

# Looking at Non-Linear Dimension Reductions as Models in the Data Space

Jayani P.G. Lakshika

Econometrics & Business Statistics, Monash University  
and

Dianne Cook

Econometrics & Business Statistics, Monash University  
and

Paul Harrison

MGBP, BDInstitute, Monash University  
and

Michael Lydeamore

Econometrics & Business Statistics, Monash University  
and

Thiyanga S. Talagala

Statistics, University of Sri Jayewardenepura

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## Abstract

Nonlinear dimension reduction (NLDR) techniques such as tSNE, and UMAP provide a low-dimensional representation of high-dimensional (high-D) data using non-linear transformation. The methods and parameter choices can create wildly different representations, making it difficult to decide which is best, or whether any or all are accurate or misleading. NLDR often exaggerates random patterns, sometimes due to the samples observed. But NLDR views have an important role in data analysis because, if done well, they provide a concise visual (and conceptual) summary of high-D distributions. To help evaluate the NLDR we have developed an algorithm to show the 2D NLDR model in the high-D space, viewed with a tour. One can see if the model fits everywhere or better in some subspaces, or completely mismatches the data. It is used to evaluate which 2D layout is the best representation of the high-D distribution and see how different methods may have similar summaries or quirks.

*Keywords:* high-dimensional data, dimension reduction, triangulation, hexagonal binning, low-dimensional manifold, manifold learning, tour, data vizualization

# 1 Introduction

Non-linear dimension reduction (NLDR) is popular for making a convenient low-dimensional ( $k - D$ ) representation of high-dimensional ( $p - D$ ) data. Recently developed methods include t-distributed stochastic neighbor embedding (tSNE) (van der Maaten & Hinton 2008), uniform manifold approximation and projection (UMAP) (McInnes & Healy 2018), potential of heat-diffusion for affinity-based trajectory embedding (PHATE) algorithm (Moon et al. 2019), large-scale dimensionality reduction Using triplets (TriMAP) (Amid & Warmuth 2022), and pairwise controlled manifold approximation (PaCMAP) (Wang et al. 2021). However, the representation generated can vary dramatically from method to method, and with different choices of parameters or random seeds made using the same method (Figure 1). The dilemma for the analyst is then, **which representation to use**. The choice might result in different procedures used in the downstream analysis, or different inferential conclusions. The research described here provides new visual tools to aid with this decision.

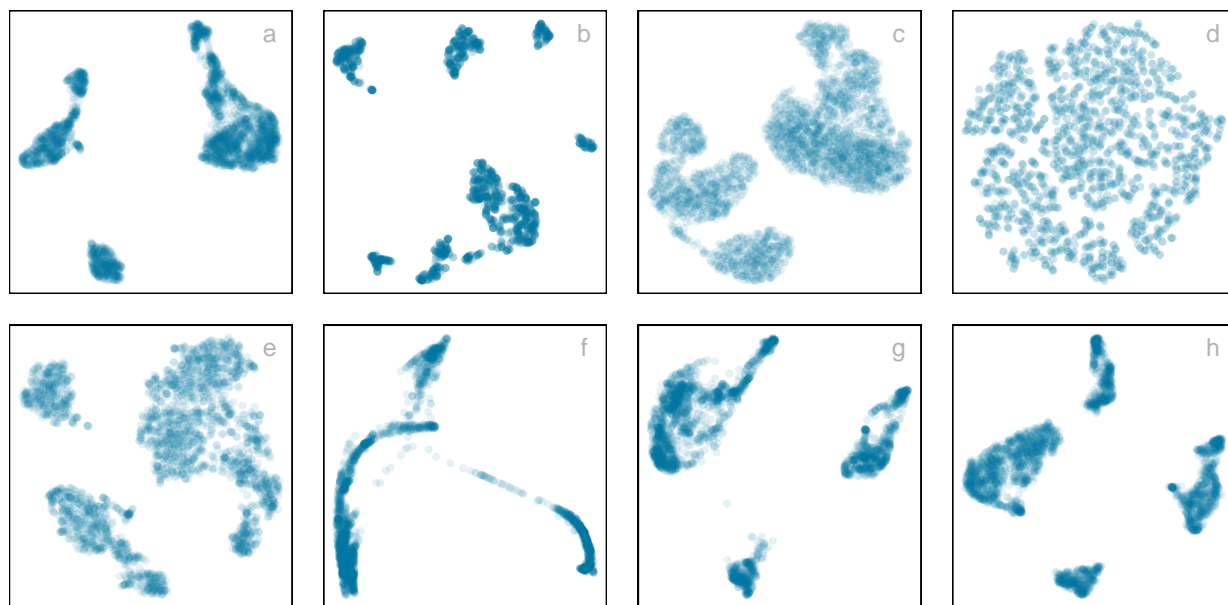


Figure 1: Six different NLDR representations of the same data. Different techniques and different parameter choices are used. Researchers may have seen any of these in their analysis of this data, depending on their choice of method, or typical parameter choice. Would they make different decisions downstream in the analysis depending on which version seen? Which is the most accurate representation of the structure in high dimensions?

