Documentation: Unit Test of Diffusion Maps with Python

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1 Introduction

1.1 Objective and methods

Although TestDiffusionMaps.py is self-explanatory, this supplementary document contains a detailed description of the Diffusion Maps framework. Further, the classes in TestDiffusionMaps.py can be used to solve problems using Diffusion Maps and to perform unit tests in the method fit. If you have interest, see the following GitHub repository https://tinyurl.com/3mxe9mzt, where Jupyter notebooks for two distinct problems and for the unit tests are provided.

The main objective of TestDiffusionMaps.py is to show how the implementation of Diffusion Maps can be simple and powerful. Moreover, two simple examples of unit tests are implemented to verify the code reliability. The classes in TestDiffusionMaps.py were implemented in Python 3.9 using the oriented-object programming (OOP) paradigm, and the examples were run on a computer with macOS. Further, the code requires the following Python toolboxes numpy, scipy, and scikit-learn.

1.2 Theory of Diffusion Maps

Nonlinear dimensionality reduction techniques consider that high-dimensional data can lie on a low-dimensional manifold. To reveal this embedded low-dimensional structure, one can resort to kernel-based techniques such as Diffusion Maps [1]; where the spectral decomposition of the transition matrix of a random walk performed on the data is used to determine a new set of coordinates, also known as diffusion coordinates, embedding this manifold into a space of reduced dimension. For example, data observed in a 3-D space can be constrained to a 2-D structure that can be revealed by the diffusion coordinates. Next, the Diffusion Maps technique is described in details.

Given a dataset $S_{\mathbf{X}} = \{\mathbf{X}_i, \dots, \mathbf{X}_N\}$ with $\mathbf{X}_i \in \mathbb{R}^n$, and a positive semi-definite kernel $k : \mathbb{R}^n \times \mathbb{R}^n \to \mathbb{R}$ such as the Gaussian kernel presented in

Eq. (1), we can construct the kernel matrix $\mathbf{K} = [k_{ij}] = [k(\mathbf{X}_i, \mathbf{X}_j)] \in \mathbb{R}^{N \times N}$ encoding the pairwise similarity of data points in $S_{\mathbf{X}}$. In this regard, the kernel attains its maximum value when $\mathbf{X}_i = \mathbf{X}_j$.

$$k(\mathbf{X}_i, \mathbf{X}_j) = \exp\left(-\frac{||\mathbf{X}_i - \mathbf{X}_j||_2^2}{4\epsilon}\right),\tag{1}$$

where ϵ is the length-scale parameter controlling the level of correlation of a point with its neighbors. Next, to obtain the coordinates embedding the high-dimensional dataset $S_{\mathbf{X}}$ into a low-dimensional space, we first build the following diagonal matrix $\mathbf{D} = [D_{ii}] \in \mathbb{R}^{N \times N}$ as

$$D_{ii} = \sum_{j=1}^{N} k_{ij}.$$
 (2)

Next, a normalized version of the kernel matrix k_{ij} is obtained as

$$\kappa_{ij} = \frac{k_{ij}}{\sqrt{D_{ii}D_{jj}}},\tag{3}$$

and the transition matrix \mathbf{P} is obtained from the following normalization

$$P_{ij} = \frac{\kappa_{ij}}{\sum_{k=1}^{N} \kappa_{ik}}.$$
 (4)

The eigendecomposition of $\mathbf{P} = [P_{ij}]$ yields a set of eigenvectors $\mathbf{\Phi} = [\phi_0, \dots, \phi_N]$ and their respective eigenvalues $\mathbf{\Lambda} = \operatorname{diag}(\lambda_0, \dots, \lambda_N)$. Thus, every element \mathbf{X}_i of $S_{\mathbf{X}}$ has a representation on a low-dimensional space defined by the **diffusion coordinates** $\psi_i = [\lambda_0 \Phi_{i0}, \dots, \lambda_r \Phi_{ir}]^T$, where $r \leq N$. For a more detailed description of the Diffusion Maps framework see [1].

${\small 2}\quad {\small Class}\,\, \textit{DiffusionMaps}$

The Diffusion Maps framework introduced in Section 1 is implemented as a class in TestDiffusionMaps.py. Next, some elements of the class DiffusionMaps are presented. First, the attributes of DiffusionMaps are the following:

- Kernel matrix K (kernel_matrix)
- Transition matrix P (transition_matrix)
- Input data $S_{\mathbf{X}}$ (X)
- Diffusion coordinates ψ_i (diffusion coordinates)

as presented in the following piece of code.

```
class DiffusionMaps:

"""

Diffusion maps is a nonlinear dimensionality reduction technique for embedding high-dimensional data into a low-dimensional Euclidean space revealing their intrinsic geometric structure.

"""

def __init__(self):
    # Attributes of ''DiffusionMaps''.
    self.kernel_matrix = None # Kernel matrix.
    self.transition_matrix = None # Kernel matrix.
    self.X = None # Input data.
    self.diffusion_coordinates = None # Diffusion Coordinates.
```

Listing 1: class DiffusionMaps.

