

# Package: topomatch (via r-universe)

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**Type** Package

**Title** Match toponyms by similarity

**Version** 0.4

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**Description** Wraps local-global alignment algorithm borrowed from package Biostrings Works better than global alignment and requires less fine-tuning (although is considerably slower too) <https://ro-che.info/articles/2016-12-11-local-alignment>. Allows to inspect the matching and act accordingly.

**Imports** Biostrings

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.2.3

**Suggests** testthat (>= 3.0.0)

**Config/testthat/edition** 3

**Repository** <https://cirad-astre.r-universe.dev>

**RemoteUrl** <https://forgemia.inra.fr/umr-astre/topomatch>

**RemoteRef** HEAD

**RemoteSha** f61ff809aede27c545121836bf34e536511d595

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best_matches	<i>Best matches</i>
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**Description**

Best matches

**Usage**

```
best_matches(x)
```

**Arguments**

x                    topomatch object

**Value**

The original list of toponyms with their best-ranked candidates.

**Examples**

```
best_matches(topomatch(c("a", "b"), c("A", "X")))
```

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exact_matches	<i>Exact toponym matches</i>
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**Description**

Exact toponym matches

**Usage**

```
exact_matches(x)
```

**Arguments**

x                    topomatch object

**Value**

named logical vector

**Examples**

```
exact_matches(topomatch(c("a", "b"), "A"))
```

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similar_matches	<i>Similar toponym matches</i>
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**Description**

Similar toponym matches

**Usage**

```
similar_matches(x)
```

**Arguments**

x                    topomatch object

**Value**

named logical vector

**Examples**

```
similar_matches(topomatch(c("a", "b"), "A"))
```

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topomatch	<i>Toponymy matching</i>
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**Description**

Match a list of toponyms against a list of candidates.

**Usage**

```
topomatch(x, candidates)
```

**Arguments**

x                    character vector of terms to match, or data.frames with the same number of columns corresponding to different nested levels of administrative units (e.g. region, department, municipality).

candidates          character vector or data.frame of terms to compare against.

**Details**

For comparison, strings are trimmed off white spaces, converted to upper-case and special characters transliterated to ASCII.

When comparing places defined at multiple hierarchical levels (using multiple columns), the level-wise scores are multiplied.

**Value**

A matrix with `x` in rows and candidates in columns with matching scores.

**Examples**

```
## Match vectors of toponyms
x <- c("abá", "Efé", "IJI", "kkk")
cand <- c("aba", "ef", "iji", "opo", "Uvu")
topomatch(x, cand)

## Match data.frames of toponyms defined at multiple levels
x <- data.frame(
  region = c("A", "A", "B", "B"),
  province = c("abá", "Efé", "IJI", "kkk")
)
cand <- data.frame(
  reg = c("a", "à", "b", "c", "b"),
  prov = c("aba", "ef", "iji", "opo", "Uvu")
)
topomatch(x, cand)
```

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transcribe

*Transcribe toponyms according to a topomatch table*

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**Description**

Build a function for transcription.

**Usage**

```
transcribe(obj, fixes = NULL)
```

**Arguments**

`obj` A topomatch object.  
`fixes` A Named character vector of manual fixes.

**Details**

If there are any unmatched toponyms, they must be manually matched in `fixes` for transcription. On the other hand, `fixes` can provide matches for incorrect associations. See examples.

**Value**

A function that takes a character vector among the original terms and returns a character vector with the corresponding matches.

**Examples**

```
(perfect <- topomatch("AFGHANISTAN", "Afghanistan"))
transcribe(perfect)("Afghanistan")

terms <- c("foo", "bar", "foobar", "X")
(mat <- topomatch(terms, c("Foo", "Bar")))
setNames(transcribe(mat, fixes = c(X = "Bar"))(terms), terms)

## You can still transcribe terms that do not need fixes
# transcribe(mat)(terms) # Error: X is not fixed
transcribe(mat)(c("foo", "bar")) # Works even if X remains unfixed.
```

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trim

*Trim whitespaces*

---

**Description**

Remove the whitespace before or after a string of character

**Usage**

```
trim(x, internal = TRUE)
```

**Arguments**

x                    character vector, matrix or array.  
internal            logical. Whether to replace sequential internal by a single space

**Value**

A character object of the same type.

**Author(s)**

Robert J. Hijmans and Jacob van Etten. The code was borrowed from function [trim](#) in package raster.

**Examples**

```
trim(c("  hi folks  !  ", "  hello  world  "))
```

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`unmatched`*Unmatched toponyms*

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**Description**

Vector of toponyms with more than one plausible candidate with the same score.

**Usage**

```
unmatched(x)
```

**Arguments**

x                    topomatch object

**Value**

named logical vector

**Examples**

```
unmatched(topomatch(c("a", "b"), c("A", "X")))
```

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