

Package: ahocorasick (via r-universe)

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Title Fast Multi-Pattern String Matching with the 'Aho-Corasick' Algorithm

Version 0.2.0

Description Provide fast multi-pattern string matching for 'R' using the 'Aho-Corasick' algorithm, powered by the 'Rust' 'aho-corasick' crate. It builds reusable automaton for detecting matches, counting matches, locating character, extracting matched text, and replacing matches in character vectors. For more details on the 'Aho-Corasick' algorithm, please see Aho and Corasick (1975) <[doi:10.1145/360825.360855](https://doi.org/10.1145/360825.360855)>.

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URL <https://yousa-mirage.github.io/r-ahocorasick/>,
<https://github.com/Yousa-Mirage/r-ahocorasick>

BugReports <https://github.com/Yousa-Mirage/r-ahocorasick/issues>

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ac_build	<i>Build an Aho-Corasick automaton</i>
----------	--

Description

ac_build() compiles a character vector of patterns into a reusable automaton backed by the Rust aho-corasick crate.

Usage

```
ac_build(
  patterns,
  match_kind = c("standard", "leftmost_first", "leftmost_longest"),
  implementation = c("auto", "noncontiguous_nfa", "contiguous_nfa", "dfa"),
  ascii_case_insensitive = FALSE,
  duplicate = c("keep", "error", "deduplicate")
)
```

Arguments

patterns	A character vector of non-empty patterns.
match_kind	Matching semantics: <ul style="list-style-type: none"> • "standard" supports overlapping search (Default). • "leftmost_first" returns leftmost non-overlapping matches, breaking ties by pattern order. • "leftmost_longest" returns leftmost non-overlapping matches, breaking ties by longest match.
implementation	Rust automaton implementation. "auto" lets the crate choose.
ascii_case_insensitive	Use ASCII-only case-insensitive matching. Default is FALSE.
duplicate	How duplicate patterns are handled: <ul style="list-style-type: none"> • "keep" preserves duplicates in their original order. • "error" fails if patterns contains duplicates. • "deduplicate" keeps the first occurrence of each pattern and drops later duplicates.

Value

An immutable <ac_automaton> object.

See Also

[ac_locate\(\)](#), [ac_locate_df\(\)](#), [ac_detect\(\)](#), [ac_count\(\)](#), [ac_extract\(\)](#), [ac_extract_df\(\)](#), [ac_replace\(\)](#), [ac_patterns\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
length(ac)
ac_info(ac)
```

ac_count

Count pattern matches in documents

Description

ac_count() returns the number of pattern matches in each document.

Usage

```
ac_count(ac, doc, overlapping = FALSE, na = c("keep", "zero", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build().
doc	A character vector of documents to search.
overlapping	Default is FALSE. If TRUE, count overlapping matches. This is only supported when ac was built with match_kind = "standard".
na	How to handle NA documents. "keep" returns NA_integer_ (default); "zero" treats missing documents as zero matches; "error" fails.

Value

An integer vector with the same length as doc.

See Also

[ac_count_file\(\)](#), [ac_detect\(\)](#), [ac_locate\(\)](#), [ac_extract\(\)](#).

Examples

```
if (requireNamespace("dplyr", quietly = TRUE)) {
  ac <- ac_build(c("hello", "world"))
  docs <- data.frame(doc = c("hello world", "nothing", "world"))
  dplyr::mutate(docs, n_matches = ac_count(ac, doc))
}
```

ac_count_file	<i>Count pattern matches in files</i>
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Description

ac_count_file() returns the number of pattern matches in each file.

Usage

```
ac_count_file(ac, path, stream = FALSE, overlapping = FALSE)
```

Arguments

ac	An <ac_automaton> object created by ac_build().
path	A vector of file paths to search.
stream	If FALSE (default), each file is read into memory before searching. If TRUE, files are searched as streams. Stream search requires an automaton built with match_kind = "standard".
overlapping	Default is FALSE. If TRUE, count overlapping matches. This is only supported when stream = FALSE and ac was built with match_kind = "standard".

Value

An integer vector with the same length as path.