These attributes are instantiated by using the method fit; where X is the input data (2-D numpy array) equivalent to $S_{\mathbf{X}}$, and epsilon is the length-scale parameter ϵ in Eq. 1 (it must be larger than zero).

```
def fit(self, X=None, epsilon=None):
```

Listing 2: Instantiating the attributes of DiffusionMaps

Once the class DiffusionMaps is instantiated, the attributes and methods are accessible from the object using OOP paradigm with Python.

3 Class MyTestCase

The file TestDiffusionMaps.py also contain the class MyTestCase for performing the unit tests in two distinct cases. The first test (Listing 3) is used to compare the number of points in diffusion_coordinates and the number of points in the the input dataset X. Therefore, the diffusion coordinates must be consistent with the input dataset.

Listing 3: Unit test 1.

The second unit test (Listing 4) check the raise of exceptions in the code. In particular, it verifies if raise ValueError is called when the shape of X is not acceptable. In this implementation of Diffusion Maps X must be an array

with two dimensions (row and columns), otherwise the code shows the following error message: Not acceptable shape for 'X'. Therefore, the code will pass this test only if it raises an exception for this particular condition.

```
# Test 2: test if the code correctly raise an exception for the shape
      of the input data 'X'.
     def test_input_shape_exception(self):
2
          # Get a random array with shape (100, 3, 1). DiffusionMaps only
       accepts len(np.shape('X')) = 2.
         X = np.random.rand(100, 3, 1)
         dfm = DiffusionMaps() # Object of 'DiffusionMaps'.
6
          # Get the exception for raise ValueError.
         with self.assertRaises(ValueError) as exception_context:
             dfm.fit(X=X, epsilon=1.0) # Instantiate the attributes of
      'DiffusionMaps' with 'epsilon' = 1.
          # The code will pass the test when it will raise the following
      exception.
         self.assertEqual(str(exception_context.exception), 'Not
      acceptable shape for 'X'.')
```

Listing 4: Unit test 2.

4 Running DiffusionMaps

This section shows how to run one example using DiffusionMaps. The problem consist of unwrapping the Swiss Roll manifold, which is defined in a 3-D space, and it is implicitly included in the unit test presented in Listing 3. Herein, the Swiss Roll manifold is sampled, and a point cloud of points in the 3-D space represent this surface, as observed in Fig. 1 when 2,000 samples are generated using the following scikit-learn command:

```
1 from sklearn.datasets import make_swiss_roll
2 X, color = make_swiss_roll(n_samples=2000, random_state=1)
```

Listing 5: Sampling the Swiss Roll manifold.

One can easily see that the Swiss Roll manifold is defined in 3-D, but it has an intrinsic 2-D structure. Thus, one can use Diffusion Maps to unwrap this 3-D structure. Using the code presented herein, one can obtain the representation of the Swiss Roll manifold in a 2-D space. To this aim, one can use the following commands to instantiate the DiffusionMaps class.

```
1 dfm = DiffusionMaps()
2 dfm.fit(X=X, epsilon=1.0)
```

Listing 6: Instantiating the class DiffusionMaps.

One can observe that the an object of DiffusionMaps (dfm) is created without input arguments. To instantiate the attributes one can use the method fit, which receives the input dataset X and the value of epsilon equal to 1.0 (which is selected by the user).

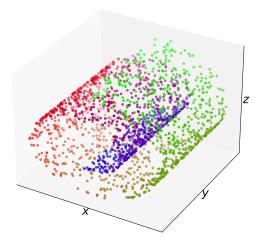


Figure 1: Example 1: Grassmannian diffusion manifold: a) training set for GH, and b) predicted Grassmannian diffusion manifold for 3,000 additional samples.

One of the attributes of DiffusionMaps is diffusion_coordinates (presented as ψ_i in Section 1.2), which stores the diffusion coordinates used to embed the Swiss Roll manifold into a 2-D space. This embedding is presented in Fig. 2, where the diffusion coordinates ψ_2 (diffusion_coordinates[:, 2]), ψ_3 (diffusion_coordinates[:, 3]), ψ_4 (diffusion_coordinates[:, 4]), and ψ_5 (diffusion_coordinates[:, 5]) are plotted with respect to ψ_1 (diffusion_coordinates[:, 1]). It is important mentioning that the 0th diffusion coordinates (ψ_0) are not used in this kind of embedding because they represent the trivial eigendirection. Therefore, when plotting the diffusion coordinates there is no need to show diffusion_coordinates[:, 0]. Based on Fig. 2 one can observe that ψ_1 and ψ_5 are the directions that unwrap the Swiss Roll manifold.

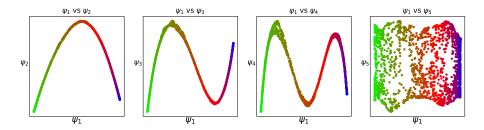


Figure 2: Example 1: Grassmannian diffusion manifold: a) training set for GH, and b) predicted Grassmannian diffusion manifold for 3,000 additional samples.

5 Running MyTestCase

To run the unit tests, one can use the following command in the directory containing TestDiffusionMaps.py:

```
1 $ python -m unittest TestDiffusionMaps
```

Listing 7: Running the unit test.

Therefore, one can expect the following outcome:

Listing 8: Outcome of the unit test.

References

[1] R. R. Coifman and S. Lafon. Diffusion maps. Applied and Computational Harmonic Analysis, 21(1):5-30, 2006. Special Issue: Diffusion Maps and Wavelets.