An Introduction to the Mapper Algorithm and Its Open-Source Implementations

Youjia Zhou

Agenda

Part I

- The Mapper Algorithm
- Kepler Mapper

Part II

- Giotto-TDA
- Mapper Interactive

Part

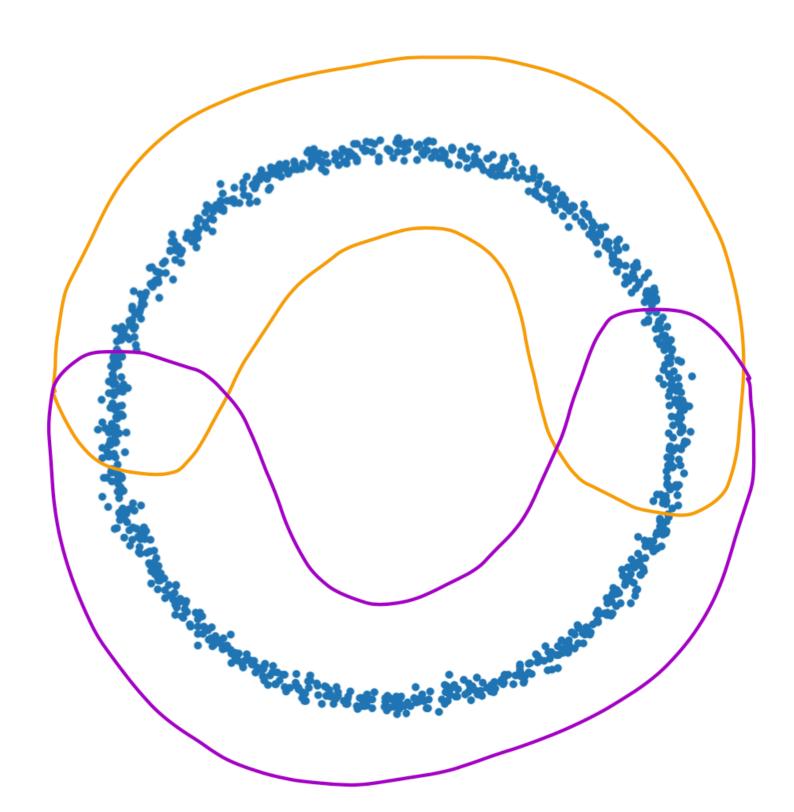
Mapper Algorithm

- First introduced by Singh et al. in 2007 [1].
- A popular framework from topological data analysis for extracting topological summaries of high-dimensional datasets.
- "Partial clustering of the data guided by a set of functions defined on the data", which captures both **clusters** and **cluster relations**.
- Combining dimensionality reduction with graph visualization
- Qualitative understanding of high-dimensional point cloud data through visualization

Mapper Algorithm: Cover and Nerve

Given a high-dimensional point cloud $\mathbb{X} \subset \mathbb{R}^d$

• A cover of \mathbb{X} is defined as a set of open sets in \mathbb{R}^d , $\mathscr{U} = \{U_i\}_{i \in I}$ such that $\mathbb{X} \subset \bigcup_{i \in I} U_i$ (I being the index set).

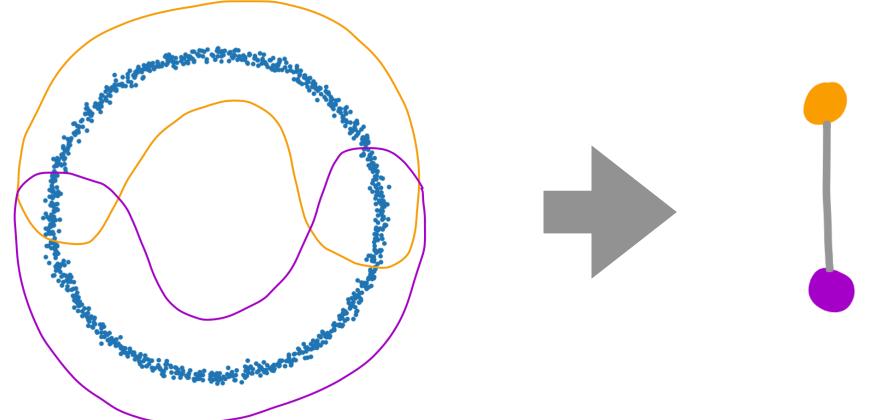


Mapper Algorithm: Cover and Nerve

• Given a cover $\mathcal{U} = \{U_i\}_{i \in I}$ of \mathbb{X} , let $\mathcal{N}(\mathcal{U})$ denote the simplicity complex that corresponds to the nerve of the cover \mathcal{U}

$$\mathcal{N}(\mathcal{U}) = \{ \sigma \subset I \mid \cap_{i \in \sigma} U_i \neq \emptyset \}$$

