# Package 'clixo'

January 29, 2021

Type Package
Title An implementation of the Clique-Extracted Ontology (CliXO) algorithm
Version 0.1.0
<b>Date</b> 2020-11-29
<b>Depends</b> R (>= $4.0.2$ )
<b>Description</b> This package facilitates application of Clique-Extracted Ontology (CliXO) algorithm in R. This algorithm was originally implemented in C++ by Michael Kramer. The input is a feature similarity matrix, while the output is an ontology source-target-similarity-relation data frame.
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LazyData true
<b>Roxygen</b> list(markdown = TRUE)
RoxygenNote 7.1.1
Imports devtools, tidyverse
Suggests BiocStyle, knitr, rmarkdown, kableExtra, magick, testthat
<pre>URL https://github.com/herdiantrisufriyana/clixo</pre>
BugReports https://github.com/herdiantrisufriyana/clixo/issues VignetteBuilder knitr
R topics documented:
clixo
Index

2 clixo

clixo

Create Clique-Extracted Ontology (CliXO)

### **Description**

This function create clique-extracted ontology from a similarity matrix.

## Usage

```
clixo(
    similarity,
    alpha = 0.01,
    beta = 0.5,
    feature_name = "feature",
    onto_prefix = "CliX0"
)
```

### Arguments

similarity Feature similarity, a square matrix of numerics containing feature-feature simi-

larity measures.

alpha A numeric of a noise parameter. Please see https://pubmed.ncbi.nlm.nih.gov/24932003/.

beta A numeric of a parameter which deals with missing edges. Please see https://pubmed.ncbi.nlm.nih.gov

feature\_name A character to annotate feature (source)-ontology (target) relation in the result-

ing ontology.

onto\_prefix A character that precedes the resulting ontology names.

### Value

clique-extracted ontology, a data frame with rows for ontologies and four columns for source, target, similarity, and relation. Feature (source)- ontology (target) relation is annotated as 'feature' as defined by default for feature\_name, while ontology-ontology relation is annotated as 'is\_a'. To differentiate between feature and ontology names, an onto\_prefix with ':' precedes an ontology name. All columns except similarity are characters. Similarity (a numeric) is a minimum threshold by which either features or ontologies (source) belong to an ontology (target).

### **Examples**

```
## Create input example
input=input_example()

## Run CliXO algorithm
ontology=clixo(input$similarity)
```

input\_example 3

input\_example

Make an input example for clixo package

## Description

This function create an input example for clixo package.

## Usage

```
input_example()
```

### Value

output A list of inputs: 1) value, a data frame with rows for instances and columns for features; 2) similarity, a square matrix of numerics containing feature-feature similarity measures.

## Examples

```
## Create input example
input=input_example()
```

# Index

```
* clixo
      clixo, 2
* data
      input_example, 3
* example
      input_example, 3
clixo, 2
input_example, 3
```