

# Topological Autoencoders

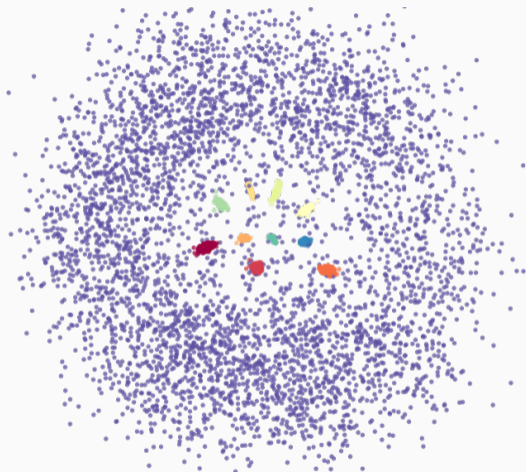
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Machine Learning and Computational Biology Group, ETH Zurich

# Motivation

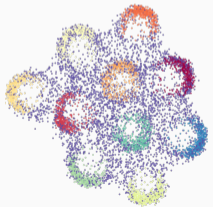


Representation of our data, but in *100 dimensional space*

# Motivation - Dimensionality reduction



PCA



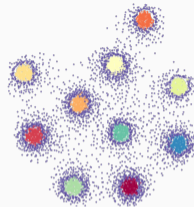
UMAP

## Issues

Most methods preserve connectivity at *local* scales

## Goal

We want to preserve connectivity at *multiple* scales

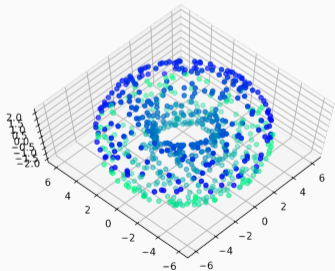
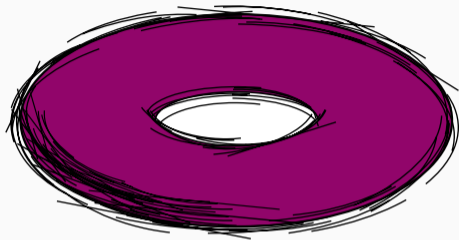


t-SNE



Autoencoder

# Topology - The study of connectivity



## Betti numbers characterize topological spaces

- $\beta_0$  connected components
- $\beta_1$  cycles
- $\beta_2$  voids

## Issues

- Great for manifolds (which are usually **unknown**)
- But instead *approximated* via samples
- Topology on samples is **noisy**

# Persistent Homology - Topology at multiple scales

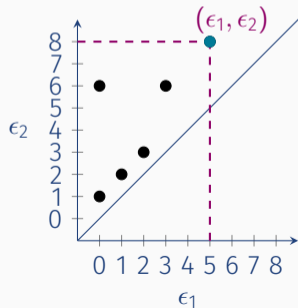
**Vietoris-Rips Complex:** Calculate neighbourhood graph (simplicial complex for higher dimensions) for all weight thresholds and keep track of the appearance and disappearance of topological features.

Filtration:

$$\emptyset = K_0 \subseteq K_1 \subseteq \dots \subseteq K_{n-1} \subseteq K_n = K$$

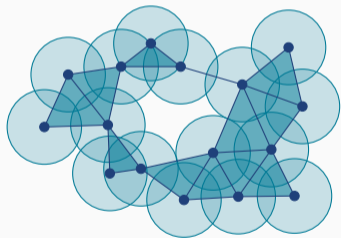


$$E := \{ (u, v) \mid \text{dist}(p_u, p_v) \leq \epsilon \}$$



# Persistent Homology - Topology at multiple scales

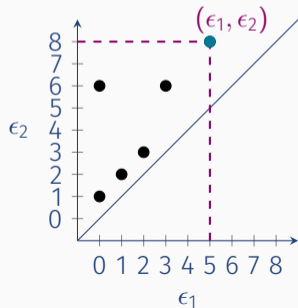
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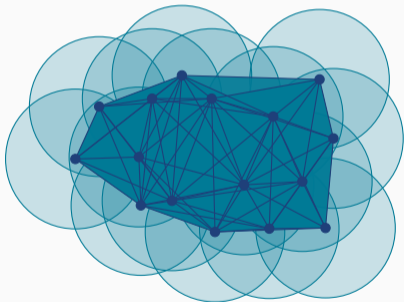
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# Persistent Homology - Topology at multiple scales

