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<https://www.sciviews.org/data.trame/>,
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Contents

data.trame-package	2
data.trame	3
dcast	5
let_data.table_to_data.trame	7
melt	8
methods	9
print.data.trame	12
subsetting	13
Index	16

data.trame-package *A better 'data.frame' for 'SciViews::R'*

Description

The 'data.trame' object is an hybrid between 'data.table', 'tibble' and 'data.frame'. It enhances the 'data.frame' with the speed of 'data.table' and the nice features of 'tibble' (petty printing, stricter rules...).

Important functions

- `data.trame()` to construct a data.trame object,
- `as.data.trame()` to coerce into a data.trame object.
- `is.data.trame()` to test for data.trame objects.
- Methods for data.frame objects.

Author(s)

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See Also

Useful links:

- <https://github.com/SciViews/data.trame>
- <https://www.sciviews.org/data.trame/>
- <https://sciviews.r-universe.dev/data.trame>
- Report bugs at <https://github.com/SciViews/data.trame/issues>

data.frame	<i>Build, coerce and test for 'data.frame' objects</i>
------------	--

Description

Build, coerce and test for 'data.frame' objects

Usage

```
data.frame(  
  ...,  
  .key = NULL,  
  .rows = NULL,  
  .name_repair = c("check_unique", "unique", "universal", "minimal")  
)  
  
as.data.frame(  
  x,  
  .key = NULL,  
  .rows = NULL,  
  .rownames = NA,  
  .name_repair = c("check_unique", "unique", "universal", "minimal"),  
  ...  
)  
  
## Default S3 method:  
as.data.frame(  
  x,  
  .key = NULL,  
  .rows = NULL,  
  .rownames = NA,  
  .name_repair = c("check_unique", "unique", "universal", "minimal"),  
  ...  
)  
  
## S3 method for class 'list'  
as.data.frame(  
  x,  
  .key = NULL,  
  .rows = NULL,  
  .rownames = NA,  
  .name_repair = c("check_unique", "unique", "universal", "minimal"),  
  ...  
)  
  
## S3 method for class 'data.frame'  
as.data.frame(  
  x,  
  .key = NULL,  
  .rows = NULL,  
  .rownames = NA,  
  .name_repair = c("check_unique", "unique", "universal", "minimal"),  
  ...  
)
```

```

    x,
    .key = NULL,
    .rows = NULL,
    .rownames = NA,
    .name_repair = c("check_unique", "unique", "universal", "minimal"),
    ...
  )

## S3 method for class 'data.frame'
as.data.frame(
  x,
  .key = NULL,
  .rows = NULL,
  .rownames = NA,
  .name_repair = c("check_unique", "unique", "universal", "minimal"),
  ...
)

## S3 method for class 'data.table'
as.data.frame(
  x,
  .key = NULL,
  .rows = NULL,
  .rownames = NA,
  .name_repair = c("check_unique", "unique", "universal", "minimal"),
  ...
)

## S3 method for class 'data.frame'
as.data.table(x, keep.rownames = FALSE, ...)

## S3 method for class 'data.frame'
as.data.frame(x, row.names = NULL, optional = TRUE, ..., keep.attr = FALSE)

## S3 method for class 'data.frame'
as_tibble(
  x,
  ...,
  .rows = NULL,
  .name_repair = c("check_unique", "unique", "universal", "minimal"),
  rownames = NULL,
  keep.attr = FALSE
)

is.data.frame(x)

```

Arguments

... A set of name-value pairs that constitute the 'data.frame'.

<code>.key</code>	A character vector with the name of the columns to use as key (sorting the data)
<code>.rows</code>	The number of rows in the final 'data.frame'. Useful to create a 0-column object, or for additional check.
<code>.name_repair</code>	Treatment of problematic column names: could be "check_unique" (default, no name repair but check they are unique), "unique" (make sure names are unique), "universal" (make sure names are unique and syntactically correct), "minimal" (no check or name repair except for existence), or a function for custom name repair, e.g., <code>.name_repair = make.names</code> for base R style.
<code>x</code>	An object.
<code>.rownames</code>	The name of the column that holds the row names of the original object, if any. If NA (default), row names are left intact. If NULL row names are removed.
<code>keep.rownames</code>	For compatibility with the generic, but not used here
<code>row.names</code>	A character vector of row names, or NULL (default).
<code>optional</code>	If TRUE, the row names are not checked for uniqueness. If FALSE, the row names are corrected with <code>make.names()</code> .
<code>keep.attr</code>	If TRUE, the attributes of the data frame are kept.
<code>rownames</code>	NULL remove row names, NA (default) leaves them, or a single string to create a column with that name.