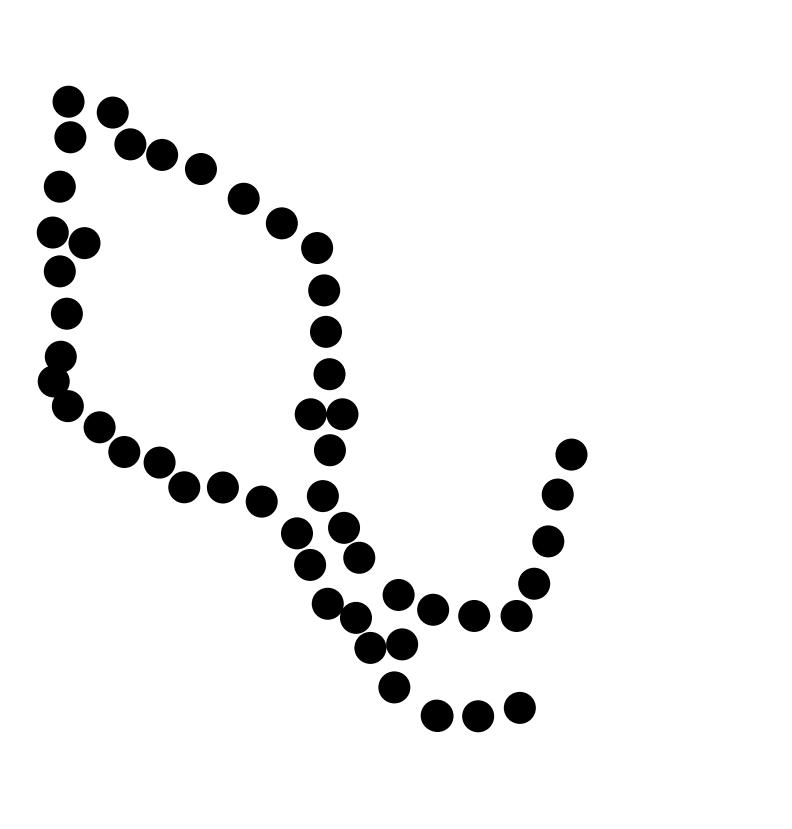
- The 1D nerve of \mathcal{U} , denoted as $\mathcal{N}_1(\mathcal{U})$, is a graph.
- Each node $i \in I$ in $\mathcal{N}_1(\mathcal{U})$ represents a cover element U_i , and there is an edge between $i,j \in I$ if $U_i \cap U_j$ is nonempty.



Mapper Algorithm: Filter Function and Clustering

- In the classic mapper construction [1], a set of scalar functions $f: \mathbb{X} \to \mathbb{R}$ (referred to as the filter functions) is used to guide obtaining a cover.
- Given a cover $\mathscr{V} = \{V_k\}$ $(1 \le k \le n)$ of $f(\mathbb{X}) \subset \mathbb{R}$, such that $f(\mathbb{X}) \subseteq \cap_k V_k$, we obtain a cover \mathscr{U} of \mathbb{X} by considering the clusters induced by points in $f^{-1}(V_k)$ for each V_k as cover elements.
- The 1D nerve of \mathcal{U} , denoted as $\mathcal{M}=\mathcal{M}(\mathbb{X},f):=\mathcal{N}_1(\mathcal{U})$, is the mapper graph of (\mathbb{X},f) .