See Also

[ac_count\(\)](#), [ac_detect_file\(\)](#), [ac_locate_bytes\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
path <- tempfile()
writeLines("hello hello world", path)
ac_count_file(ac, path)
```

ac_detect

Detect pattern matches in documents

Description

`ac_detect()` returns whether each document has at least one pattern match.

Usage

```
ac_detect(ac, doc, na = c("keep", "false", "error"))
```

Arguments

ac	An <code><ac_automaton></code> object created by <code>ac_build()</code> .
doc	A character vector of documents to search.
na	How to handle NA documents. "keep" returns NA (default); "false" treats missing documents as not matched; "error" fails.

Value

A logical vector with the same length as doc.

See Also

[ac_detect_file\(\)](#), [ac_count\(\)](#), [ac_locate\(\)](#), [ac_extract\(\)](#).

Examples

```
if (requireNamespace("dplyr", quietly = TRUE)) {
  ac <- ac_build(c("hello", "world"))
  docs <- data.frame(doc = c("hello world", "nothing", "world"))
  dplyr::mutate(docs, matched = ac_detect(ac, doc))
}
```

ac_detect_file *Detect pattern matches in files*

Description

ac_detect_file() returns whether each file has at least one pattern match.

Usage

```
ac_detect_file(ac, path, stream = FALSE)
```

Arguments

ac	An <ac_automaton> object created by ac_build().
path	A vector of file paths to search.
stream	If FALSE (default), each file is read into memory before searching. If TRUE, files are searched as streams. Stream search requires an automaton built with match_kind = "standard".

Value

A logical vector with the same length as path.

See Also

[ac_detect\(\)](#), [ac_count_file\(\)](#), [ac_locate_bytes\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
path <- tempfile()
writeLines("hello world", path)
ac_detect_file(ac, path)
```

ac_extract *Extract pattern matches from documents*

Description

ac_extract() returns one list element per document. Each element contains the matched text and the corresponding pattern values.

Usage

```
ac_extract(ac, doc, overlapping = FALSE, na = c("keep", "empty", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build().
doc	A character vector of documents to search.
overlapping	Default is FALSE. If TRUE, extract overlapping matches. This is only supported when ac was built with match_kind = "standard".
na	How to handle NA documents. "keep" returns one row with missing matches and patterns values (default); "empty" treats missing documents as no matches; "error" fails.

Value

A list with the same length as doc. Each element is a data frame with one row per match and two columns:

- matches: Text matched in the document.
- patterns: Pattern values corresponding to each match.

See Also

[ac_extract_df\(\)](#), [ac_locate\(\)](#), [ac_detect\(\)](#), [ac_count\(\)](#).

Examples

```
if (
  requireNamespace("dplyr", quietly = TRUE) &&
  requireNamespace("tibble", quietly = TRUE) &&
  requireNamespace("tidyr", quietly = TRUE)
) {
  ac <- ac_build(c("hello", "world"))
  tibble::tibble(doc = c("hello world", "nothing", "world")) |>
    dplyr::mutate(extracted = ac_extract(ac, doc)) |>
    tidyr::unnest(extracted)
}
```

ac_extract_df

Extract pattern matches as a data frame

Description

ac_extract_df() is the data-frame form of [ac_extract\(\)](#). It is useful when you want one row per match instead of one list element per document.

Usage

```
ac_extract_df(ac, doc, overlapping = FALSE, na = c("omit", "keep", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build().
doc	A character vector of documents to search.
overlapping	Default is FALSE. If TRUE, extract overlapping matches. This is only supported when ac was built with match_kind = "standard".
na	How to handle NA documents. "omit" drops missing documents (default); "keep" returns one row with missing result columns for each missing document; "error" fails.

Value

A data frame with one row per match and three columns: doc_id, matches, and patterns.

See Also

[ac_extract\(\)](#), [ac_locate_df\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
doc <- c("hello world", "nothing", "world hello")
ac_extract_df(ac, doc)
```

ac_extract_file	<i>Extract pattern matches from files</i>
-----------------	---

Description

ac_extract_file() returns one list element per file. Each element contains the matched text and the corresponding pattern values.

Usage

