

Package ‘x.ent’

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Type Package

Title eXtraction of ENTity

Description Provides a tool for extracting information (entities and relations between them) in text datasets. It also emphasizes the results exploration with graphical displays. It is a rule-based system and works with hand-made dictionaries and local grammars defined by users. 'x.ent' uses parsing with Perl functions and JavaScript to define user preferences through a browser and R to display and support analysis of the results extracted. Local grammars are defined and compiled with the tool Unitex, a tool developed by University Paris Est that supports multiple languages. See ?xconfig for an introduction.

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Depends R (>= 3.0.0)

Imports stringr, xtable, jsonlite, ggplot2, statmod

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URL <https://github.com/win-stub/x.ent>

BugReports <https://github.com/win-stub/x.ent/issues>

SystemRequirements Perl (>= 5.0), Unitex (>= 3.0
<http://www-igm.univ-mlv.fr/~unitex/>)

NeedsCompilation no

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add_unique*Add a value to a current list that every value is unique***Description**

Add a value to an existing list of values. These values are unique in the list.

Usage

```
add_unique(list, value)
```

Arguments

list	: a list of values
value	: a value that we want to add to the list

Value

list	return a list that elements in the list aren't duplicated
------	---

Examples

```
list1= c("a", "b", "c")
value = "a"
list1 <- add_unique(list1,value)
```

str_count*Count words in a text*

Description

Count words of characters in the string which satisfy a regular expression

Usage

```
str_count(x, pattern, sep)
```

Arguments

x	input string
pattern	regular expression
sep	a string used to separate columns, default is "".

Value

number	return a number of words that satisfies a regular expression
--------	--

Examples

```
x = "file_1:b::$carbonate:c:dimethylsulfide:coccoliths:co2:aragonite:calcite:"  
str_count(x, pattern=":co2:", sep="")
```

trim*Remove whitespace from both sides of a string*

Description

Remove all spaces from text except for single spaces between words

Usage

```
trim(x)
```

Arguments

x	is a string that we want to delete whitespace from both sides
---	---

Examples

```
str = " Hello World! "  
trim(str)
```

upload_dico	<i>Upload file</i>
-------------	--------------------

Description

Copy file from a local folder to a folder on the system

Usage

```
upload_dico(file)
```

Arguments

file : the path of local file

xdata	<i>Transform the results to data frame</i>
-------	--

Description

This is a function using transformation of results to data frame.

Usage

```
xdata(e = NULL)
xdata_value(v, sort = "a")
```

Arguments

e a vector of a entity or a list of entities, if it is nul, it shows all entities and relations that it is configured in the configuration file

v a entity

sort with the function xdata_value, variable "sort" allows you to sort values following frequency or alphabetically

Details

The data frame contains the columns of the name of entity or relationship and the rows of values of named entity.

Value

data frame return a data frame

Author(s)

Tien T. Phan

See Also

[xparses](#) call the main function of module extraction written by Perl

Examples

```
xdata() #show all entities
xdata(c("p","b")) #show two entities: "p", "b"
xdata_value("p") #show only values of entity "p"
#there are two columns "value" et "freq" in this data frame
xdata_value("p")[[["value"]]] #convert to a vector
```

xentity

List of entities or relations

Description

Show all entities or relations

Usage

```
xentity()
xrelation()
```

Value

list return a list of entities or relations

See Also

[xshow](#) display results

Examples

```
xentity()
xrelation()
```

xhist*Graph xhist***Description**

The function **xhist** in **x.ent** is a graphical representation of the distribution of entities with time.

Usage

```
xhist(v = "")
```

Arguments

v	a value of entity or the relations between entities
---	---

Details

Result after calling the function **xparse** has the following format:

1. file_name:entity:\$:list_value_found
2. ...
3. file_name:entity1:entity2:....:\$:value_e1:value_e2:....:negation

Function **xhist** will convert the data format above to a data frame. The histogram uses this data frame to display graphs. The data frame format:

1. column file : name of file
2. column date : (format mm.yyyy)
3. column value_date, this value is used for creating histogram
4. column visible: if visible = 1 then this record will be used in histogram

Value

This function returns a data frame so that users can check or use it to create new graphs.

dataframe	return a data frame
------------------	---------------------

See Also

[xplot](#) type graphique plot
[xshow](#) display the results of extracted data
[xshow](#) display results

Examples

```
xhist() #all documents
xhist(v="colza") #only documents contain "colza"
xhist(v="colza:altise") #only documents contain a relation "colza:altise"
```

xparse*Call script Perl for extracting data from corpus*

Description

Call script Perl for extracting data from corpus. Before you run, you must configure a configuration file ini.json in the folder config as: dictionaries, graphs of grammar (Use tools Unitex for creating)...

Usage

```
xparse(json_path = "", verbose=FALSE)
```

Arguments

json_path	path of configuration file (*.json)
verbose	logical. Should R report extra information on progress? Set to TRUE by the command-line option –verbose.