The paper is organised as follows. Section 2 provides a summary of the literature on NLDR, and high-dimensional data visualization methods. Section 3 contains the details of the new methodology, including simulated data examples. Two applications illustrating the use of the new methodology for bioinformatics and image classification are in Section 4. Limitations and future directions are provided in Section 5.

## 2 Background

Historically,  $k-D$  representations of  $p-D$  data have been computed using multidimensional Scaling (MDS), which includes principal components analysis (PCA) as a special case. The  $k-D$  representation can be considered to be a layout of points in  $k-D$  produced by an embedding procedure that maps the data from  $p-D$ . In MDS, the  $k-D$  layout is constructed by minimizing a stress function that differences distances between points in  $p-D$  with potential distances between points in  $k-D$ . Various formulations of the stress function result in non-metric scaling (Kruskal 1964) and isomap (Silva & Tenenbaum 2002). Challenges in working with high-dimensional data, including visualization, are outlined in Johnstone & Titterton (2009).

Many new methods for NLDR have emerged in recent years, all designed to better capture specific structures potentially existing in  $p-D$ . Here we focus on five currently popular techniques, tSNE, UMAP, PHATE, TriMAP and PaCMAP. tSNE and UMAP can be considered to produce the  $k-D$  minimizing the divergence between two distributions, where the distributions are modeling the inter-point distances. PHATE, TriMAP and PaCMAP are examples of diffusion processes (Coifman et al. 2005) spreading to capture geometric shapes, that include both global and local structure.

The array of layouts in Figure 1 illustrate what can emerge from the choices of method and parameters, and the random seed that initiates the computation. Key structures interpreted from these views suggest:

- **highly separated clusters** (a, b, e, g, h) with the number ranging from 3-6,
- **stringy branches** (f), and
- **barely separated clusters** (c, d) which apparently **contradicts** the other representations.

It happens because these methods and parameter choices provide different lenses on the interpoint distances in the data.

The alternative approach to visualizing the high-dimensional data is to use linear projections. PCA is the classical approach, resulting in a set of new variables which are linear combinations of the original variables. Tours, defined by Asimov (1985), broaden the scope by providing movies of linear projections, that provide views the data from all directions. Lee et al. (2021) provides an review of the main developments in tours. There are many tour algorithms implemented, with many available in the R package `tourr` (Wickham et al. 2011), and versions enabling better interactivity in `langevitour` (Paul Harrison 2022) and `detourr` (Hart & Wang 2022). Linear projections are a safe way to view high-dimensional data, because they do not warp the space, so they are more faithful representations of the structure. However, linear projections can be cluttered, and global patterns can obscure local structure. The simple activity of projecting data from  $p-D$  suffers from piling (Laa et al. 2022), where data concentrates in the center of projections. NLDR is designed to escape these issues, to exaggerate structure so that it can be observed. But as a result NLDR can hallucinate wildly, to suggest patterns that are not actually present in the data.

The solution is to use the tour to examine how the NLDR is warping the space.

- Model-in-the-data-space: how can we represent the model, eg plane for PCA, grid of values for classification boundaries.

## 3 Method

- Overview: Generate a form that maps the model, that is the interpoint distances. What is the model?
- Notation
- Create a representation of the model
  - using hex-binning in 2D,
  - parameters,
  - tuning,
  - pre-processing
- How does this map to the representation in high-d
  - Centroids,
  - Edges
- Measuring fit
  - Fitted values
  - Error calculation
- What is learned about simulated examples
  - Interesting organisation of points in UMAP
  -

## 4 Applications

### 4.1 pbmc

- NLDR view used to illustrate clusters
- Use our method to assess is it a reasonable representation
- Demonstrate that it is not
- Illustrate how to use our method to get a better representation

### 4.2 digits: 1

- NLDR is used to illustrate different ways 1's are drawn
- Use our method to assess is it a reasonable representation
- Demonstrate that it is, except for the anomalies

## 5 Discussion

- Summarise contributions
- Explain where it is expected or not expected to work, eg higher dimensional relationships
- Diagnostic app to explore differences in distances
- What might be useful enhancements

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