```
ac_extract_file(ac, path, stream = FALSE, overlapping = FALSE)
```

Arguments

ac	An <ac_automaton> object created by ac_build().
path	A vector of file paths to search.
stream	If FALSE (default), each file is read into memory before searching. If TRUE, files are searched as streams. Stream search requires an automaton built with match_kind = "standard".
overlapping	Default is FALSE. If TRUE, extract overlapping matches. This is only supported when stream = FALSE and ac was built with match_kind = "standard".

Value

A list with the same length as path. Each element is a data frame with one row per match and two columns:

- matches: Text matched in the file.
- patterns: Pattern values corresponding to each match.

See Also

[ac_extract\(\)](#), [ac_detect_file\(\)](#), [ac_count_file\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
path <- tempfile()
writeLines("hello world", path)
ac_extract_file(ac, path)
```

ac_info	<i>Return automaton metadata</i>
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Description

Return automaton metadata

Usage

```
ac_info(ac)
```

Arguments

ac An <ac_automaton> object created by `ac_build()`.

Value

A list of automaton metadata.

See Also

[ac_build\(\)](#), [ac_patterns\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
ac_info(ac)
```

ac_locate *Locate pattern matches in strings*

Description

ac_locate() searches a character vector with a compiled automaton and returns one list element per document. Character offsets are 1-based and inclusive, so they can be used directly with substr().

Usage

```
ac_locate(ac, doc, overlapping = FALSE, na = c("keep", "empty", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build().
doc	A character vector of documents to search.
overlapping	Default is FALSE. If TRUE, report overlapping matches. This is only supported when ac was built with match_kind = "standard".
na	How to handle NA documents. "keep" returns one row with missing pattern_id, start, and end values (default); "empty" treats missing documents as no matches; "error" fails.

Value

A list with the same length as doc. Each element is a data frame with one row per match and three columns:

- pattern_id: Index of the matched pattern in ac_patterns(ac).
- start: 1-based index of the first character in each match.
- end: 1-based index of the last character in each match.

See Also

[ac_locate_df\(\)](#), [ac_locate_bytes\(\)](#), [ac_extract\(\)](#), [ac_detect\(\)](#), [ac_count\(\)](#).

Examples

```
if (
  requireNamespace("dplyr", quietly = TRUE) &&
  requireNamespace("tibble", quietly = TRUE) &&
  requireNamespace("tidyr", quietly = TRUE)
) {
  ac <- ac_build(c("hello", "world"))
  tibble::tibble(doc = c("hello world", "nothing", "world")) |>
    dplyr::mutate(hits = ac_locate(ac, doc)) |>
    tidyr::unnest(hits)
}
```

ac_locate_bytes	<i>Locate pattern matches with byte offsets</i>
-----------------	---

Description

ac_locate_bytes() searches a character vector with a compiled automaton and returns byte offsets from the Rust aho-corasick crate. Byte offsets are 0-based, and byte_end is end-exclusive.

Usage

```
ac_locate_bytes(ac, doc, overlapping = FALSE, na = c("omit", "keep", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build().
doc	A character vector of documents to search.
overlapping	Default is FALSE. If TRUE, report overlapping matches. This is only supported when ac was built with match_kind = "standard".
na	How to handle NA documents. "omit" drops missing documents (default); "keep" returns one row with missing result columns for each missing document; "error" fails.

Value

A data frame with one row per match and four columns: doc_id, pattern_id, byte_start, and byte_end.

See Also

[ac_locate\(\)](#), [ac_locate_df\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
doc <- c("hello world", "nothing", "world hello")
ac_locate_bytes(ac, doc)
```

ac_locate_df	<i>Locate pattern matches as a data frame</i>
--------------	---

Description

ac_locate_df() is the data-frame form of [ac_locate\(\)](#). It is useful when you want one row per match instead of one list element per document.

Usage

```
ac_locate_df(ac, doc, overlapping = FALSE, na = c("omit", "keep", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build() .
doc	A character vector of documents to search.
overlapping	Default is FALSE. If TRUE, report overlapping matches. This is only supported when ac was built with <code>match_kind = "standard"</code> .
na	How to handle NA documents. "omit" drops missing documents (default); "keep" returns one row with missing result columns for each missing document; "error" fails.

Value

A data frame with one row per match and four columns: doc_id, pattern_id, start, and end.

See Also

[ac_locate\(\)](#), [ac_locate_bytes\(\)](#), [ac_extract_df\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
doc <- c("hello world", "nothing", "world hello")
ac_locate_df(ac, doc)
```

ac_locate_file	<i>Locate pattern matches in files</i>
----------------	--

Description

ac_locate_file() searches files with a compiled automaton and returns one list element per file. Character offsets are 1-based and inclusive, so they can be used directly with substr().