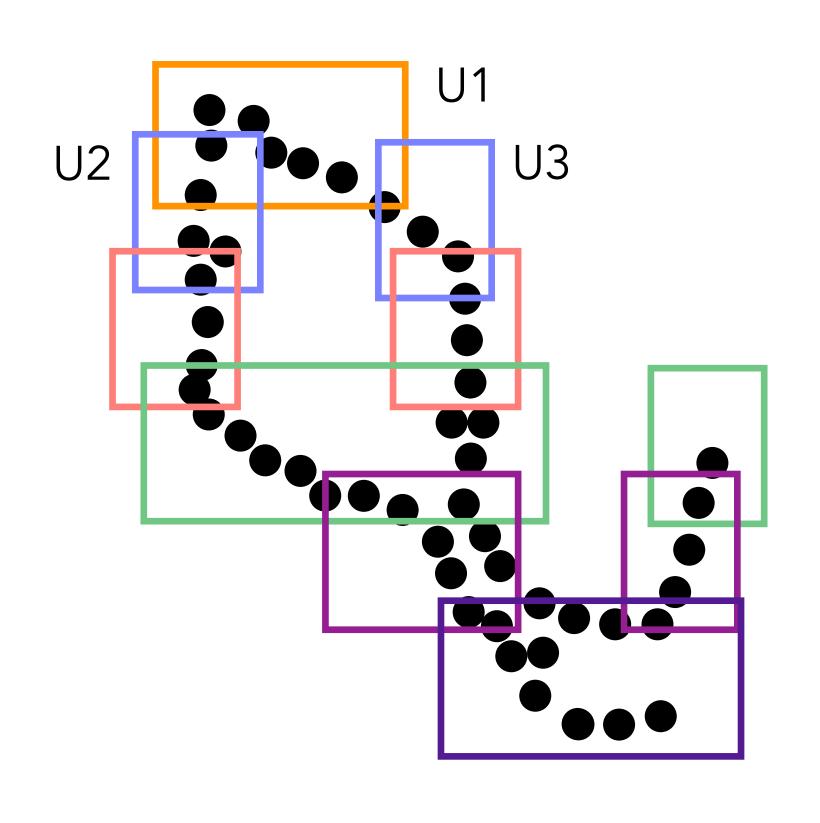
Mapper Graph

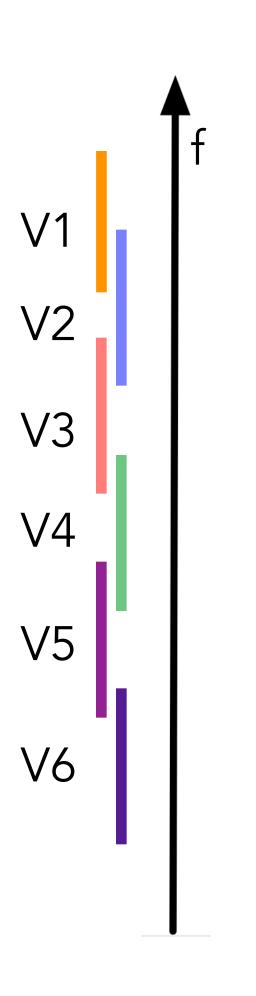


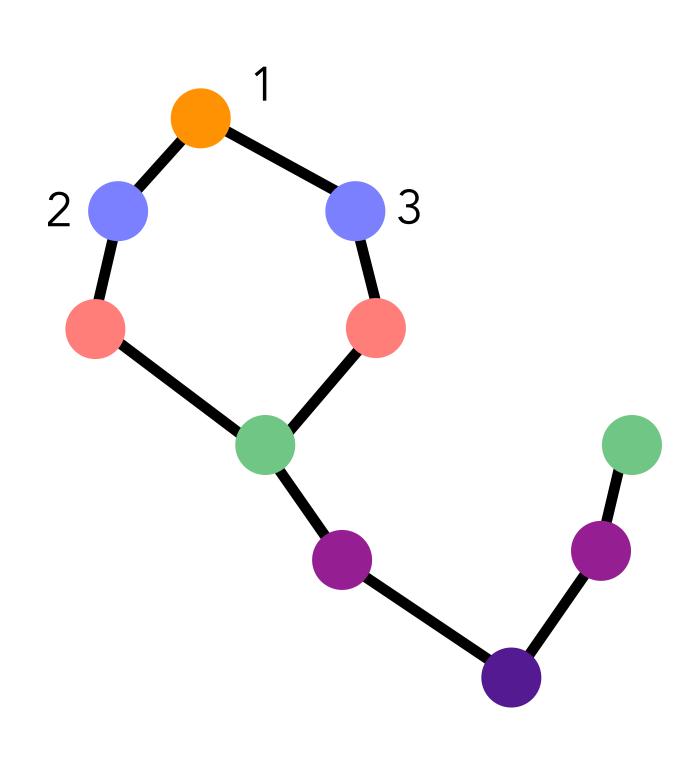
V4

V5

Mapper Graph







Mapper Algorithm: Input, Output, Parameters

Input:

- Point cloud data X
- Distance metric on the point cloud d_{X}
- Filter functions (lens) $f: \mathbb{X} \to \mathbb{R}^d$

Output:

- The mapper graph G_X : a summary of the data as a graph (or a simplicial complex)
- Visualization, statistics and ML

Parameters:

- Parameters for the chosen clustering algorithms
- Filter functions f_1, f_2, \dots , etc.
- Covering parameters
- Visualization parameters: color functions, etc.

Mapper Algorithm: Clustering Details

- Almost any clustering algorithm can be used
- Assume there is a notion of distance (metric) between a pair of points in the data domain (distance can be computed or provided)
- Clustering is equivalent to a notion of connected component in the point cloud setting
- Commonly used clustering algorithms:
 - Density-based spatial clustering of applications with noise (DBSCAN) [2]
 - Single-linkage clustering
 - K-means, etc.
- Desirable properties:
 - Not restricted to Euclidean distance; can take distance matrix input
 - Do not require specifying the number of clusters beforehand

Mapper Algorithm: Parameters for Covering

- Number of intervals: *m*
- Percentage of overlap: p
- Uniform Cover: intervals have equal length, equal percentage of overlap
- Balanced Cover: approximately the same number of points are contained in each cover interval

Mapper Algorithm: Choose Filter functions

- A filter function can be a given prior, e.g. one feature of the point cloud
- It can also be derived from the properties of the point cloud it self
 - Density estimation

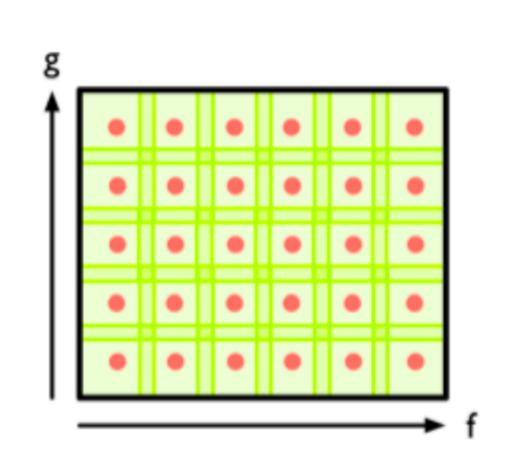
$$f_{\epsilon}(x) = C_{\epsilon} \sum_{y \in X} exp \frac{-d(x, y)^{2}}{\epsilon}$$