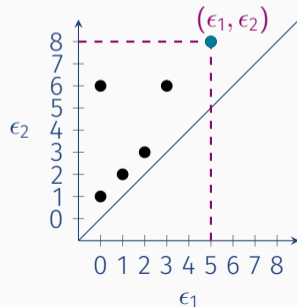
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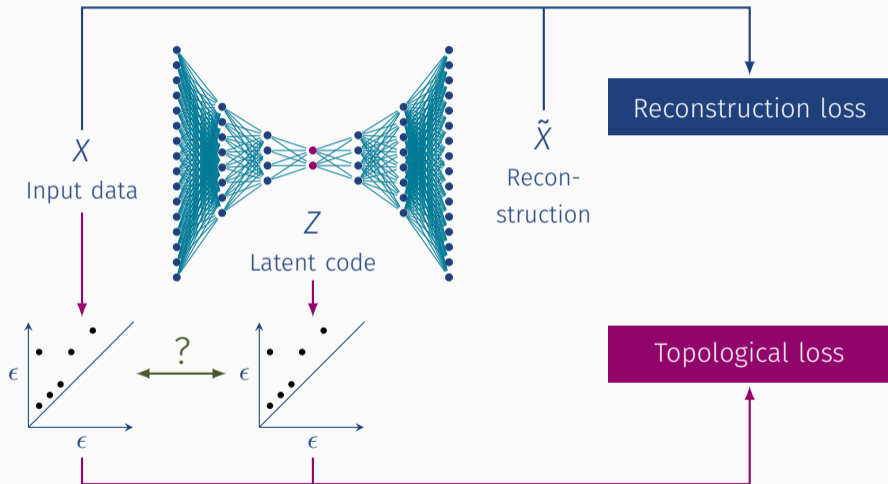
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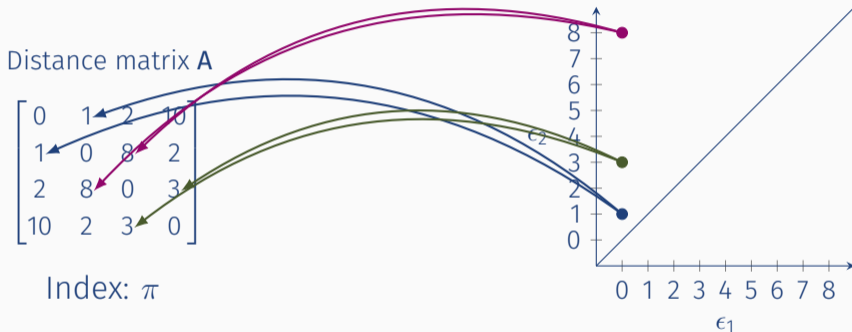
# Method - Overview





# Distance matrix and relation to persistence diagrams

While the persistence computation is an inherently discrete process, we can nevertheless compute gradients due to one key observation.



## Insight

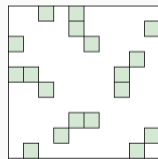
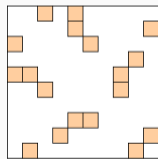
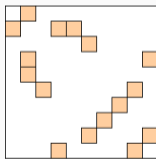
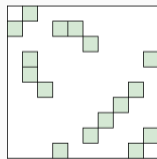
We can map all Persistent Homology computations of Flag complexes to individual edges of the distance matrix!

# Topological loss term

$$\mathcal{L}_t = \mathcal{L}_{\mathcal{X} \rightarrow \mathcal{Z}} + \mathcal{L}_{\mathcal{Z} \rightarrow \mathcal{X}}$$

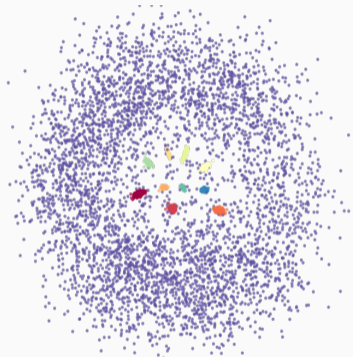
$$\mathcal{L}_{\mathcal{X} \rightarrow \mathcal{Z}} := \frac{1}{2} \|\mathbf{A}^{\mathcal{X}}[\pi^{\mathcal{X}}] - \mathbf{A}^{\mathcal{Z}}[\pi^{\mathcal{X}}]\|^2$$

$$\mathcal{L}_{\mathcal{Z} \rightarrow \mathcal{X}} := \frac{1}{2} \|\mathbf{A}^{\mathcal{Z}}[\pi^{\mathcal{Z}}] - \mathbf{A}^{\mathcal{X}}[\pi^{\mathcal{Z}}]\|^2$$



