{rrapply}: Revisiting R-base rapply()

The minimal rrapply-package contains a single function rrapply(), providing an extended implementation of R-base's rapply() function. rrapply() recursively applies a function f to elements of a nested list and controls how to structure the returned result.

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Function signature

```
rrapply(
  object,
  condition,
  f,
  classes = "ANY",
  deflt = NULL,
  how = c("replace", "list", "unlist",
        "prune", "flatten", "melt", "bind",
        "recurse", "unmelt", "names"),
  options,
  ...
)
```

```
• object a "list-like" object;
```

```
    condition a condition for application of f;
    f a function to recursively apply to each list element;
    classes classes to which f is applied, can include "list" or "data.frame";
    deflt a default return value;
    how how to structure the result;
    options
```

Example data

• ...

```
library(rrapply)
# data: renewable energy per country 2016
# as % of total energy consumption
data("renewable_energy_by_country")
# data: pokemon properties in pokemon G0
data("pokedex")
```

condition functions;

additional arguments for f and

How to structure the result

• how = "replace"

replaces elements x satisfying condition and classes by f(x) and maintains list structure:

```
# replace all missing values by 0
rrapply(
  renewable_energy_by_country,
  condition = \(x\) is.na(x),
  f = \(x\) 0,
  how = "replace"
)
```

• how = "list"

replaces elements x satisfying condition and classes by f(x) and others by deflt maintaining list structure:

```
# replace all missing values by 0
rrapply(
  renewable_energy_by_country,
  condition = \(x) !is.na(x),
  delft = 0,
  how = "list"
)
```

• how = "unlist"

similar to how = "list" unlisting the returned
result:

```
# replace missing values by 0 and unlist
rrapply(
   renewable_energy_by_country,
   classes = "numeric",
   delft = 0,
   how = "unlist"
)
```

• how = "prune"

similar to **how = "replace"** pruning all elements not subject to f:

```
# prune all missing values and maintain
# list structure
rrapply(
   renewable_energy_by_country,
   condition = \(x) !is.na(x),
   how = "prune"
)
```

• how = "flatten"

similar to how = "prune" returning a flattened
unnested pruned list. Coercion is the same as
how = "unlist" (using the default options):

```
# prune all missing values and return
# flattened list
rrapply(
  renewable_energy_by_country,
  condition = \(x\) !is.na(x),
  how = "flatten"
)
```

• how = "melt"

similar to **how = "prune"** returning a melted data.frame of the pruned list with columns L1, L2, ..., value. Each row contains the path and value of an element x:

```
# prune all missing values and melt list
l <- rrapply(
  renewable_energy_by_country,
  condition = \(x) !is.na(x),
  how = "melt"
)</pre>
```

• how = "unmelt"

reconstructs a nested list from a melted data.frame as returned by **how = "melt"**:

```
# unmelt data.frame back to nested list
rrapply(l, how = "unmelt")
```

• how = "bind"

similar to **how** = "**prune**" unnesting repeated list elements into a wide data.frame. Each repeated element expands to a single row with columns aligned by names:

```
# unnest to wide data.frame and include
# parent node names as columns L1, L2, ...
rrapply(
   pokedex,
```

unnest repeated list to wide data.frame

• how = "recurse"

how = "bind"

similar to **how** = **"replace"** but recurses further into modified elements satisfying condition and classes after application of f:

options = list(namecols = T)

```
# recursively remove all list attributes
rrapply(
  renewable_energy_by_country,
  f = \(x) c(x),
  classes = c("list", "ANY"),
  how = "recurse"
)
```

• how = "names"

similar to how = "recurse" replacing the name
of element x by f(x) instead of its content using
classes = c("list", "ANY") by default:

```
## recursively capitalize all names in list
rrapply(
   renewable_energy_by_country,
   f = \(x, .xname) toupper(.xname),
   how = "names"
)
```

Special arguments .xname, .xpos, .xparents and .xsiblings

The f and condition functions accept four special arguments in addition to the principal argument:

• .xname evaluates to the name of the current list element:

```
# filter list elements by name
rrapply(
  renewable_energy_by_country,
  condition = \(x, .xname\) .xname == "Belgium"
  how = "prune"
)
```

• .xpos evaluates to the position of the element in the nested list as an integer vector:

```
# return position of element in list
rrapply(
  renewable_energy_by_country,
  condition = \((x, .xname) .xname == "Belgium",
  f = \((x, .xpos) .xpos,
  how = "flatten"
)
```

•.xparents evaluates to the vector of parent names of the current element. .xsiblings evaluates to the parent list containing the current element and its direct siblings: