#### TARDIS: Topological Algorithm for Robust Discovery of Singularities

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### Topological Singularity Detection at Multiple Scales

Julius Von Rohrscheidt, Bastian Rieck Proceedings of the 40th International Conference on Machine Learning, PMLR 202:35175-35197, 2023.

#### Abstract

The manifold hypothesis, which assumes that data lies on or close to an unknown manifold of low intrinsic dimension, is a staple of modern machine learning research. However, recent work thas shown that real-world data exhibits distinct non-manifold structures, i.e. singularities, that can lead to erroneous findings. Detecting such singularities is therefore crucial as a percensor to interpolation and inference tasks. We address this issue by developing a topological transwork that (i) quantifies the docal intrinsic dimension, and (ii) yields a Lucidicity score for sassessing the "manifoldness" of a point along multiple scales. Our approach identifies singularities of complex spaces, while also capturing singular structures and local geometric complexity in image data.

#### https://proceedings.mlr.press/v202/von-rohrscheidt23a.html

#### Manifolds in a nutshell



A **manifold** is a space that resembles Euclidean space locally, i.e. *every* point admits a neighbourhood that looks like a Euclidean ball.

#### Manifolds in a nutshell



- Manifolds can encode complex global behaviour
- However, locally they look 'trivial'.

- Manifolds are widely studied objects in mathematics
- In Data Science, most non-linear dimensionality reduction techniques (UMAP, t-SNE, ...) make use of the manifold hypothesis:

The **manifold hypothesis** assumes that the given data lies on a lower dimensional manifold.

 Performance of these algorithms depends on the correctness of the manifold hyothesis.

#### Singularities

A **singularity** is a point in a space that violates the assumption of being *locally Euclidean*.

A singular space is a space that may admit singularities.



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### Why singularities?

- Recently, Brown et. al.<sup>1</sup> found evidence that popular datasets (MNIST, FashionMNIST, ...) do not satisfy the manifold hypothesis.
- Moreover, Perea et. al.<sup>2</sup> showed empirically that manifold learning algorithms fail in general, when the underlying data does not stem from a manifold.

 $\Rightarrow$  Let's test the manifold hypothesis!

<sup>1</sup>Brown, Bradley CA, et al. "The Union of Manifolds Hypothesis and its Implications for Deep Generative Modelling." arXiv preprint arXiv:2207.02862 (2022). <sup>2</sup>Mike, Joshua Lee, and Jose Perea. "TALLEM: Topological Assembly of Locally Euclidean Models." 2022 Spring Western Sectional Meeting. AMS, 2022.

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•  $c^{\circ}S^{1} \cong D^{2}$  (2-dimensional disk) •  $c^{\circ}(S^{1} \sqcup S^{1}) \cong$  double cone

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- $X X_{n-2}$  is dense in X.
- For each point x ∈ X<sub>n-k</sub> − X<sub>n-k-1</sub>, there exists an open neighborhood U of x in X and a compact (PL) stratified pseudomanifold L of dimension k − 1 and a (PL) homeomorphism

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(which is stratum-preserving.)

 For a point *x* ∈ *X*, its *i*-th local homology *H<sub>i</sub>(X, X − x)* captures homological information of an infinitesimal small neighborhood of *x*, relative to an infinitesimal punctured neighbourhood of *x* (in *X*).

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- Let *X* be a (stratified) pseudomanifold and  $x \in X$ . Then *x* has a distinguished neighborhood  $U \cong \mathbb{R}^k \times c^{\circ}L$ , where *L* is called the **link** of *x*.

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- If  $U \cong c^{\circ}L$ , one can show that

$$H_i(X, X - x) = \tilde{H}_{i-1}(L)$$

for all  $i \ge 0$ .

- Let X be a (stratified) pseudomanifold and  $x \in X$ . Then x has a distinguished neighborhood  $U \cong \mathbb{R}^k \times c^{\circ}L$ , where L is called the **link** of x.
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- The motivation to use local homology for singularity detection stems from the following fact:
- If  $U \cong c^{\circ}L$ , one can show that

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• In particular, if X = M is a manifold of dimension *n*, one obtains

$$H_i(M, M-x) = \tilde{H}_{i-1}(S^{n-1}) = \begin{cases} \mathbb{Z}, i = n \\ 0, i \neq n \end{cases}$$

- As we have already seen, manifolds are characterised by a local property.
- **Idea:** Test the 'manifoldness' of each point in the data space, individually.



- From your given dataset X, choose a point  $x \in X$ .
- For two fixed radius parameters r < s, let B<sup>s</sup><sub>r</sub>(x) denote the set of data points with distance to x at least r, and at most s.
- Let  $\mathcal{V}(B_r^s(x), t)$  denote the *Vietoris-Rips* construction w.r.t.  $B_r^s(x)$  at filtration step *t*.

