# Experiments

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SPHERES



MNIST

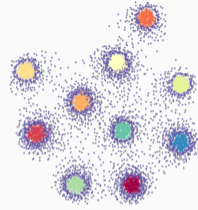


FASHION-MNIST

# Spheres



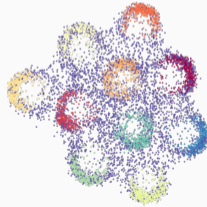
PCA



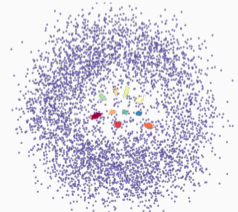
t-SNE



Autoencoder



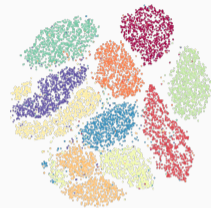
UMAP



Topo-AE



PCA



t-SNE



Autoencoder



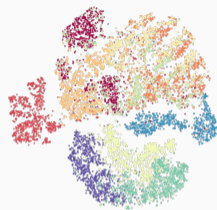
UMAP



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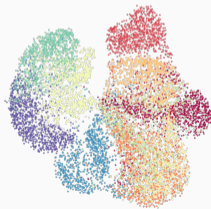
PCA



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Autoencoder



UMAP



Topo-AE

- Novel method for **preserving topological information** of the input space in dimensionality reduction
- Under weak theoretical assumptions our loss term is **differentiable** and allowing the training of MLPs via backpropagation
- Our method **was uniquely able to capture spatial relationships** of nested high-dimensional spheres



For further information and more theory

Check out our paper on ArXiv!



<https://arxiv.org/abs/1906.00722>

# Appendix

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# Bound of bottleneck distance between persistence diagrams on subsampled data

## Theorem

Let  $X$  be a point cloud of cardinality  $n$  and  $X^{(m)}$  be one subsample of  $X$  of cardinality  $m$ , i.e.  $X^{(m)} \subseteq X$ , sampled without replacement. We can bound the probability of the persistence diagrams of  $X^{(m)}$  exceeding a threshold in terms of the bottleneck distance as

$$\mathbb{P}\left(d_b\left(\mathcal{D}^X, \mathcal{D}^{X^{(m)}}\right) > \epsilon\right) \leq \mathbb{P}\left(d_H\left(X, X^{(m)}\right) > 2\epsilon\right),$$

where  $d_H$  refers to the Hausdorff distance between the point cloud and its subsample.

# Expected value of Hausdorff distance

## Theorem

Let  $\mathbf{A} \in \mathbb{R}^{n \times m}$  be the distance matrix between samples of  $X$  and  $X^{(m)}$ , where the rows are sorted such that the first  $m$  rows correspond to the columns of the  $m$  subsampled points with diagonal elements  $a_{ij} = 0$ . Assume that the entries  $a_{ij}$  with  $i > m$  are random samples following a distance distribution  $F_D$  with  $\text{supp}(f_D) \in \mathbb{R}_{\geq 0}$ . The minimal distances  $\delta_i$  for rows with  $i > m$  follow a distribution  $F_\Delta$ . Letting  $Z := \max_{1 \leq i \leq n} \delta_i$  with a corresponding distribution  $F_Z$ , the expected Hausdorff distance between  $X$  and  $X^{(m)}$  for  $m < n$  is bounded by:

$$\mathbb{E} \left[ d_H(X, X^{(m)}) \right] = \mathbb{E}_{Z \sim F_Z} [Z] \leq \int_0^{+\infty} \left( 1 - F_D(z)^{(n-1)} \right) dz \leq \int_0^{+\infty} \left( 1 - F_D(z)^{m(n-m)} \right) dz$$

## Explicit Gradient Derivation

Letting  $\theta$  refer to the parameters of the *encoder*, we have

$$\begin{aligned}\frac{\partial}{\partial \theta} \mathcal{L}_{X \rightarrow Z} &= \frac{\partial}{\partial \theta} \left( \frac{1}{2} \|\mathbf{A}^X[\pi^X] - \mathbf{A}^Z[\pi^X]\|^2 \right) \\ &= -(\mathbf{A}^X[\pi^X] - \mathbf{A}^Z[\pi^X])^\top \left( \frac{\partial \mathbf{A}^Z[\pi^X]}{\partial \theta} \right) \\ &= -(\mathbf{A}^X[\pi^X] - \mathbf{A}^Z[\pi^X])^\top \left( \sum_{i=1}^{|\pi^X|} \frac{\partial \mathbf{A}^Z[\pi^X]_i}{\partial \theta} \right),\end{aligned}$$

where  $|\pi^X|$  denotes the cardinality of a persistence pairing and  $\mathbf{A}^Z[\pi^X]_i$  refers to the  $i$ th entry of the vector of paired distances.