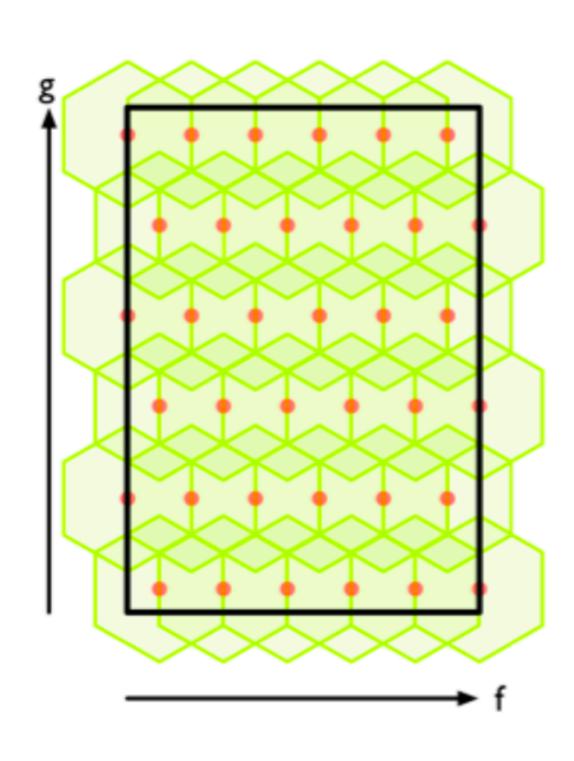
Eccentricity

$$E_p(x) = \left(\frac{\sum_{y \in X} d(x, y)^p}{N}\right)^{1/p}$$

- Distance to a point in the data
- Graph laplacians
- •

Mapper Algorithm: 1D vs 2D Mapper

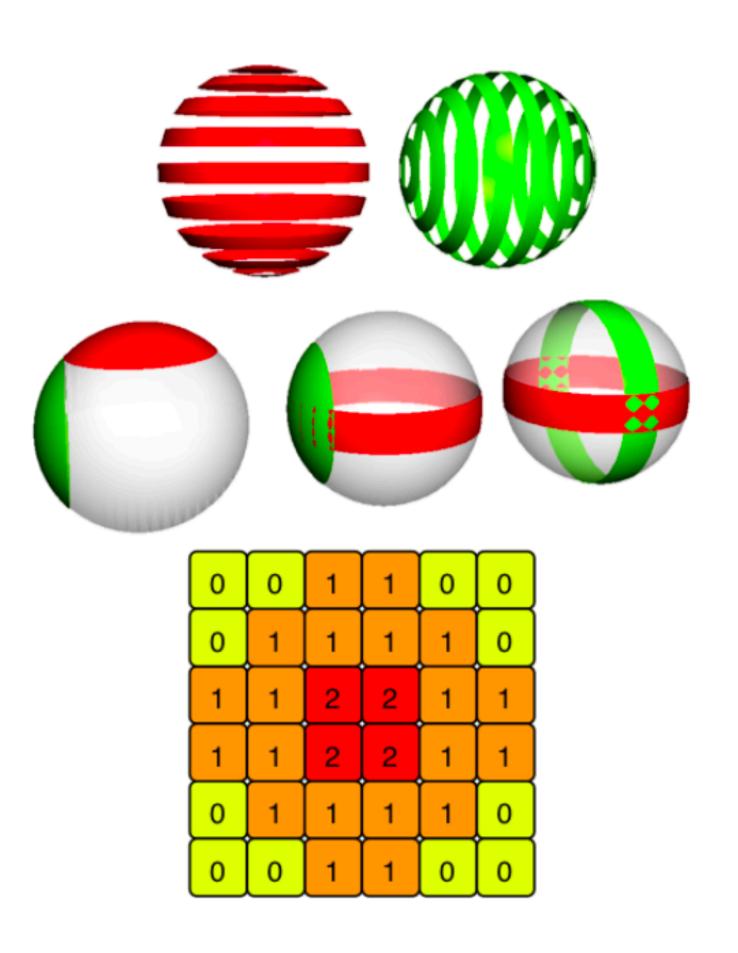




- 1 vs. 2 filter function(s)
- 1D intervals vs. 2D intervals
- For 2D Mapper, the covering of the domain of the function is no longer by intervals, instead, by rectangles or other geometric shapes, etc.

source: [Singh et al. 2007]

Mapper Algorithm: 2D Mapper Example



source: [Singh et al. 2007]

State-of-the-art Tools

- KeplerMapper [3, 4]
- Giotto-TDA [5]
- Mapper Interactive [6]
- Gudhi [7]
- Hyppo-X [8]
- PhenoMapper [9]

•

Kepler Mapper

- A library implementing the Mapper algorithm in Python
- Provides control over the clustering algorithm, scaling algorithms, covering scheme, and nerve scheme
- Scikit-Learn-API-compatible clustering and scaling algorithms
- Interactive capabilities for visual exploration
- Installation
 - Install with pip: pip install kmapper
 - Install from source: git clone https://github.com/MLWave/kepler-mapper cd kepler-mapper pip install -e .

