Package 'calACS'

March 31, 2016

Type Package

Title Calculations for All Common Subsequences

Version 2.2.2
Date 2016-3-31
Author Alan Gu
Maintainer Alan Gu <alan.on.ca@gmail.com></alan.on.ca@gmail.com>
Description Implements several string comparison algorithms, including calACS (count all common subsequences), lenACS (calculate the lengths of all common subsequence). Some algorithms differentiate between the more strict definition of subsequence, where a common subsequence cannot be separated by any other items, from its looser counterpart, where a common subsequence can be interrupted by other items. This difference is shown in the suffix of the algorithm (-Strict vs -Loose). For example, q-w is a common subsequence of q-w-er and q-e-w-r on the looser definition, but not on the more strict definition. calACSLoose Algorithm from Wang, H. All common subsequences (2007) IJCAI International Joint Conference on Artificial Intelligence, pp. 635-640.
License GPL
LazyData TRUE
RoxygenNote 5.0.1
NeedsCompilation no
Repository CRAN
Date/Publication 2016-03-31 19:35:54
R topics documented:
calACSLoose 2 calACSStrict 2 lenACSStrict 3 lenLCSStrict 4 longestVec 5
Index 6
1

2 calACSStrict

calACSLoose	Calculate the total number of all common subsequences between a string and a vector/list of strings. Subsequences can be interrupted by
	items, i.e. q-w is considered a subsequence of q-e-w-r

Description

Calculate the total number of all common subsequences between a string and a vector/list of strings. Subsequences can be interrupted by items, i.e. q-w is considered a subsequence of q-e-w-r

Usage

```
calACSLoose(vecA, listB, sep = "-", dropFirstItem = FALSE)
```

Arguments

vecA The single string

1istB The vector/list of 1 or more strings

sep Delimiter separating each items in a sequence

dropFirstItem Boolean. If true, the first item in each sequence is excluded from counting all

subsequences

Value

The total number of all common subsequences as an integer in a vector

Examples

```
 {\tt calACSLoose("q-w-e-r", c("q-e-w-r", "q-r-e-w"), "-") } \\ {\tt calACSLoose("itemToBeDropped-q-w-e-r", "itemToBeDroped-q-e-w-r", "-", dropFirstItem=TRUE) } \\ {\tt calACSLoose("itemToBeDropped-q-w-e-r", "-", dropFirstItem=TRUE) } \\ {\tt calACSLoose("ite
```

calACSStrict Count the total number of all common subsequences between a string

and a vector/list of strings. Subsequences cannot be interrupted by any item, i.e. q-w is not considered a subsequence of q-e-w-r due to

the interrupting 'e'

Description

Count the total number of all common subsequences between a string and a vector/list of strings. Subsequences cannot be interrupted by any item, i.e. q-w is not considered a subsequence of q-e-w-r due to the interrupting 'e'

lenACSStrict 3

Usage

```
calACSStrict(vecA, listB, sep = "-", dropFirstItem = FALSE,
  ignoreLenOneSubseq = FALSE, ignoreLenZeroSubseq = FALSE)
```

Arguments

vecA The single string

listB The vector/list of 1 or more strings

sep Delimiter separating each items in a sequence

dropFirstItem Boolean. If true, the first item in each sequence is excluded from counting all

subsequences

ignoreLenOneSubseq

Boolean. If true, all length one subequences are not counted as common subse-

quences

ignoreLenZeroSubseq

Boolean. If true, the length zero subsequence (empty set) is not counted as a

common subsequence

Value

The total number of all common subsequences as an integer in a vector

Examples

```
calACSStrict("q-w-e-r", c("q-e-w-r","q-r-e-w"), "-")
calACSStrict("itemToBeDropped-q-w-e-r", "itemToBeDropped-q-e-w-r", "-", dropFirstItem=TRUE)
```

lenACSStrict

Calculate the length of each common subsequences between a string and a vector/list of strings. Subsequences cannot be interrupted by any item, i.e. q-w is not considered a subsequence of q-e-w-r due to the interrupting 'e'

Description

Calculate the length of each common subsequences between a string and a vector/list of strings. Subsequences cannot be interrupted by any item, i.e. q-w is not considered a subsequence of q-e-w-r due to the interrupting 'e'

Usage

```
lenACSStrict(vecA, listB, sep = "-", dropFirstItem = FALSE,
  ignoreLenOneSubseq = FALSE)
```

4 lenLCSStrict

Arguments

vecA The single string

1istB The vector/list of 1 or more strings

sep Delimiter separating each items in a sequence

dropFirstItem Boolean. If true, the first item in each sequence is excluded from counting all

subsequences

ignoreLenOneSubseq

Boolean. If true, all length one subequences are not counted as common subse-

quences

Value

A list of vectors of the length of each common subsequence

Examples

```
lenACSStrict("q-w-e-r", c("q-e-w-r","q-r-e-w","q-w-r-e"), "-")
lenACSStrict("itemToBeDropped-q-w-e-r", "itemToBeDropped-q-e-w-r", "-", dropFirstItem=TRUE)
```

lenLCSStrict Calculate the length of the longest common subsequence (KCS) be-

tween a string and a vector/list of strings. Subsequences cannot be interrupted by any item, i.e. q-w is not considered a subsequence of

q-e-w-r due to the interrupting 'e'

Description

Calculate the length of the longest common subsequence (KCS) between a string and a vector/list of strings. Subsequences cannot be interrupted by any item, i.e. q-w is not considered a subsequence of q-e-w-r due to the interrupting 'e'

Usage

```
lenLCSStrict(vecA, listB, sep = "-", dropFirstItem = FALSE)
```

Arguments

vecA The single string

1istB The vector/list of 1 or more strings

sep Delimiter separating each items in a sequence

dropFirstItem Boolean. If true, the first item in each sequence is excluded from counting all

subsequences

longestVec 5

Value

A list of vectors of the length of each common subsequence

Examples

```
lenACSStrict("q-w-e-r", c("q-e-w-r","q-r-e-w","q-w-r-e"), "-")
lenACSStrict("itemToBeDropped-q-w-e-r", "itemToBeDropped-q-e-w-r", "-", dropFirstItem=TRUE)
```

longestVec

The function takes in multiple vectors of any length, and returns the one with the longest length. The tieBreaker variable controls if the first or the last of the longest vectors gets returned in case there are multiple

Description

The function takes in multiple vectors of any length, and returns the one with the longest length. The tieBreaker variable controls if the first or the last of the longest vectors gets returned in case there are multiple

Usage

```
longestVec(..., tieBreaker = "last")
```

Arguments

... vectors of any length

tieBreaker decides if the first or the last longest vector gets returned if there are multiple

longest vectors. Can be either 'first' or 'last'. Default to 'last'.

Examples

```
longestVec(1:5, c('a', 'b'))
```

Index

```
calACSLoose, 2
calACSStrict, 2
lenACSStrict, 3
lenLCSStrict, 4
longestVec, 5
```