal{U}$  of X:  $\mathcal{U} = \{f^{-1}(I)\}_{I \in \mathcal{I}}$
- $\bullet$  Refine  ${\mathcal U}$  by separating each of its elements into its various connected components in  $X\to$  connected cover  ${\mathcal V}$
- The Mapper is the *nerve* of  $\mathcal{V}$ :
  - 1 vertex per element  $V \in \mathcal{V}$
  - 1 edge per intersection  $V \cap V' \neq \emptyset$ ,  $V,V' \in \mathcal{V}$
  - 1 k-simplex per (k+1)-fold intersection  $\bigcap_{i=0}^{k} V_i \neq \emptyset, V_0, \cdots, V_k \in \mathcal{V}$

Input:

- point cloud  $P \subseteq X$  with metric  $d_P$
- continuous function  $f: \textbf{\textit{P}} \rightarrow \mathbb{R}$
- cover  ${\mathcal I}$  of  $\operatorname{im}(f)$  by open intervals:  $\operatorname{im} f \subseteq \bigcup_{I \in {\mathcal I}} I$

Method:

- Compute *pullback cover*  $\mathcal{U}$  of P:  $\mathcal{U} = \{f^{-1}(I)\}_{I \in \mathcal{I}}$
- Refine  $\mathcal{U}$  by separating each of its elements into its various clusters, as identified by a clustering algorithm  $\rightarrow$  connected cover  $\mathcal{V}$
- The Mapper is the *nerve* of  $\mathcal{V}$ : intersections are assessed by the
  - 1 vertex per element  $V \in \mathcal{V}$
- presence of common data points
- 1 edge per intersection  $V \cap V' \neq \emptyset$ ,  $V,V' \in \mathcal{V}$
- 1 k-simplex per (k+1)-fold intersection  $\bigcap_{i=0}^k V_i \neq \emptyset$ ,  $V_0, \cdots, V_k \in \mathcal{V}$



In [22]: cover\_complex = MapperComplex(
 input\_type='point cloud', min\_points\_per\_node=0,
 clustering=None, N=100, beta=0., C=10,
 filter\_bnds=None, resolutions=[20,2], gains=None, verbose=verbose)

In [23]: \_ = cover\_complex.fit(X, filters=filt2d, colors=filt2d)





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#### Computation with filtrations and matrix reduction

Algorithms for computing the homology groups of a simplicial complex work by *decomposing* it with a so-called *filtration*.

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**Def:** A filtered simplicial complex S is a family  $\{S_u\}_{u \in \mathbb{R}}$  of subcomplexes of some fixed simplicial complex S s.t.  $S_a \subseteq S_b$  for any  $a \leq b$ .

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**Def:** The persistence barcode (resp. diagram) D is a set of points in the plane (resp. intervals) encoding the topological features that appeared and disappeared in the filtration.

















Persistence of sublevel sets of function

 $H_0$  (connected components)










#### $H_0$ (connected components)

 $\mathbb{R} \land \qquad \text{When two components merge, stop the} \\ \text{bar of the most recent one ($ *elder rule* $).}$ 









## Persistence of images





 $H_1$  (loops)





```
In []: st = get_simplex_tree_from_faces(faces)
filtration = geodesic_distances(base_vertex)
for v in range(len(vertices)):
    st.assign_filtration([v], filtration[v])
st.make_filtration_non_decreasing()
st.persistence()
dgm = st.persistence_intervals_in_dimension(1)
```







**Thm:**  $d_b(D(f), D(g)) \le ||f - g||_{\infty}$ 



# Stability and distance between PDs



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#### **Thm:** $d_b(D(f), D(g)) \le ||f - g||_{\infty}$

```
In []: BD = gudhi.representations.BottleneckDistance(epsilon=.001)
BD.fit([st1.persistence_intervals_in_dimension(1)])
bd = BD.transform([st2.persistence_intervals_in_dimension(1)])
WD = gudhi.representations.WassersteinDistance(internal_p=2, order=2)
WD.fit([st1.persistence_intervals_in_dimension(1)])
wd = WD.transform([st2.persistence_intervals_in_dimension(1)])
```

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# Persistence diagrams and ML



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0

0

 $\mathcal{H}$ 

0

- Dim. red. (PCA, MDS, UMAP, t-SNE)
- Clustering (DBSCAN, K-means, etc.)