Details

Input: dictionaries, grammars (build with software Unitex). Output: a result file of every entity and relation

Value

Result file includes:

comp1	data of every entity such as: file1:entity1:\$:data1:data2:
comp2	data of every relation of every entity for example: file1:entity1:entity2:\$\$:data1:data2:1

See Also

[xshow](#) display results

Examples

```
xparse()
```

xplot*Graph xplot***Description**

Graph xplot, this graph compares the appearance of entities or relations during one period

Usage

```
xplot(v1 = "", v2 = "", t = "")
```

Arguments

v1	O or 1 entity1 value
v2	a vector of entity2 value
t	a time value, format (mm.yyyy) or interval of time value, for example: t=c("02.2010","02.2012")

Details

Result after calling the function `xparse` has the following format:

1. file_name:entity:\$:list_value_found
2. ...
3. file_name:entity1:entity2:...:\$:value_e1:value_e2:....:negation

Function `xplot` will convert the data format above to a data frame. The `xplot` uses this data frame to display graphs. The data frame format:

1. column file : name of file
2. column date : (format mm.yyyy)
3. column value_date, this value is used for creating graph
4. column visible: if visible = 1 then this record will be used in graph
5. column value of entite v1 or v2 or v1 combined with v2

Value

This function returns a data frame so that users can check or use it to create new graphs.

dataframe	return a data frame
-----------	---------------------

See Also

[xhist](#) type graphique histogram
[xprop](#) type graphique proportion
[xshow](#) displays results of extracted data

Examples

```
xplot(v1="colza")
xplot(v1="colza",v2=c("altice","rouille"))
xplot(v1="colza",v2=c("altice","rouille"),t="09.2010")
xplot(v1="colza",v2=c("altice","rouille"),t=c("09.2010","02.2011"))
```

xprop

Graph xprop

Description

This visualization is a type of 100% stacked histogramme. The graph xprop shows the distribution of the relationship between entities in the corpus. The total of the bar represents 100%.

Usage

```
xprop(v1,v2,type=1)
```

Arguments

v1	a vector of values
v2	a vector of values
type	type of graph

Details

After calling the function xparses, the result has the following format:

1. file_name:entity:\$:list_value_found
2. ...
3. file_name:entity1:entity2:...:\$:value_e1:value_e2:....:negation

Function xprop will convert the data format above to a data frame such as:

1. a list of columns that call the values of v2. Those columns will contain a value 0 or 1.
2. a column has a name "cat" - categorie.
3. a column has a name "val" - value.

Each line describes the relevant information between values of vector v1 and values of vector v2. If there exists a relationship between a value of v1 with a value of v2 then the column of value v2 will be 1, the column "cat" carrying value is the value of v2 and the column "val" has the value current of v1.

Author(s)

Tien T. Phan

See Also

[xhist](#) type graphique histogram
[xplot](#) type graphique plot

Examples

```
xprop(v1=c("chou", "colza"), v2=c("mouche du chou", "rouille"))
v1 = as.vector(xdata_value("p")[[["value"]]])
v2 = as.vector(xdata_value("b")[[["value"]]])
xprop(v1, v2, type=2)
```

xshow

*Show results***Description**

Show results after calling the function `xparse`.

Usage

```
xshow(e=NULL, sort="a")
```

Arguments

- | | |
|------|---|
| e | an entity or a list of entities that you want display, default e = NULL => display all columns |
| sort | type sort of data, default sort = "a" => sorted by alphabet, sort = "f" => sorted by frequency. |

Details

Show results after calling function `xparse`. The result file has format:

1. entity file1:entity1:\$:data1:data2:data3:
2. relation file1:entity1:entity2:\$\$:data_e1:data_e2:negation

Author(s)

Tien T. Phan

See Also

[xparse](#) call the main function of module extraction written by Perl
[xshow](#) display results

Examples

```
xfile() #show all names of files in corpus  
xshow() #all columns  
xshow(e="p",sort="a") #show result of entity "p", sorted by alphabet  
xshow(e="p",sort="f")  
xshow(e=c("p", "m"))
```

xtest *Test each pair relations*

Description

We recommend four testings distribution to compare two samples:

1. Kolmogorov Smirnov test
2. Wilcoxon signed rank test
3. Student's t test
4. Compare Groups of Growth Curves

Usage

```
xtest(v1, v2)
```

Arguments

- | | |
|----|-------------------------------|
| v1 | a vector of the first entity |
| v2 | a vector of the second entity |

Details

The function xtest will combine the values in the first entity with the values in the second entity, each pair relations will be looking in documents. If this relationship exists, it will bring a value 1 otherwise 0

Author(s)

Tien T. Phan

See Also

- [ks.test](#) Kolmogorov Smirnov test
[wilcox.test](#) Wilcoxon signed rank test
[t.test](#) Student's t test
[compareGrowthCurves](#) Compare Groups of Growth Curves

Examples

```
#get all values of entity bioaggressor
b <- as.vector(xdata_value("b")[[ "value"]])
xtest("colza",b)
```

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