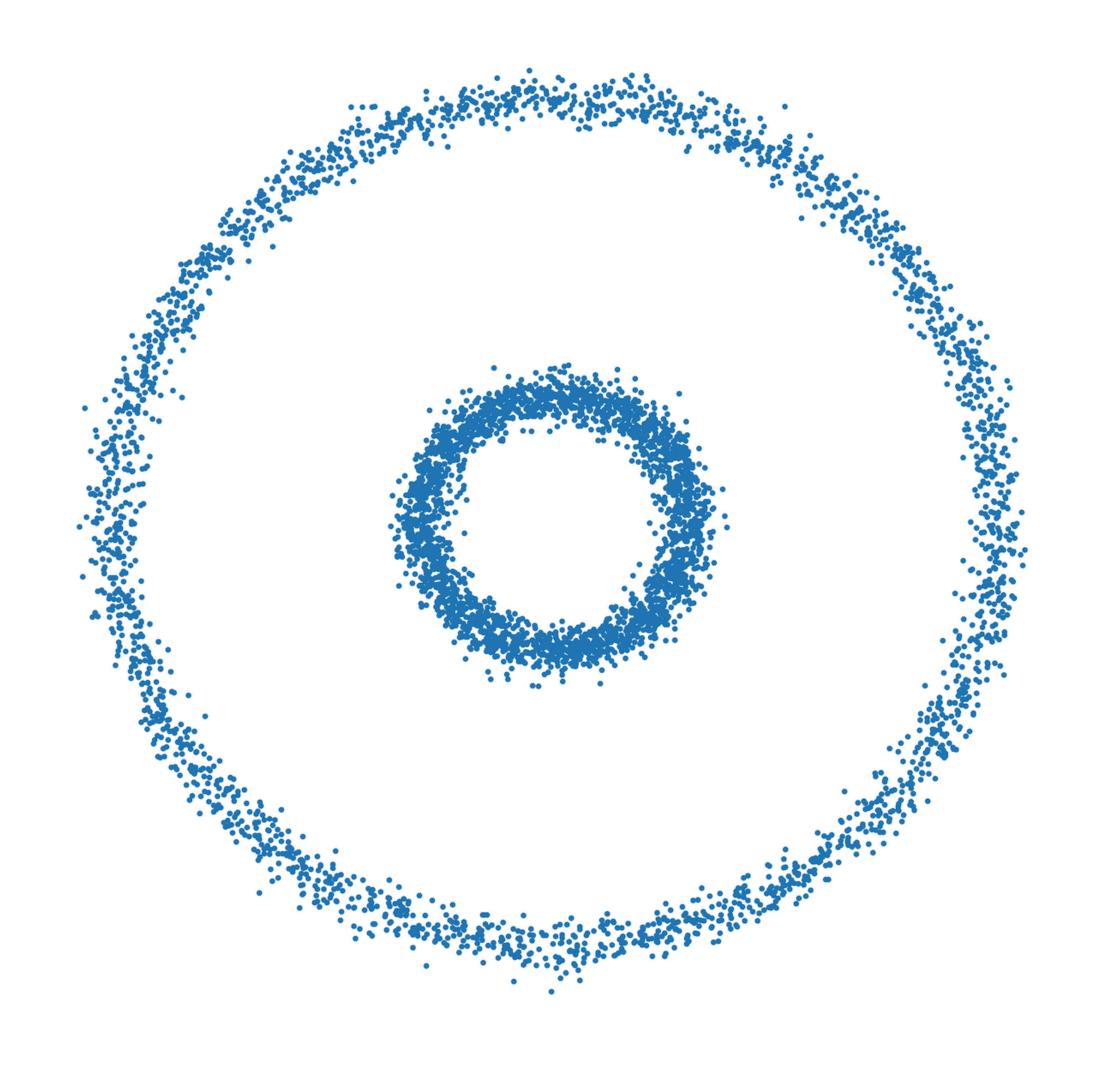
Kepler Mapper: kmapper.KeplerMapper

The class used to build topological networks from (high-dimensional) point cloud data

- Fit a filter function (lens) to a dataset and transform it.
- Map the filter function with overlapping intervals, cluster the points inside the interval, and connect two clusters with an edge if they intersect with each other.
- Visualize the network with HTML and D3.js

Example: two circles

• Data: 2D point cloud



Kepler Mapper: Choose a lens

- fit_transform(X[, projection, scaler, distance_matrix]). The projection parameter can be:
 - a list of dimension indices
 - a string from ["sum", "mean", "median", "max", "min", "std", "dist_mean", "l2norm", "knn_distance_n"]
 - a Scikit-learn class with fit_transform, e.g., manifold.TSNE()
- An n x d array generated manually

