

# Package ‘maditr’

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**Type** Package

**Title** Fast Data Aggregation, Modification, and Filtering with Pipes and 'data.table'

**Version** 0.7.4

**Maintainer** Gregory Demin <gdemin@gmail.com>

**Depends** R (>= 3.3.0)

**Imports** data.table (>= 1.12.6), magrittr (>= 1.5)

**Suggests** knitr, tinytest, utils, rmarkdown

## Description

Provides pipe-style interface for 'data.table'. Package preserves all 'data.table' features without significant impact on performance. 'let' and 'take' functions are simplified interfaces for most common data manipulation tasks. For example, you can write 'take(mtcars, mean(mpg), by = am)' for aggregation or 'let(mtcars, hp\_wt = hp/wt, hp\_wt\_mpg = hp\_wt/mpg)' for modification. Use 'take\_if/let\_if' for conditional aggregation/modification. 'query\_if' function translates its arguments one-to-one to '['.data.table' method. Additionally there are some conveniences such as automatic 'data.frame' conversion to 'data.table'.

**License** GPL-2

**URL** <https://github.com/gdemin/maditr>

**BugReports** <https://github.com/gdemin/maditr/issues>

**VignetteBuilder** knitr

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**Author** Gregory Demin [aut, cre]

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coalesce	<i>Return first non-missing element</i>
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**Description**

It is an alias for `data.table::fcoalesce`. For details see [fcoalesce](#)

**Usage**

```
coalesce(...)
```

**Arguments**

...           vectors

**Value**

A vector the same length as the first ... argument with NA values replaced by the first non-missing value.

**Examples**

```
# examples from dplyr
x = sample(c(1:5, NA, NA, NA))
coalesce(x, 0L)

y = c(1, 2, NA, NA, 5)
z = c(NA, NA, 3, 4, 5)
coalesce(y, z)
```

---

`dcast`*Convert data between wide and long forms.*

---

### Description

The `dcast` formula takes the form `LHS ~ RHS`, ex: `var1 + var2 ~ var3`. The order of entries in the formula is essential. There are two special variables: `.` and `...`. `.` represents no variable; `...` represents all variables not otherwise mentioned in formula. LHS variable values will be in rows. RHS variables values will become column names. `fun.aggregate(value.var)` will be cell values. For details see [dcast](#) and [melt](#).

### Usage

```
dcast(  
  data,  
  formula,  
  fun.aggregate = NULL,  
  sep = "_",  
  ...,  
  margins = NULL,  
  subset = NULL,  
  fill = NULL,  
  drop = TRUE,  
  value.var = guess(data),  
  verbose = getOption("datatable.verbose")  
)
```

```
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")  
)
```

```
guess(data)
```

### Arguments

`data` A `data.table`/`data.frame`. `data.frame` will be automatically converted to `data.table`.  
`formula` A formula of the form `LHS ~ RHS` to cast. For details see [dcast](#).

<code>fun.aggregate</code>	Should the data be aggregated before casting? If the formula doesn't identify a single observation for each cell, then aggregation defaults to length with a message.
<code>sep</code>	Character vector of length 1, indicating the separating character in variable names generated during casting. Default is <code>_</code> for backwards compatibility.
<code>...</code>	Any other arguments that may be passed to the aggregating function.
<code>margins</code>	For details see <a href="#">dcast</a> .
<code>subset</code>	Specified if casting should be done on a subset of the data.
<code>fill</code>	Value with which to fill missing cells. If <code>fun.aggregate</code> is present, takes the value by applying the function on a 0-length vector.
<code>drop</code>	<code>FALSE</code> will cast by including all missing combinations. <code>c(FALSE, TRUE)</code> will only include all missing combinations of formula LHS. And <code>c(TRUE, FALSE)</code> will only include all missing combinations of formula RHS.
<code>value.var</code>	Name of the column whose values will be filled to cast. Function <code>'guess()'</code> tries to, well, guess this column automatically, if none is provided. It is possible to cast multiple <code>'value.var'</code> columns simultaneously. For details see <a href="#">dcast</a> .
<code>verbose</code>	For details see <a href="#">dcast</a> .
<code>id.vars</code>	vector of id variables. Can be integer (corresponding id column numbers) or character (id column names) vector. If missing, all non-measure columns will be assigned to it. If integer, must be positive; see Details.
<code>measure.vars</code>	Measure variables for melting. Can be missing, vector, list, or pattern-based. For details see <a href="#">dcast</a> .
<code>variable.name</code>	name for the measured variable names column. The default name is <code>'variable'</code> .
<code>value.name</code>	name for the molten data values column(s). The default name is <code>'value'</code> . Multiple names can be provided here for the case when <code>measure.vars</code> is a list, though note well that the names provided in <code>measure.vars</code> take precedence.
<code>na.rm</code>	If <code>TRUE</code> , NA values will be removed from the molten data.
<code>variable.factor</code>	If <code>TRUE</code> , the variable column will be converted to factor, else it will be a character column.
<code>value.factor</code>	If <code>TRUE</code> , the value column will be converted to factor, else the molten value type is left unchanged.