Value

A 'data.frame' object, which is indeed a 'data.frame'/'data.frame' object, thus subclassing 'data.frame'. `is.data.frame()` returns TRUE if the object is a 'data.frame', and FALSE otherwise.

Examples

```
dtrm <- data.frame(
  a = 1:3,
  b = letters[1:3],
  c = factor(LETTERS[1:3]), .key = c('a', 'b'), .rows = 3
)
is.data.frame(dtrm)
```

dcast

Fast dcast for data.frame

Description

`dcast()` transforms a `data.frame` from long to wide format, a little bit like `tidyr::pivot_wider()`. See `data.table::dcast()` for explanations.

Usage

```
dcast(
  data,
  formula,
  fun.aggregate = NULL,
  ...,
  margins = NULL,
  subset = NULL,
  fill = NULL,
  value.var = guess(data)
)

## S3 method for class 'data.frame'
dcast(
  data,
  formula,
  fun.aggregate = NULL,
  sep = "_",
  ...,
  margins = NULL,
  subset = NULL,
  fill = NULL,
  drop = TRUE,
  value.var = guess(data),
  verbose = getOption("datatable.verbose")
)
```

Arguments

<code>data</code>	A <code>data.frame</code> object.
<code>formula</code>	A formula LHS ~ RHS, see <code>data.table::dcast()</code> for details.
<code>fun.aggregate</code>	The function to aggregate multiple data before dcasting.
<code>...</code>	Arguments passed to the aggregating function.
<code>margins</code>	Note implemented yet.
<code>subset</code>	Should the dcasting be done on a subset of the data?
<code>fill</code>	Value with which to fill missing cells.
<code>value.var</code>	The name of the column to use as value variable. If not provided it is "guessed" the <code>guess()</code> internal function is used to build a good default name.
<code>sep</code>	Character vector of length 1, used to separate parts of the variable names (<code>_</code> by default).
<code>drop</code>	FALSE should dcast include all missing combinations?
<code>verbose</code>	Not used yet.

Value

A keyed `data.frame` is returned with the dcasted data.

See Also

[data.table::dcast\(\)](#), [melt\(\)](#)

Examples

```
# Adapted from first example of ?dcast.data.table
ChickWeight = as.data.frame(ChickWeight)
ChickWeight <- set_names(ChickWeight, tolower(names(ChickWeight)))
dtrm <- melt(ChickWeight, id.vars = 2:4)
dcast(dtrm, time ~ variable, fun.aggregate = mean)
```

```
let_data.table_to_data.frame
```

Switch back and forth quickly between data.frame and data.table by reference

Description

These functions just change the class of the object by reference (data.frame and data.table objects are, internally, identical except for their class). These functions are intended only for programmers! There is no check that the object is correct before the change.

Usage

```
let_data.table_to_data.frame(x)
```

```
let_data.frame_to_data.table(x)
```

Arguments

`x` A valid data.frame or data.table object

Value

A data.table or data.frame object.

Examples

```
dtrm <- data.frame(a = 1:3, b = letters[1:3], c = factor(LETTERS[1:3]))
class(dtrm)
let_data.frame_to_data.table(dtrm)
class(dtrm)
let_data.table_to_data.frame(dtrm)
class(dtrm)
```

```
# Whenever you need to use a data.table method on a data.frame in a function,
# you can do something like this:
test_fun <- function(x, i, expr) {
```

```

    force(i) # Make sure i is evaluated before changing the class of x
    # value will be evaluated with x being a data.table here!
    let_data.frame_to_data.table(x)
    on.exit(let_data.table_to_data.frame(x))
    print(class(x)) # Internally, it is now a data.table
    expr
  }
  test_fun(dtrm, 1:2, class(dtrm))
  class(dtrm) # Back to data.frame

```

melt

Fast melt for data.frame

Description

`melt()` is reshaping wide-to-long, notb unlike `tidyr::pivot_longer()` for `data.frame` objects. See [data.table::melt\(\)](#) for explanations.