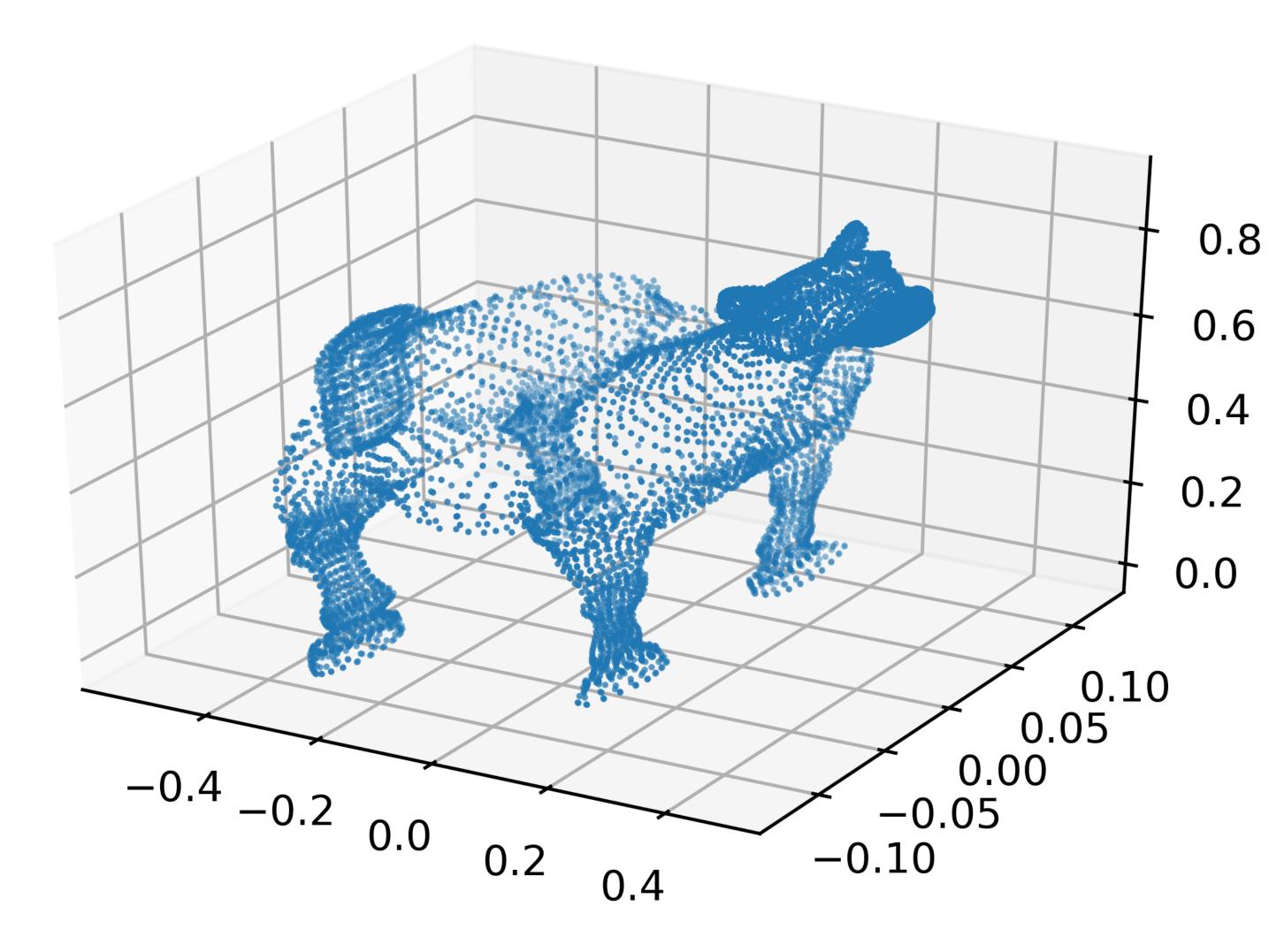
Kepler Mapper: Visualization

- HTML visualization
- Visualization adapters
 - networkx adapter
- Plotlyviz

Example: 3D horse

Data: 3D point cloud

```
0.596860 -0.379364
     -0.013562
     -0.008161 0.602476 -0.378438
     -0.014804 0.590327 -0.383195
     -0.011898 0.582940 -0.390050
     -0.002709 0.575463 -0.396369
      0.027072 \quad 0.222551 \quad -0.542017
8426
8427
      0.027611 \quad 0.220884 \quad -0.545835
8428
      0.033172 \quad 0.221070 \quad -0.543778
8429
      0.032512 0.223522 - 0.540444
8430 -0.054765 0.552121 -0.378869
[8431 rows x 3 columns]
```



Part II

Giotto-TDA

- A high-performance topological machine learning toolbox in Python
- Create a MapperPipeline object that interfaces with scikit-learn for downstream analysis
- Provide an interactive visualization that can be configured in real time
- Installation: pip install -U giotto-tda

Giotto-TDA: MapperPipeline

Basic steps

- Choose a filter function
 A variety of filter functions can be imported as follows:
 from gtda.mapper.filter import FilterFunctionName
- Construct a cover
 A choice of cover can be imported as follow: from gtda.mapper.cover import CoverName
- 3. Choose a clustering algorithm
 - scikit-learn method: from sklearn.cluster import ClusteringAlgorithm
 - giotto-tda method: from gtda.mapper.cluster import FirstSimpleGap
- 4. The resulting mapper graph will be generated automatically by giotto-tda

Giotto-TDA: Visualization

Static

- plot_static_mapper_graph
- Configure the layout, coloring, dimension, etc.

