

quollr: An R Package for Visualizing 2D Models from Nonlinear Dimension Reductions in High Dimensional Space

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Abstract An abstract of less than 150 words.

```
library(quollr)
library(tibble)
library(knitr)
library(kableExtra)
library(ggplot2)
```

```
set.seed(20230531)
```

1 Introduction

2 Methodology

3 Implementation

The package can be installed from CRAN:

```
install.packages("quollr")
```

The development version can be installed from GitHub:

```
devtools::install_github("JayaniLakshika/quollr")
```

The following demonstration of the package's functionality assumes quollr has been loaded. We also want to load the built-in data sets `s_curve_noise_training` and `s_curve_noise_umap`.

The main steps for the algorithm can be executed by the main function `fit_highd_model()`, or can be run separately for more flexibility. When constructing the 2D model, the user can choose either to fit the 2D model with hexagonal bin centroids or bin means using `is_bin_centroid` argument.

Constructing the 2D model

Preprocessing

The function `gen_scaled_data()` is used to prepare the NLDR data to fit within the bounds required for regular hexagonal binning, ensuring effective visualization.

Binning data

The main steps for the hexagonal binning algorithm can be executed by the main function `hex_binning()`, or can be run separately for more flexibility.

Step 1: Compute hexagonal grid configurations

Step 2: Generate the hexagonal bin centroids

Step 3: Create the hexagonal grid

Step 4: Assign NLDR data to hexagons

Obtain bin centroids/ means

To obtain hexagonal bin centroids, `extract_hexbin_centroids()` is used. The function `extract_hexbin_mean()` is used to obtain average within each hexagon.

Triangulate bin centroids/ means

The `tri_bin_centroids()` is used to do triangulate bin centroids/ bin means. Then, `gen_edges()` compute the edges to obtain the triangular mesh which is the 2D model.

Lifting the model into high dimensions

The p -D model is generated by `avg_highd_data()` by passing the p -D data.

Tuning model parameters

Mainly there are two model parameter need to consider: benchmark value to remove the low-density hexagons, and benchmark value to remove the long edges. `find_low_dens_hex()` is used to find the hexagons which contains less number of points by considering the density of their neighboring points as well. In here, user can first decide which are the low-density hexagons and pass them to this function to check whether these removal of low-density hexagons can affect the model fit by looking at the the neighbors. On the other hand, `find_lg_benchmark()` function is used to compute the threshold for removing long edges.

Compute residuals and Mean Square Error (MSE)

As a Goodness of fit statistics for the model, `gen_summary()` is used to compute residuals and MSE. For the fitted model, how residuals and MSE is changing is shown in Figure 2.

Visualizations

The package provides four basic visualizations which includes three visualizations regarding 2D model (static vis) and one is regrading p -D model (dynamic vis). Each visualization can be done their own functions which describes in this section.

2D model visualization

To visualize the 2D model, mainly three functions are used. As shown in Figure 1, `geom_trimesh()` to visualize the triangular mesh by adding a new layer to `ggplot()`. After identifying benchmark value to remove long edge, `vis_lg_mesh()` is used to visualize the triangular mesh by coloring the small and long edges. As shown in Figure 1, the small and long edges are colored by black and red respectively. Following this, `vis_rmlg_mesh()` is used to visualize smoothed 2D model which is the 2D model after removing the long edges (see Figure 1). In `vis_lg_mesh()` and `vis_rmlg_mesh()`, `benchmark_value` argument controls the edge removal in 2D.

p -D model visualization

The p -D model overlaid on data is visualized by the function `show_langevi tour()`. This is a useful dynamic visualization result to validate visually whether the model fits the data well or not.

Prediction

`predict_emb()` function is used to predict 2D embedding for a new data point using the fitted model. This function is useful to predict 2D embedding irrespective of the NLDR technique.

Tests

All functions have tests written and implemented using the `testthat` (Wickham 2011) in R.

4 Application

5 Conclusion

6 Acknowledgements

This article is created using [knitr](#) (Xie 2015) and [rmarkdown](#) (Xie, Allaire, and Golemund 2018) in R with the `rjtools::rjournal_article` template. The source code for reproducing this paper can be found at: <https://github.com/JayaniLakshika/paper-quollr>.

References

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