Usage

```
melt(data, ..., na.rm = FALSE, value.name = "value")
```

```

## S3 method for class 'data.frame'
melt(
  data,
  id.vars,
  measure.vars,
  variable.name = "variable",
  value.name = "value",
  ...,
  na.rm = FALSE,
  variable.factor = TRUE,
  value.factor = FALSE,
  verbose = getOption("datatable.verbose")
)

```

Arguments

<code>data</code>	A <code>data.frame</code> object.
<code>...</code>	Arguments passed to other methods.
<code>na.rm</code>	Should NA values be removed? Default is FALSE.
<code>value.name</code>	Name for the molten data values column(s).
<code>id.vars</code>	Vector of id variables.
<code>measure.vars</code>	Measure variables for melting. Can be missing.
<code>variable.name</code>	Name (default <code>variable</code>) of output column containing the information about melted columns.

variable.factor	If TRUE, the variable column is converted to a factor, otherwise, it is a character column.
value.factor	If TRUE, the value column is converted to a factor, else, it is left unchanged.
verbose	TRUE turns on status and information messages.

Value

An unkeyed data.frame containing the molten data.

See Also

[data.table::melt\(\)](#), [dcast\(\)](#)

Examples

```
# Adapted from first example of ?melt.data.table
set.seed(45)
library(data.table)
dtrm <- data.frame(
  i_1 = c(1:5, NA),
  n_1 = c(NA, 6, 7, 8, 9, 10),
  f_1 = factor(sample(c(letters[1:3], NA), 6L, TRUE)),
  f_2 = ordered(c("z", "a", "x", "c", "x", "x")),
  c_1 = sample(c(letters[1:3], NA), 6L, TRUE),
  c_2 = sample(c(LETTERS[1:2], NA), 6L, TRUE),
  d_1 = as.Date(c(1:3, NA, 4:5), origin = "2013-09-01"),
  d_2 = as.Date(6:1, origin = "2012-01-01")
)
# add a couple of list cols
dtrm$l_1 <- dtrm[, ~list(c = list(rep(i_1, sample(5, 1L))))], by = ~i_1$c
dtrm$l_2 <- dtrm[, ~list(c = list(rep(c_1, sample(5, 1L))))], by = ~i_1$c

# id.vars, measure.vars as character/integer/numeric vectors
melt(dtrm, id.vars = 1:2, measure.vars = "f_1")
```

Description

These methods handle data.frame objects correctly, so that they remain internally consistent with data.tables.

Usage

```
## S3 method for class 'data.frame'
head(x, n = 6L, ...)

## S3 method for class 'data.frame'
tail(x, n = 6L, ...)

## S3 replacement method for class 'data.frame'
names(x) <- value

set_names()

let_names(x, value)

## S3 replacement method for class 'data.frame'
row.names(x) <- value

set_row_names(x, value)

let_row_names(x, value)

## S3 method for class 'data.frame'
edit(name, ...)

## S3 method for class 'data.frame'
cbind(
  x,
  ...,
  keep.rownames = FALSE,
  check.names = FALSE,
  key = NULL,
  stringsAsFactors = FALSE
)

## S3 method for class 'data.frame'
rbind(
  x,
  ...,
  use.names = TRUE,
  fill = FALSE,
  idcol = NULL,
  ignore.attr = FALSE
)
```

Arguments

x	A data.frame object.
n	The number of rows to keep.

...	Further parameters (not used yet).
value	The value passed to the method.
name	The name of the data.frame to edit.
keep.rownames	If TRUE, the row names are kept as a column
check.names	If TRUE, the names of the columns are checked
key	The key to set on the resulting data.frame. If NULL, no key is set.
stringsAsFactors	If TRUE, character columns are converted to factors.
use.names	If TRUE, the names of the columns are matched. If FALSE, match is done by position. if "check", warn if names do not match.
fill	If TRUE, fill missing columns with NA. By default FALSE.
idcol	Create a column with ids showing where the data come from. With TRUE, the column is named id. With idcol = "col_name", it has that name.
ignore.attr	If TRUE, ignore attributes when binding.

Value

head() and tail() return truncated data.frame objects (n first or last rows). names(dtrm) <- value and set_names() both set names (colnames). let_names() change names by reference. row.names(dtrm) <- value and set_row_names() both set the row names of a data.frame. However, only the second one keeps the selfref pointer integrity. let_row_names() is a faster version, but it changes the row names by reference. With all let_xxx() functions, you need to take extra care to avoid unexpected side effects, see examples. cbind() combines data.frames by columns, rbind() combines them by rows.