#### Etc.

[Persistence Images: A Stable Vector Representation of Persistent Homology, Adams et al., JMLR, 2017]









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In [20]:	<pre>from sklearn.preprocessing from sklearn.pipeline from sklearn.svm from sklearn.ensemble from sklearn.neighbors</pre>	<pre>import MinMaxScaler import Pipeline import SVC import RandomForestClassifier import KNeighborsClassifier</pre>				
	<pre># Definition of pipeline pipe = Pipeline([("Separator", gd.representations.DiagramSelector(limit=np.inf, point_type="finite")),</pre>					
	<pre># Parameters of pipeline. The param = [{"Scaler_use":     "TDA":     "TDAbandwidth"     "TDAnum_direct"     "Estimator":</pre>	<pre>is is the place where you specify the methods you want to use to handle diagrams     [False],     [gd.representations.SlicedWassersteinKernel()], ': [0.1, 1.0], tions": [20],     [SVC(kernel="precomputed", gamma="auto")]},</pre>				
	{"Scaleruse": "TDA": "TDAbandwidth" "TDAweight": "Estimator":	<pre>[False], [gd.representations.PersistenceWeightedGaussianKernel()], [0.1, 0.01], [lambda x: np.arctan(x[1]-x[0])], [SVC(kernel="precomputed", gamma="auto")]},</pre>				
	<pre>{"Scaleruse":     "TDA":     "TDAresolution     "TDAbandwidth'     "Estimator":</pre>	<pre>[True], [gd.representations.PersistenceImage()], "": [5,5], [6,6] ], ": [0.01, 0.1, 1.0, 10.0], [SVC()]},</pre>				
	{"Scaleruse": "TDA": "TDAresolution "Estimator":	<pre>[True], [gd.representations.Landscape()], [100], [RandomForestClassifier()]},</pre>				
	<pre>{"Scaler_use":     "TDA":     "TDAepsilon":     "Estimator": ]</pre>	[ <b>False</b> ], [gd.representations.BottleneckDistance()], [0.1], [KNeighborsClassifier(metric="precomputed")]}				



[PersLay: A Neural Network Layer for Persistence Diagrams and New Graph Topological Signatures, C., Chazal, Ike, Lacombe, Royer, Umeda, AISTATS, 2019]



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Weight function learnt

A persistence diagram D is made of all  $(\mathcal{F}(\sigma_+), \mathcal{F}(\sigma_-)) \in \mathbb{R}^2$  where  $\sigma_+$  (resp.  $\sigma_-$ ) is positive (resp. negative), and  $\mathcal{F}$  is the filtration function.

Thus we can define the gradient of a point  $p = (\mathcal{F}(\sigma_+), \mathcal{F}(\sigma_-)) \in D$  as

 $\nabla p = [\nabla \mathcal{F}(\sigma_+), \nabla \mathcal{F}(\sigma_-)]$ 

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Ex: VR filtration parametrized by a point cloud

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```
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```

In []: optimizer = tensorflow.keras.optimizers.SGD(learning\_rate=0.1)
for epoch in tqdm(range(200+1)):
 with tensorflow.GradientTape() as tape:
 dgm = gudhi.tensorflow.RipsLayer(maximum\_edge\_length=1., homology\_dimensions=[1]).call(X)[0][0]
 # Opposite of the squared distances to the diagonal
 persistence\_loss = -tensorflow.math.reduce\_sum(tf.square(.5\*(dgm[:,1]-dgm[:,0])))
 # Unit square regularization
 regularization = tensorflow.reduce\_sum(tf.maximum(tf.abs(X)-1, 0))
 loss = persistence\_loss + regularization
 gradients = tape.gradient(loss, [X])
 # We also apply a small random noise to the gradient to ensure convergence
 np.random.seed(epoch)
 gradients[0] = gradients[0] + numpy.random.normal(loc=0., scale=.001, size=gradients[0].shape)

```
optimizer.apply_gradients(zip(gradients, [X]))
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**Ex:** images filtered by a direction parameterized by angle



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Dataset	Baseline	Before	After	Difference	Dataset	Baseline	Before	After	Difference
vs01	100.0	61.3	99.0	+37.6	vs26	99.7	98.8	98.2	-0.6
vs02	99.4	98.8	97.2	-1.6	vs28	99.1	96.8	96.8	0.0
vs06	99.4	87.3	98.2	+10.9	vs29	99.1	91.6	98.6	+7.0
vs09	99.4	86.8	98.3	+11.5	vs34	99.8	99.4	99.1	-0.3
vs16	99.7	89.0	97.3	+8.3	vs36	99.7	99.3	99.3	-0.1
vs19	99.6	84.8	98.0	+13.2	vs37	98.9	94.9	97.5	+2.6
vs24	99.4	98.7	98.7	0.0	vs57	99.7	90.5	97.2	+6.7
vs25	99.4	80.6	97.2	+16.6	vs79	99.1	85.3	96.9	+11.5



# What's next?

• Representative cycles


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- Representative cycles
- Zigzag persistence



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- Multi-parameter persistence



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Thanks!!