Interactive

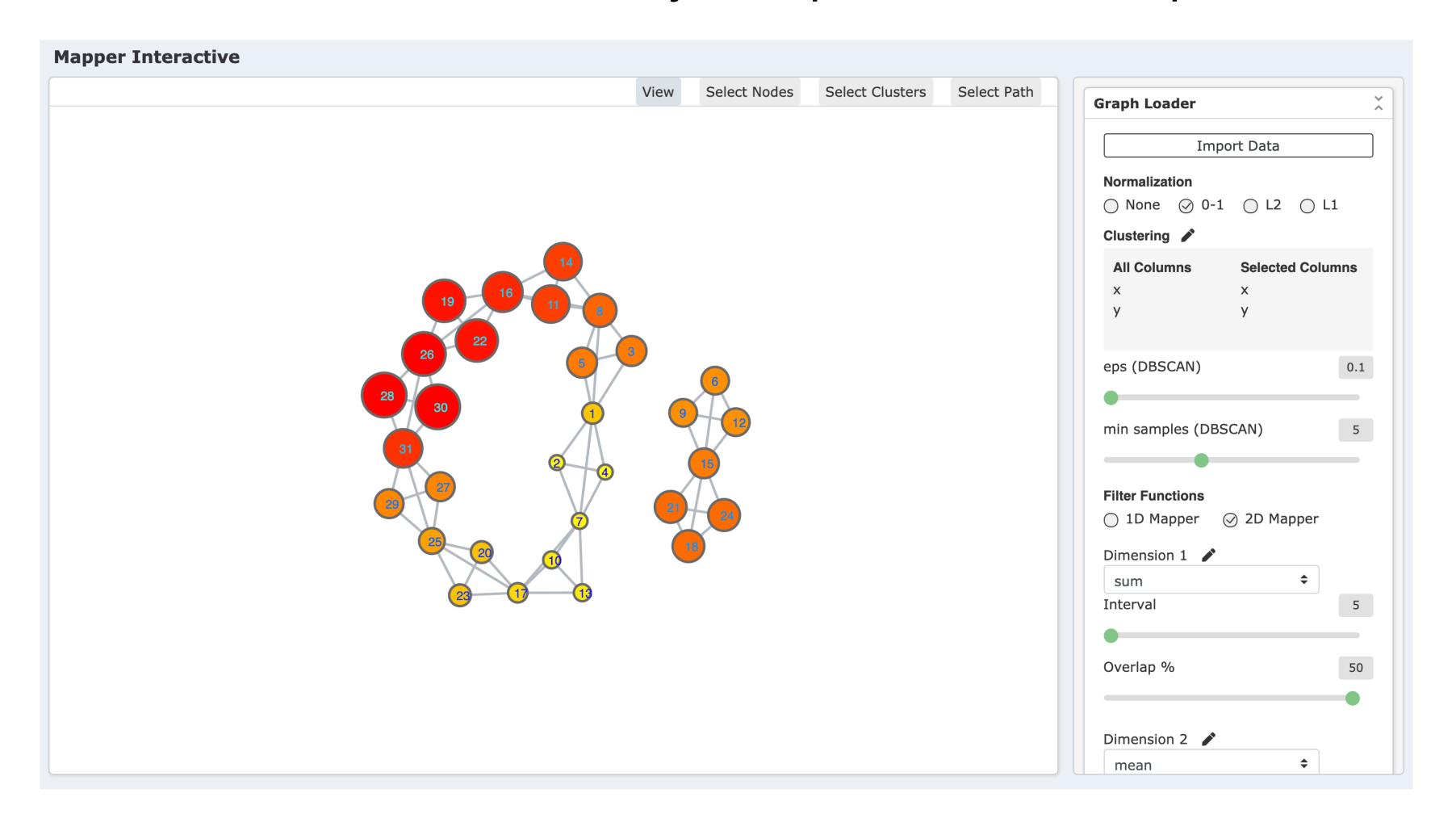
- Compute mapper graph on-the-fly
- Live Jupyter session needed

Mapper Interactive

- A scalable, extendable, and interactive toolbox for the visual exploration of highdimensional point clouds via the mapper graph
- Web-interface for online computation
- Command-line API for offline computation
- The mapper graph computation is modified from the KeplerMapper implementation with an effective speedup strategy
- Installation
 git clone https://github.com/MapperInteractive/MapperInteractive.git
 cd MapperInteractive
 python3 run.py

Mapper Interactive: Interactivity

A web-based interface for on-the-fly computation and exploration.



Mapper Interactive: Extendability

Two modes for different user groups to extend the framework

Novice user mode:

- Describe the new module information within the new modules.json file.
- The addition of supervised and unsupervised learning algorithms is allowed via scikit-learn.

Expert user mode:

 Modify the template function for customizable and multistep analysis pipelines.

Mapper Interactive: Scalability

Key implementational idea

- The computational bottleneck happens during the DBSCAN clustering stage when querying all pairwise distances.
- Modifications:
 - Parallelize individual clustering instances.
 - Precompute the distance matrix of points within each interval.
- This strategy is applicable to any mapper framework employing DBSCAN as a clustering subroutine.

Example: COVID-19 Trends

- Daily records of COVID-19 cases in 9 selected states from April 12, 2020 to September 18, 2020
- 9240 points (rows): a daily record for a given state
- 7 columns for clustering: number of confirmed cases, death cases, active cases, people tested, as well as the testing rate, mortality rate, and incidence rate (i.e., the number of cases per 100K persons).
- Filter function: days of recording
- Parameters
 - DBSCAN: eps = 0.15, min_samples = 5
 - Covering: n = 20, p = 0.5

