

# Package ‘NESS’

April 3, 2025

**Title** What the Package Does (One Line, Title Case)

**Version** 0.0.0.9000

**Description** What the package does (one paragraph).

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**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

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## R topics documented:

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NESS	<i>NESS: Neighbor Embedding Stability Scoring</i>
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## Description

Performs dimensionality reduction (t-SNE, UMAP, or PHATE) multiple times to evaluate local neighbor stability across repeated embeddings. This function is useful for evaluating the robustness of low-dimensional embeddings for high-dimensional data.

## Usage

```
NESS(  
  data,  
  ...,  
  data.name = "",  
  GCP = NULL,  
  cell_type = NULL,  
  rareness = FALSE,  
  method = "tsne",  
  initialization = 1,  
  stability_threshold = 0.75,  
  early_stop = TRUE,  
  seed_base = 1e+10,  
  N = 30,
```

```

k = 50,
svd_cutoff_ratio = 1.1,
svd_max_k = 30,
stop_global_stability_threshold = 0.9,
stop_relative_change_threshold = 0.05
)

```

## Arguments

<code>data</code>	A numeric matrix or data frame with rows as observations and columns as features.
<code>...</code>	Additional arguments passed to the dimensionality reduction methods ( <code>Rtsne</code> , <code>uwot::umap</code> , or <code>phateR::phate</code> ), such as <code>theta</code> for t-SNE, <code>min_dist</code> for UMAP, or <code>decay</code> for PHATE.
<code>data.name</code>	Character string used in plot titles to label the dataset.
<code>GCP</code>	Optional numeric vector of neighborhood sizes (e.g., perplexity for t-SNE or number of neighbors for UMAP/PHATE). If <code>NULL</code> , a default sequence is generated.
<code>cell_type</code>	Optional vector of cell type labels for coloring the embedding plots.
<code>rareness</code>	Logical; if <code>TRUE</code> , computes rareness metrics based on neighbor consistency across embeddings.
<code>method</code>	Dimensionality reduction method to use. One of <code>"tsne"</code> , <code>"umap"</code> , or <code>"phateR"</code> .
<code>initialization</code>	Initialization method: 1 for random, 2 for PCA-based initialization.
<code>stability_threshold</code>	Quantile threshold (default = 0.75) for determining local neighbor stability.
<code>early_stop</code>	Logical; if <code>TRUE</code> , stops early if global stability saturates.
<code>seed_base</code>	Base random seed used for reproducibility.
<code>N</code>	Number of repeated embeddings runs.
<code>k</code>	Number of nearest neighbors to use when computing stability metrics (default=50).
<code>svd_cutoff_ratio</code>	Threshold ratio used to estimate intrinsic dimensionality via SVD (default = 1.1).
<code>svd_max_k</code>	Maximum number of SVD components to check when estimating dimensionality (default = 30).
<code>stop_global_stability_threshold</code>	Early stopping threshold for global stability (default = 0.9).
<code>stop_relative_change_threshold</code>	Early stopping threshold for relative improvement in global stability (default = 0.05).

## Value

A list containing:

- local\_stability** A vector of local kNN stability scores for the selected GCP value.
- plot\_list\_stability** A ggplot2 object showing the embedding colored by stability.
- global\_stability\_plot** A ggplot2 line plot showing global stability across GCP values.
- plot\_list\_cell\_type** (optional) Embedding plot colored by cell type if `cell_type` is provided.
- rareness\_mean** (optional) A ggplot2 plot of rareness score (mean) if `rareness = TRUE`.