Package 'unnest'

December 4, 2023

Title Unnest Hierarchical Data Structures

Version 0.0.5

Description Fast flattening of hierarchical data structures (e.g. JSON, XML) into data.frames with a flexible spec language.

License GPL (>= 2)

Encoding UTF-8

RoxygenNote 7.1.1

Suggests data.table, dplyr, knitr, repurrrsive, rmarkdown, roxygen2, testthat, tibble, tidyr

VignetteBuilder knitr

URL https://github.com/vspinu/unnest

BugReports https://github.com/vspinu/unnest/issues

NeedsCompilation yes

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Repository CRAN

Date/Publication 2023-12-03 23:10:02 UTC

R topics documented:

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spec   *Unnest spec*

Description

Unnest spec is a nested list with the same structure as the nested json. It specifies how the deeply nested lists ought to be unnested. `spec()` is a handy constructor for spec lists. `s()` is a shorthand alias for `spec()`.

Usage

```
spec(
  selector = NULL,
  ...,
  as = NULL,
  children = NULL,
  groups = NULL,
  include = NULL,
  exclude = NULL,
  stack = NULL,
  process = NULL,
  default = NULL
)

s(
  selector = NULL,
  ...,
  as = NULL,
  children = NULL,
  groups = NULL,
  include = NULL,
  exclude = NULL,
  stack = NULL,
  process = NULL,
  default = NULL
)
```

Arguments

selector   A shorthand syntax for an `include` parameter. Can be a list or a character vector.

1. When `selector` is a list or a character vector with length greater than 1, each element is an include parameter at the corresponding level. For example `s(c("a", "b"), ...)` is equivalent to `s(include = "a", s(include = "b", ...))`

2. When `selector` is a character of length 1 and contains "/" characters it is split with "/" first. For instance `s(c("a", "b"), ...)`, `s("a/b", ...)`

and `s("a", s("b", ...))` are all equivalent to the canonical `s(include = "a", s(include = "b", ...))`. Components consisting entirely of digits are converted to integers. For example `s("a/2/b" ...)` is equivalent to `s("a", s(2, s("b", ...)))`

3. Multiple `include` fields can be separated with `,`. For example `s("a/b,c/d")` is equivalent to `s("a", s(include = c("b", "c"), s("d", ...)))`

`as`	name for this field in the extracted data.frame
`children, ...`	Unnamed list of children spec. `...` is merged into `children`. `children` is part of the canonical spec.
`groups`	Named list of specs to be processed in parallel. The return value is a named list of unnested data.frames. The results is the same as when each spec is `unnested` separately except that dedupe parameter of `unnest()` will work across groups and execution is faster because the nested list is traversed once regardless of the number of groups.
`include, exclude`	
	A list, a numeric vector or a character vector specifying components to include or exclude. A list can combine numeric indexes and character elements to extract.
`stack`	Whether to stack this node (TRUE) or to spread it (FALSE). When `stack` is a string an index column is created with that name.
`process`	Extra processing step for this element. Either NULL for no processing (the default), "as_is" to return the entire element in a list column, "paste" to paste elements together into a character column.
`default`	Default value to insert if the `include` specification hasn't matched.

Value

`s()`: a canonical spec - a list consumed by C++ unnesting routines.

Examples

```
s("a")
s("a//c2")
s("a/2/c2,cid")
```

---

unnest	*Unnest lists*

Description

Unnest nested lists into a flat data.frames.

Usage

```
unnest(
  x,
  spec = NULL,
  dedupe = FALSE,
  stack_atomic = NULL,
  process_atomic = NULL,
  process_unnamed_lists = NULL,
  cross_join = TRUE
)
```

Arguments

x	a nested list to unnest
spec	spec to use for unnesting. See [spec()]().
dedupe	whether to dedupe repeated elements. If TRUE, if a node is visited for a second time and is not explicitly declared in the spec the node is skipped. This is particularly useful with grouped specs.
stack_atomic	Whether atomic leaf vectors should be stacked or not. If NULL, the default, data.frame vectors are stacked, all others are spread.
process_atomic	Process spec for atomic leaf vectors. Either NULL for no processing (the default), "as_is" to return the entire element in a list column, "paste" to paste elements together into a character column.
process_unnamed_lists	
	How to process unnamed lists. Can be one of "as_is" - return a list column, "exclude" - drop these elements unless they are explicitly included in the spec, "paste" - return a character column, "stack" - automatically stack. If NULL (the default), do nothing - process them normally according to the specs.
cross_join	Specifies how the results from sibling nodes are joined (cbinded) together. The shorter data.frames (fewer rows) can be either recycled to the max number of rows across all joined components with cross_join = FALSE. Or, the results are cross joined (produce all combinations of rows across all components) with cross_join = TRUE. cross_join = TRUE is the default because of no data loss and it is more conducive for earlier error detection with incorrect specs

Value

A data.frame, data.table or a tibble as specified by the option unnest.return.type. Defaults to data.frame.

Examples

```
x <- list(a = list(b = list(x = 1, y = 1:2, z = 10),
                   c = list(x = 2, y = 100:102)))
xxx <- list(x, x, x)
```

```
## spreading
unnest(x, s("a"))
unnest(x, s("a"), stack_atomic = TRUE)
unnest(x, s("a/b"), stack_atomic = TRUE)
unnest(x, s("a/c"), stack_atomic = TRUE)
unnest(x, s("a"), stack_atomic = TRUE, cross_join = TRUE)
unnest(x, s("a//x"))
unnest(x, s("a//x,z"))
unnest(x, s("a/2/x,y"))

## stacking
unnest(x, s("a/", stack = TRUE))
unnest(x, s("a/", stack = TRUE, as = "A"))
unnest(x, s("a/", stack = TRUE, as = "A"), stack_atomic = TRUE)
unnest(x, s("a/", stack = "id"), stack_atomic = TRUE)
unnest(x, s("a/", stack = "id", as = ""), stack_atomic = TRUE)

unnest(xxx, s(stack = "id"))
unnest(xxx, s(stack = "id"), stack_atomic = TRUE)
unnest(xxx, s(stack = "id", s("a/b/y/", stack = TRUE)))

## exclusion
unnest(x, s("a/b/", exclude = "x"))

## dedupe
unnest(x, s("a", s("b/y"), s("b")), stack_atomic = TRUE)
unnest(x, s("a", s("b/y"), s("b")), dedupe = TRUE, stack_atomic = TRUE)

## grouping
unnest(xxx, stack_atomic = TRUE,
       s(stack = TRUE,
         groups = list(first = s("a/b/x,y"),
                       second = s("a/b"))))

unnest(xxx, stack_atomic = TRUE, dedupe = TRUE,
       s(stack = TRUE,
         groups = list(first = s("a/b/x,y"),
                       second = s("a/b"))))

## processing as_is
str(unnest(xxx, s(stack = "id",
                  s("a/b/y", process = "as_is"),
                  s("a/c", process = "as_is"))))
str(unnest(xxx, s(stack = "id", s("a/b/", process = "as_is"))))
str(unnest(xxx, s(stack = "id", s("a/b", process = "as_is"))))

## processing paste
str(unnest(x, s("a/b/y", process = "paste")))
str(unnest(xxx, s(stack = TRUE, s("a/b/", process = "paste"))))
str(unnest(xxx, s(stack = TRUE, s("a/b", process = "paste"))))

## default
unnest(x, s("a/b/c/", s("b", default = 100)))
```

```
unnest(x, s("a/b/c/", stack = "ix", s("b", default = 100)))
```

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