Package 'WOAkMedoids'

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Type Package

| Depends R (>= 3.5.0) |
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| Title Whale Optimization Algorithm for K-Medoids Clustering |
| Version 0.1.0 |
| Date 2024-08-03 |
| Encoding UTF-8 |
| Description Implements the Whale Optimization Algorithm(WOA) for k-medoids clustering, providing tools for effective and efficient cluster analysis in various data sets. The methodology is based on ``The Whale Optimization Algorithm" by Mirjalili and Lewis (2016) <doi:10.1016 j.advengsoft.2016.01.008="">.</doi:10.1016> |
| License GPL (>= 2) |
| Imports dtwclust, proxy, cluster, Rcpp (>= 1.0.11) |
| LinkingTo Rcpp |
| RoxygenNote 7.2.1 |
| LazyData true |
| NeedsCompilation yes |
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| Repository CRAN |
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Lightning7

Lightning7 Data for Testing

Description

A dataset containing example data for testing purposes from the UCR Time Series Classification Archive. This dataset is a time series dataset with correct classifications in the first column. There are 7 classes in this dataset. It contains 73 series, each with 319 time points, and the best DTW window length for this dataset is 5.

Usage

```
data(Lightning7)
```

Format

A data frame with 73 rows and 320 columns. The first column (V1) is a factor vector of correct classifications, and the remaining 319 columns (V2 to V320) are numeric vectors of time series data.

Source

UCR Time Series Classification Archive

References

- Eads, Damian R., et al. "Genetic algorithms and support vector machines for time series classification." Applications and Science of Neural Networks, Fuzzy Systems, and Evolutionary Computation V. Vol. 4787. International Society for Optics and Photonics, 2002.
- http://www.timeseriesclassification.com/description.php?Dataset=Lightning7

Examples

```
data(Lightning7)
head(Lightning7)
```

woa_kmedoids

Whale Optimization Algorithm for K-Medoids Clustering

Description

This function implements the Whale Optimization Algorithm (WOA) for K-Medoids clustering. Supported distance measures are Dynamic Time Warping (DTW) and Euclidean Distance (ED).

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Usage

```
woa_kmedoids(
  data,
  ClusNum,
  distance_method = c("dtw", "ed"),
  learned_w = NULL,
  Max_iter = 20,
  n = 5
)
```

Arguments

data Data matrix

ClusNum Number of clusters

distance_method

Distance calculation method, either "dtw" or "ed"

learned_w Window size for DTW (only used if distance_method is "dtw")

Max_iter Maximum number of iterations (default is 20, it can be adjusted according to the

size of the dataset)

Population size (number of whales, default is 5, it can be adjusted according to

the size of the dataset)

Value

The 'woa_clustering' object containing the clustering result and medoids

Author(s)

Chenan Huang, Narumasa Tsutsumida

References

Mirjalili, S., & Lewis, A. (2016). The whale optimization algorithm. Advances in engineering software, 95, 51-67.

Examples

```
# NOTE: This example only shows how to implement woa_kmedoids using sample data.
# Results do not suggest any meanings.
data(Lightning7)
Lightning7_data <- Lightning7[, -1] # Remove the first column of classification data
  result <- woa_kmedoids(Lightning7_data, ClusNum = 7, distance_method = "dtw", learned_w = 5)
  print(result)</pre>
```

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