# Density distribution error

## Definition (Density distribution error)

Let  $\sigma \in_{>0}$ . For a finite metric space  $\mathcal{S}$  with an associated distance  $\text{dist}(\cdot, \cdot)$ , we evaluate the density at each point  $x \in \mathcal{S}$  as

$$f_{\sigma}^{\mathcal{S}}(x) := \sum_{y \in \mathcal{S}} \exp\left(-\sigma^{-1} \text{dist}(x, y)^2\right),$$

where we assume without loss of generality that  $\max \text{dist}(x, y) = 1$ . We then calculate  $f_{\sigma}^X(\cdot)$  and  $f_{\sigma}^Z(\cdot)$ , normalise them such that they sum to 1, and evaluate

$$\text{KL}_{\sigma} := \text{KL}\left(f_{\sigma}^X \parallel f_{\sigma}^Z\right), \quad (1)$$

i.e. the Kullback–Leibler divergence between the two density estimates.

# Quantification of performance

Data set	Method	KL <sub>0.01</sub>	KL <sub>0.1</sub>	KL <sub>1</sub>	$\ell$ -MRRE	$\ell$ -Cont	$\ell$ -Trust	$\ell$ -RMSE	Data MSE
SPHERES	Isomap	0.181	<b>0.420</b>	<b>0.00881</b>	<b>0.246</b>	<b>0.790</b>	<b>0.676</b>	10.4	-
	PCA	0.332	0.651	0.01530	0.294	0.747	0.626	11.8	0.9610
	TSNE	<b>0.152</b>	0.527	0.01271	<u><b>0.217</b></u>	0.773	<u><b>0.679</b></u>	<b>8.1</b>	-
	UMAP	0.157	0.613	0.01658	0.250	0.752	0.635	<b>9.3</b>	-
	AE	0.566	0.746	0.01664	0.349	0.607	0.588	13.3	<u><b>0.8155</b></u>
	TopoAE	<u><b>0.085</b></u>	<u><b>0.326</b></u>	<u><b>0.00694</b></u>	0.272	<u><b>0.822</b></u>	0.658	13.5	<b>0.8681</b>
F-MNIST	PCA	<u><b>0.356</b></u>	<u><b>0.052</b></u>	<u><b>0.00069</b></u>	0.057	0.968	0.917	<b>9.1</b>	0.1844
	TSNE	0.405	0.071	0.00198	<u><b>0.020</b></u>	0.967	<b>0.974</b>	41.3	-
	UMAP	0.424	0.065	0.00163	0.029	<u><b>0.981</b></u>	0.959	<b>13.7</b>	-
	AE	0.478	0.068	0.00125	<b>0.026</b>	0.968	<u><b>0.974</b></u>	20.7	<u><b>0.1020</b></u>
	TopoAE	<b>0.392</b>	<b>0.054</b>	<b>0.00100</b>	0.032	<b>0.980</b>	0.956	20.5	<b>0.1207</b>
MNIST	PCA	0.389	0.163	0.00160	0.166	0.901	0.745	<b>13.2</b>	0.2227
	TSNE	<u><b>0.277</b></u>	<b>0.133</b>	0.00214	<u><b>0.040</b></u>	0.921	<u><b>0.946</b></u>	22.9	-
	UMAP	<b>0.321</b>	0.146	0.00234	<b>0.051</b>	<u><b>0.940</b></u>	<b>0.938</b>	<b>14.6</b>	-
	AE	0.620	0.155	<b>0.00156</b>	0.058	0.913	0.937	18.2	<u><b>0.1373</b></u>
	TopoAE	0.341	<u><b>0.110</b></u>	<u><b>0.00114</b></u>	0.056	<b>0.932</b>	0.928	19.6	<b>0.1388</b>

## Quantification of performance - 2

Data set	Method	$KL_{0.01}$	$KL_{0.1}$	$KL_1$	$\ell$ -MRRE	$\ell$ -Cont	$\ell$ -Trust	$\ell$ -RMSE	Data MSE
CIFAR	PCA	<b>0.591</b>	<b>0.020</b>	<u>0.00023</u>	0.119	<u>0.931</u>	0.821	<u>17.7</u>	0.1482
	TSNE	0.627	0.030	0.00073	<u>0.103</u>	0.903	<b>0.863</b>	<b>25.6</b>	-
	UMAP	0.617	0.026	0.00050	0.127	0.920	0.817	33.6	-
	AE	0.668	0.035	0.00062	0.132	0.851	<u>0.864</u>	36.3	<b>0.1403</b>
	TopoAE	<u>0.556</u>	<u>0.019</u>	<b>0.00031</b>	<b>0.108</b>	<b>0.927</b>	0.845	37.9	<u>0.1398</u>