#### Back to data: Persistent homology

Given a finite metric space (X, d), the Vietoris–Rips complex at step t is defined as the abstract simplicial complex V(X, t), in which an abstract k-simplex (x<sub>0</sub>,..., x<sub>k</sub>) of points in X is spanned if and only if d(x<sub>i</sub>, x<sub>j</sub>) ≤ t for all 0 ≤ i ≤ j ≤ k.

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- This leads to  $H_i(\mathcal{V}(\mathbb{X}, t_1)) \to H_i(\mathcal{V}(\mathbb{X}, t_2))$  for any  $t_1 \leq t_2$

The *i*-th **persistent homology (PH)** of  $\mathbb{X}$  with respect to the Vietoris-Rips construction is defined to be the collection of all these *i*-th homology groups, together with the respective induced maps between them, and denoted by  $PH_i(\mathcal{V}(\mathbb{X}, \bullet))$ 

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- We denote the resulting persistence information by  $PH(\mathcal{V}(B_r^s(x), \bullet)).$
- The idea is now to compare the topological information of  $B_r^s(x)$ with the one of a known Euclidean model space  $EucB_r^s(x)$ :  $d_B^{r,s} := d_B \left[ PH(\mathcal{V}(B_r^s(x), \bullet)), PH(\mathcal{V}(EucB_r^s(x), \bullet)) \right]$
- Finally, we vary *r* and *s* and take the average of these distances:  $\mathfrak{E}(x) := \frac{1}{C} \sum_{(r,s)} d_{B}^{r,s}$

 $\mathfrak{E}(x)$  is called the **Euclidicity** of *x* (w.r.t. the ambient data).

#### Euclidicity enjoys theoretical guarantees



 $\mathfrak{E}(x)$  is called the **Euclidicity** of *x* (w.r.t. the given data  $\mathbb{X}$ ).

When the dataset X is sampled from a manifold,  $\mathfrak{E}(x)$  will be small, for any point *x*.

#### Euclidicity tends to zero for 'manifold points'

#### Theorem

Let  $M \subset \mathbb{R}^N$  be a smooth n-dimensional manifold and let  $\mathbb{X} \subset M$  be a finite sample of size  $S := |\mathbb{X}|$ . For a given  $\epsilon > 0$ , sufficiently large S and a point  $x \in \mathbb{X}$ , there exists  $s_{\epsilon} > 0$  that (up to a constant) only depends on  $\epsilon$ , such that  $\mathfrak{E}(x)$  is bounded above by  $\epsilon$ , for any radius configuration with maximum outer radius at most  $s_{\epsilon}$ .

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However,  $\mathfrak{E}(x)$  will usually *not* tend to zero when x is a singularity! (Homology of the link of x is usually different to the homology of a sphere.)

### Euclidicity detects singularities



Input space with singularities



Euclidicity

Euclidicity scores of *singular* points are higher than for *non-singular* points.



#### Real-world data admits singular regions

 The following are embeddings of tokens of a Large Language Model (RoBERTa)



#### Euclidicity detects non-linearities in image datasets

- By flattening images, we obtain point cloud representations of image datasets in order to calculate Euclidicity scores.
- It turns out that high Euclidicity values correspond to images that possess a high degree of geometric complexity **inside of** the image.



Figure 6: Left to right: samples images exhibiting low, median, and high Euclidicity, respectively.

#### Misclassified samples admit higher Euclidicity scores

We trained a simple neural network to analyse the Euclidicity scores of misclassified vs. correctly classified samples.

**Misclassified** samples admit significantly **higher** Euclidicity scores than correctly classified samples.

Acknowledgement: This experiment was conducted together with Francesco Conti (Università di Pisa)



Figure 8: A comparison of Euclidicity scores for misclassified and correctly classified samples in two image data sets.

 We have already seen that if X = M is a manifold of dimension n and x ∈ M, its local homology reads

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• This means that we can deduce the intrinsic dimension of *M*, by looking at its local homology!



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- This maximum homology degree is *n*.
- In practice, data may be noisy. We therefore only consider homology generators that exceed a certain persistence threshold.
- Finally, we vary *r* and *s*, and average the resulting dimension estimates. This is called the **persistent intrinsic dimension** (**PID**) of *x*.

#### Theorem

Let  $M \subset \mathbb{R}^N$  be an n-dimensional compact smooth manifold and let  $\mathbb{X} := \{x_1, \ldots, x_S\}$  be a collection of uniform samples from M. For a sufficiently large S, PID calculates the correct intrinsic dimension of M in a small neighbourhood around x, for any  $x \in M$ . Moreover, this neighbourhood can be chosen arbitrarily small by increasing S.



- Dimensionality estimates: twoNN vs. PID.
- PID is more nuanced in capturing changes in dimensionality, assigning 1 to almost all points of the circle, i.e. S<sup>1</sup>, while highlighting that points closer to S<sup>2</sup> exhibit an increase in dimensionality.

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- The given framework can be used to estimate the intrinsic dimension around the data point, locally.
- Experiments suggest that singularities have meaning: can we regularise for singularities, how?

# TARDIS: Topological Algorithms for Robust Discovery of Singularities



https://github.com/aidos-lab/TARDIS