Examples

```
dtrm <- data.frame(
  a = 1:10,
  b = letters[1:10],
  c = factor(LETTERS[1:10]), .key = c('a', 'b')
)
head(dtrm)
tail(dtrm, n = 3L)
cbind(dtrm, dtrm)
rbind(dtrm, dtrm)
dtrm2 <- set_row_names(dtrm, paste("row", letters[1:10]))
dtrm2
dtrm # Not changed
# Take care with let_xxx() functions: it propagates changes to other
# data.frames if you did not used copy()!
dtrm2 <- dtrm
dtrm3 <- data.table::copy(dtrm)
let_row_names(dtrm2, paste("row", letters[11:20]))
dtrm2 # OK
dtrm # Also changed!
dtrm3 # Not changed, because created using copy()
```

print.data.trame *Printing data.frames*

Description

A data.frame prints almost like a tibble.

Usage

```
## S3 method for class 'data.frame'
print(
  x,
  width = NULL,
  ...,
  n = NULL,
  max_extra_cols = NULL,
  max_footer_lines = NULL
)

## S3 method for class 'data.frame'
format(
  x,
  width = NULL,
  ...,
  n = NULL,
  max_extra_cols = NULL,
  max_footer_lines = NULL
)

## S3 method for class 'datatrame'
obj_sum(x)

## S3 method for class 'data.trame'
obj_sum(x)

## S3 method for class 'datatrame'
tbl_sum(x, ...)

## S3 method for class 'data.trame'
tbl_sum(x, ...)

## S3 method for class 'data.trame'
tbl_nrow(x, ...)
```

```
## S3 method for class 'data.frame'
str(object, ..., indent.str = " ", nest.lev = 0)
```

Arguments

<code>x</code>	A <code>data.frame</code> object.
<code>width</code>	The width of the text output. If <code>NULL</code> , the default, <code>getOption("width")</code> is used.
<code>...</code>	Additional arguments passed to <code>format()</code> .
<code>n</code>	The number of rows to print. If <code>NULL</code> , a default number is used.
<code>max_extra_cols</code>	The maximum number of extra columns to print abbreviated. If <code>NULL</code> by default, a reasonable default value is used.
<code>max_footer_lines</code>	Maximum number of lines in the footer. If <code>NULL</code> , a reasonable default value is used.
<code>object</code>	A <code>data.frame</code> .
<code>indent.str</code>	The string used for indentation.
<code>nest.lev</code>	The current nesting level, used for recursive printing.

Examples

```
dtrm <- data.frame(
  a = -1:3,
  b = letters[1:5],
  c = factor(LETTERS[1:5]),
  d = c(TRUE, FALSE, TRUE, NA, TRUE), .key = c('a', 'b'), .rows = 5
)
dtrm
str(dtrm)
```

subsetting

Subsetting data.frames

Description

Subsetting `data.frames` uses a syntax similar to `tibble`, or formulas for `i`, `j`, and possibly `by` or `keyby` to use the `data.table` syntax instead.

Usage

```
## S3 method for class 'data.frame'
x[i, j, by, keyby, with = TRUE, drop = FALSE, ...]

set_(x, i, j, value, byref = FALSE)

let_(x, i = NULL, j = seq_along(x), value)
```

Arguments

x	A data.frame object.
i	Selection of rows by indices, negative indices, logical or a formula
j	Selection of columns by indices, negative indices, logical, names or a formula (both i and j must be formulas simultaneously). If := is used in the formula to create one or more new variables by reference, the expression must be placed between {} to avoid operators precedence issues, or better: := could be just replaced by ~.
by	Grouping columns (must be a formula and j must be also provided as a formula)
keyby	Either TRUE/FALSE if by is provided, or a formula (and j must also be provided as a formula)
with	Logical, whether to evaluate j in the data.frame if TRUE or in the calling environment if FALSE (default is TRUE). with = FALSE is similar to tibble subsetting and it is forced when i or j are not formulas.
drop	Coerce to a vector if the returned data.frame only has one column
...	Further arguments passed to the underlying data.table subsetting
value	The value to insert as subassignment in a data.frame object.
byref	Logical, whether to use by reference or not (FALSE by default).