Usage

```
ac_locate_file(ac, path, overlapping = FALSE)
```

Arguments

ac	An <ac_automaton> object created by ac_build().
path	A vector of file paths to search.
overlapping	Default is FALSE. If TRUE, report overlapping matches. This is only supported when ac was built with match_kind = "standard".

Details

File location search is always non-streaming. Converting byte offsets from a streaming search into R-facing character offsets would require a second pass over the same file to reconstruct UTF-8 character boundaries. Keeping ac_locate_file() as a simple in-memory search is the clearest implementation.

Value

A list with the same length as path. Each element is a data frame with one row per match and three columns:

- pattern_id: Index of the matched pattern in ac_patterns(ac).
- start: 1-based index of the first character in each match.
- end: 1-based index of the last character in each match.

See Also

[ac_locate\(\)](#), [ac_detect_file\(\)](#), [ac_count_file\(\)](#), [ac_extract_file\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
path <- tempfile()
writeLines("hello world", path)
ac_locate_file(ac, path)
```

ac_patterns *Return patterns stored in an automaton*

Description

Return patterns stored in an automaton

Usage

```
ac_patterns(ac)
```

Arguments

ac An <ac_automaton> object created by `ac_build()`.

Value

A character vector of stored patterns.

See Also

[ac_build\(\)](#), [ac_info\(\)](#).

Examples

```
ac <- ac_build(c("hello", "world"))
ac_patterns(ac)
```

ac_replace *Replace pattern matches in documents*

Description

`ac_replace()` replaces all non-overlapping matches in each document with the corresponding replacement string.

Usage

```
ac_replace(ac, doc, replace_with, na = c("keep", "empty", "error"))
```

Arguments

ac	An <ac_automaton> object created by ac_build().
doc	A character vector of documents to search and replace.
replace_with	A character vector of replacements. If length 1, the same replacement is used for every pattern. Otherwise, it MUST have the same length as ac_patterns(ac), and replacements are matched to patterns by position.
na	How to handle NA documents. "keep" returns NA_character_ (default); "empty" treats missing documents as empty strings; "error" fails.

Value

A character vector with the same length and names as doc.

See Also

[ac_build\(\)](#), [ac_detect\(\)](#), [ac_count\(\)](#), [ac_extract\(\)](#), [ac_locate\(\)](#).

Examples

```
ac <- ac_build(c("fox", "brown", "quick"))
ac_replace(
  ac,
  "The quick brown fox.",
  c("sloth", "grey", "slow")
)

ac <- ac_build(c("append", "appendage", "app"), match_kind = "leftmost_first")
ac_replace(ac, "append the app to the appendage", c("x", "y", "z"))
```

ac_replace_file	<i>Replace pattern matches in files</i>
-----------------	---

Description

ac_replace_file() replaces all non-overlapping matches in input files and writes the result to output files.

Usage

```
ac_replace_file(ac, path, replace_with, output = NULL, stream = FALSE)
```

Arguments

ac	An <ac_automaton> object created by <code>ac_build()</code> .
path	A vector of input file paths to search and replace.
replace_with	A character vector of replacements. If length 1, the same replacement is used for every pattern. Otherwise, it MUST have the same length as <code>ac_patterns(ac)</code> , and replacements are matched to patterns by position.
output	A vector of output file paths. It must have the same length as <code>path</code> . If <code>NULL</code> , output paths are created by adding <code>"_replaced"</code> suffix. Existing output files are overwritten.
stream	If <code>FALSE</code> (default), each file is read into memory before replacement. If <code>TRUE</code> , files are searched and replaced as streams. Stream replacement requires an automaton built with <code>match_kind = "standard"</code> .

Value

A character vector of output file paths with the same length as `path`.

See Also

[ac_replace\(\)](#), [ac_detect_file\(\)](#), [ac_count_file\(\)](#).

Examples

```
ac <- ac_build(c("fox", "brown", "quick"))
path <- tempfile(fileext = ".txt")
writeLines("The quick brown fox.", path)
ac_replace_file(path = path, ac = ac, replace_with = c("sloth", "grey", "slow"))
```

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