

# Package: calACS (via r-universe)

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**Type** Package

**Title** Calculations for All Common Subsequences

**Version** 2.2.2

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**Description** Implements several string comparison algorithms, including calACS (count all common subsequences), lenACS (calculate the lengths of all common subsequences), and lenLCS (calculate the length of the longest 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-e-r 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

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**Repository** <https://agu-1.r-universe.dev>

**RemoteUrl** <https://github.com/cran/calACS>

**RemoteRef** HEAD

**RemoteSha** c152cf80f7c20c141e6ec5b88a22e4ca8ceb252b

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calACSLoose	<i>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</i>
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### 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
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

### Value

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

### Examples

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

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calACSstrict	<i>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'</i>
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### 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'

### 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 subsequences are not counted as common subsequences
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)
```

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lenACSStrict	<i>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'</i>
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### 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)
```

### 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 subsequences are not counted as common subsequences

### 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)
```

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lenLCSStrict	<i>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'</i>
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**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
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

**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)
```

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longestVec	<i>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</i>
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**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'))
```

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