Value

A data.frame object, or a vector if drop = TRUE and the result has only one column.

Examples

```
dtrm <- data.frame(
  a = 1:3,
  b = letters[1:3],
  c = factor(LETTERS[1:3])
)
# Subsetting rows, the tibble-way
dtrm[1:2, ]
dtrm[-1, ]
dtrm[c(TRUE, FALSE, TRUE), ]
# On the contrary to data.table, providing only one arg, means subsetting
# columns (like for data.frame or tibble)
dtrm[c(TRUE, FALSE, TRUE)]
dtrm[dtrm$a > 1, ] # Must fully qualify the column name
# Subsetting the data.table way, with formulas: no fully qualification needed
dtrm[~a > 1, ]

# Subsetting the columns, the tibble way
dtrm[, 1:2]
dtrm[, -1]
dtrm[, c(TRUE, FALSE, TRUE)]
dtrm[, c("a", "b")]
# You must set drop = TRUE explicitly to return a vector
dtrm[, 2] # Still a data.frame, like tibble, but unlike the data.frame method
```

```

dtrm[, 2, drop = TRUE] # Now a vector
# The selection is referentially transparent, i.e., you can do:
sel <- c("c", "b")
dtrm[, sel]
# Subsetting the columns, the data.table way, with formulas
dtrm[~1:2, ~(b)]
dtrm[~1:2, ~b] # If not enclosed in .(), returns a vector instead
# Precautions are needed here because it is NOT referentially transparent:
dtrm[, ~..sel] # In data.table language, this is how you access `sel`

# Extended data.table syntax using i, j, by, or keyby with formulas
# Warning: due to precedence of operators, you must use braces here!
dtrm[, ~{d := paste0(b, c)}] # Changed in place (by reference!)
# Another form that does not need braces, but is less readable:
dtrm[, ~`:=`(e, paste0(b, a))]
# or equivalently:
dtrm[, ~let(e = paste0(b, a))]
# In this case, it is much better to just replace `:=` by `~`, but internally
# it uses set(). It is faster, but much more limited and cannot use by or
# or keyby:
dtrm[, f ~ paste0(c, a)]
# One can also use standard evaluation in that case using with = FALSE
dtrm[, f ~ paste0(dtrm$c, dtrm$a), with = FALSE]
#
# Take care when you provide only one argument:
# If it is a formula, the data.table syntax is used (select rows)
# otherwise, the data.frame syntax applies, and columns are selected!
dtrm[1:2] # All rows and 2 first columns
dtrm[~1:2] # All columns and 2 first rows!

# For $, on the contrary to data.frame/data.table, but like tibble,
# no partial match is allowed (returns NULL with a warning)
dtrm$count <- dtrm$c
names(dtrm)
dtrm$count #OK
#dtrm$co # Not OK, no partial match allowed

```

Index

[.data.frame (subsetting), 13

as.data.frame.data.frame (data.frame), 3

as.data.table.data.frame (data.frame), 3

as.data.frame (data.frame), 3

as.data.frame(), 2

as_tibble.data.frame (data.frame), 3

cbind.data.frame (methods), 9

data.table::dcast(), 5–7

data.table::melt(), 8, 9

data.frame, 3

data.frame(), 2

data.frame-package, 2

dcast, 5

dcast(), 9

edit.data.frame (methods), 9

format.data.frame (print.data.frame), 12

head.data.frame (methods), 9

is.data.frame (data.frame), 3

is.data.frame(), 2

let_ (subsetting), 13

let_data.table_to_data.frame, 7

let_data.frame_to_data.table
(let_data.table_to_data.frame),
7

let_names (methods), 9

let_row_names (methods), 9

make.names(), 5

melt, 8

melt(), 7

methods, 9

names<- .data.frame (methods), 9

obj_sum.data.frame (print.data.frame),
12

obj_sum.datatframe (print.data.frame), 12

print.data.frame, 12

rbind.data.frame (methods), 9

row.names<- .data.frame (methods), 9

set_ (subsetting), 13

set_names (methods), 9

set_row_names (methods), 9

str.data.frame (print.data.frame), 12

subsetting, 13

tail.data.frame (methods), 9

tbl_nrow.data.frame (print.data.frame),
12

tbl_sum.data.frame (print.data.frame),
12

tbl_sum.datatframe (print.data.frame), 12