				1			1	_		I							1			
1	Province_Sta Cou	ntry_Reg	Last_Update La	t	Long_	Confirmed	Deaths	Recovered	Active	FIPS	Incident_Ra	t People_Test Pe	ople_Hosp	Mortality_Ra	UID	ISO3	Testing_Rate H	lospitalizati st	tart_date d	lays
2	Arizona US		6/8/20 3:33	33.7298	-111.4312	26989	1051	5517	20421	4	370.793369	281621	3352	3.89417911	84000004	USA	3869.10221	12.4198748	4/12/20	57
3	California US		6/8/20 3:33	36.1162	-119.6816	130615	4632		125983	6	330.568594	4 2362218		3.5463002	84000006	USA	5978.44874		4/12/20	57
4	Florida US		6/8/20 3:33	27.7663	-81.6868	63938	2700		61238	12	297.694306	1216158	11215	4.22284088	84000012	USA	5662.4122	17.5404298	4/12/20	57
5	Georgia US		6/8/20 3:33	33.0406	-83.6431	51898	2180		49718	13	488.800343	539884	8685	4.20054723	84000013	USA	5084.88736	16.7347489	4/12/20	57
6	Illinois US		6/8/20 3:33	40.3495	-88.9861	127757	5904		121853	17	1008.19764	1042774		4.6212732	84000017	USA	8229.07773		4/12/20	57
7	New Jersey US		6/8/20 3:33	40.2989	-74.521	164164	12176	27824	124164	34	1848.23788	960425	18050	7.41697327	84000034	USA	10812.9301	10.9951025	4/12/20	57
8	New York US		6/8/20 3:33	42.1657	-74.9481	378097	30374	67544	280179	36	1943.5876	2497842	89995	8.03338826	84000036	USA	12840.0245	23.8020931	4/12/20	57
9	North Carolir US		6/8/20 3:33	35.6301	-79.8064	35625	1032	18860	15733	37	339.671193	511226		2.89684211	84000037	USA	4874.35074		4/12/20	57
10	Texas US		6/8/20 3:33	31.0545	-97.5635	75408	1841	49758	23809	48	260.064524	1100446		2.44138553	84000048	USA	3795.18043		4/12/20	57
11	Arizona US		6/7/20 3:53	33.7298	-111.4312	25451	1043	5399	19009	4	349.663272	2 271646	3320	4.0980708	84000004	USA	3732.05882	13.0446741	4/12/20	56
12	California US		6/7/20 3:53	36.1162	-119.6816	128593	4607		123986	6	325.45119	2308300		3.58184349	84000006	USA	5841.98971		4/12/20	56
13	Florida US		6/7/20 3:53	27.7663	-81.6868	62758	2688		60070	12	292.200244	1174185	11163	4.28311928	84000012	USA	5466.98658	17.7873737	4/12/20	56
14	Georgia US		6/7/20 3:53	33.0406	-83.6431	51359	2178		49181	13	483.723781	522857	8662	4.24073677	84000013	USA	4924.51888	16.8655932	4/12/20	56
15	Illinois US		6/7/20 3:53	40.3495	-88.9861	126890	5864		121026	17	1001.35569	1022074		4.62132556	84000017	USA	8065.72315		4/12/20	56
16	New Jersey US		6/7/20 3:53	40.2989	-74.521	163893	12106	27641	124146	34	1845.18683	919448	18023	7.38652658	84000034	USA	10351.5912	10.9968089	4/12/20	56
17	Now York IIC		6/7/20 2.52	12 1657	74 0491	277216	20200	67261	270775	26	1020 E7201	1 2/27/07	90005	0.02510262	94000026	LICA	12520 2616	22 0512607	4/12/20	56

Example - Neuron Activations

- The Cifar dataset is created by passing input images from CIFAR-10 [10] to ResNet-18 neural network.
- Treat the activation vectors collected from the last layer of the network as the input high-dimensional point cloud
- 50K images (rows) from 10 image classes
- 512 dimensions (columns)
- Filter function: I2-norm
- Parameters
 - No normalization
 - DBSCAN: eps = 10, min_samples = 5
 - Covering: n = 40, p = 0.2

References

- [1] G. Singh, F. M´emoli, and G. Carlsson. Topological methods for the analysis of high dimensional data sets and 3D object recognition. Eurographics Symposium on Point-Based Graphics, pages 91–100, 2007.
- [2] M. Ester, H.-P. Kriegel, J. Sander, and X. Xu. A density-based algorithm for discovering clusters in large spatial databases with noise. Proceedings of the 2nd International Conference on Knowledge Discovery and Data Mining, pages 226–231, 1996.
- [3] van Veen et al., (2019). Kepler Mapper: A flexible Python implementation of the Mapper algorithm. Journal of Open Source Software, 4(42), 1315, https://doi.org/10.21105/joss.01315
- [4] Hendrik Jacob van Veen, Nathaniel Saul, David Eargle, & Sam W. Mangham. (2019, October 14). Kepler Mapper: A flexible Python implementation of the Mapper algorithm (Version 1.4.1). Zenodo. http://doi.org/10.5281/ zenodo.4077395
- [5] Tauzin, Guillaume, et al. "giotto-tda:: A Topological Data Analysis Toolkit for Machine Learning and Data Exploration." J. Mach. Learn. Res. 22 (2021): 39-1.
- [6] Youjia Zhou, Nithin Chalapathi, Archit Rathore, Yaodong Zhao, and Bei Wang. 2021. Mapper Interactive: A Scalable, Extendable, and Interactive Toolbox for the Visual Exploration of High-Dimensional Data. In IEEE 14th Pacific Visualization Symposium. 101–110.

References

[7] C. Maria, J.-D. Boissonnat, M. Glisse, and M. Yvinec, "The GUDHI library: simplicial complexes and persistent homology," in International congress on mathematical software. Springer, 2014, pp. 167–174.

[8] Methun Kamruzzaman, Ananth Kalyanaraman, Bala Krishnamoorthy, Stefan Hey, and Pat Schnable. 2019. Hyppo-X: A scalable exploratory framework for analyzing complex phenomics data. *IEEE/ACM Transactions on Computational Biology and Bioinformatics* (2019).

[9] Y. Zhou, M. Kamruzzaman, P. Schnable, B. Krishnamoorthy, A. Kalyanaraman, and B. Wang. Pheno-mapper: an interactive toolbox for the visual exploration of phenomics data. In Proceedings of the 12th ACM Conference on Bioinformatics, Computational Biology, and Health Informatics, pages 1–10, 2021.

[10] Krizhevsky, Alex, and Geoffrey Hinton. "Learning multiple layers of features from tiny images." (2009): 7.