**Value**

`data.table`

**Author(s)**

Matt Dowle [mattjdowle@gmail.com](mailto:mattjdowle@gmail.com)

**Examples**

```
# examples from 'tidyr' package
stocks = data.frame(
  time = as.Date('2009-01-01') + 0:9,
  X = rnorm(10, 0, 1),
  Y = rnorm(10, 0, 2),
  Z = rnorm(10, 0, 4)
)
stocksm = stocks %>%
  melt(id.vars = "time", variable.name = "stock", value.name = "price")
stocksm %>% dcast(time ~ stock)
stocksm %>% dcast(stock ~ time)

# dcast and melt are complements
df = data.frame(x = c("a", "b"), y = c(3, 4), z = c(5, 6))
df %>%
  dcast(z ~ x, value.var = "y") %>%
  melt(id.vars = "z", variable.name = "x", value.name = "y", na.rm = TRUE)
```

dt\_count

*Additional useful functions***Description**

- dt\_count calculates number of cases by groups, possibly weighted. dt\_add\_count adds number of cases to existing dataset.
- dt\_top\_n returns top n rows from each group.

**Usage**

```
dt_count(data, ..., weight = NULL, sort = FALSE, name = "n")
```

```
dt_add_count(data, ..., weight = NULL, sort = FALSE, name = "n")
```

```
dt_top_n(data, n, by, order_by = NULL)
```

**Arguments**

data	data.table/data.frame data.frame will be automatically converted to data.table.
...	variables to group by.
weight	optional. Unquoted variable name. If provided result will be the sum of this variable by groups.
sort	logical. If TRUE result will be sorted in descending order by resulting variable.
name	character. Name of resulting variable.
n	numeric. number of top cases. If n is negative then bottom values will be returned.

by list or vector of grouping variables  
 order\_by unquoted variable name by which result will be sorted. If not specified, defaults to the last variable in the dataset.

### Value

data.table

### Examples

```
data(mtcars)

# dt_count
dt_count(mtcars, am, vs)
dt_add_count(mtcars, am, vs, name = "am_vs")[] # [] for autoprinting

# dt_top_n
dt_top_n(mtcars, 2, by = list(am, vs))
dt_top_n(mtcars, 2, order_by = mpg, by = list(am, vs))
```

---

dt\_left\_join *Join two data.frames by common columns.*

---

### Description

Do different versions of SQL join operations. See examples.

### Usage

```
dt_left_join(x, y, by = NULL, suffix = c(".x", ".y"))
dt_right_join(x, y, by = NULL, suffix = c(".x", ".y"))
dt_inner_join(x, y, by = NULL, suffix = c(".x", ".y"))
dt_full_join(x, y, by = NULL, suffix = c(".x", ".y"))
dt_semi_join(x, y, by = NULL)
dt_anti_join(x, y, by = NULL)
```

### Arguments

x data.frame or data.table  
 y data.frame or data.table

by	a character vector of variables to join by. If NULL, the default, *_join() will do a natural join, using all variables with common names across the two tables. A message lists the variables so that you can check they're right (to suppress the message, simply explicitly list the variables that you want to join). To join by different variables on x and y use a named vector. For example, by = c("a" = "b") will match x.a to y.b.
suffix	If there are non-joined duplicate variables in x and y, these suffixes will be added to the output to disambiguate them. Should be a character vector of length 2.

**Value**

data.table

**Examples**

```
workers = fread("
  name company
  Nick Acme
  John Ajax
  Daniela Ajax
")

positions = fread("
  name position
  John designer
  Daniela engineer
  Cathie manager
")

workers %>% dt_inner_join(positions)
workers %>% dt_left_join(positions)
workers %>% dt_right_join(positions)
workers %>% dt_full_join(positions)

# filtering joins
workers %>% dt_anti_join(positions)
workers %>% dt_semi_join(positions)

# To suppress the message, supply 'by' argument
workers %>% dt_left_join(positions, by = "name")

# Use a named 'by' if the join variables have different names
positions2 = setNames(positions, c("worker", "position")) # rename first column in 'positions'
workers %>% dt_inner_join(positions2, by = c("name" = "worker"))
```

## Description

Subset of 'dplyr' verbs to work with `data.table`. Note that there is no `group_by` verb - use `by` or `keyby` argument when needed.

- `dt_mutate` adds new variables or modify existing variables. If `data` is `data.table` then it modifies in-place.
- `dt_summarize` computes summary statistics. Splits the data into subsets, computes summary statistics for each, and returns the result in the "data.table" form.
- `dt_summarize_all` is the same as `dt_summarize` but work over all non-grouping variables.
- `dt_filter` selects rows/cases where conditions are true. Rows where the condition evaluates to NA are dropped.
- `dt_select` selects column/variables from the data set. Range of variables are supported, e.g. `vs:carb`. Characters which start with '^' or end with '\$' considered as Perl-style regular expression patterns. For example, '^Petal' returns all variables started with 'Petal'. 'Width\$' returns all variables which end with 'Width'. Pattern '^.' matches all variables and pattern '^.\*my\_str' is equivalent to contains "my\_str". See examples.
- `dt_arrange` sorts dataset by variable(-s). Use '-' to sort in descending order. If `data` is `data.table` then it modifies in-place.

## Usage

```
dt_mutate(data, ..., by)

dt_summarize(data, ..., by, keyby, fun = NULL)

dt_summarize_all(data, fun, by, keyby)

dt_summarise(data, ..., by, keyby, fun = NULL)

dt_summarise_all(data, fun, by, keyby)

dt_select(data, ...)

dt_filter(data, ...)

dt_arrange(data, ..., na.last = FALSE)
```

## Arguments

<code>data</code>	<code>data.table</code> / <code>data.frame</code> <code>data.frame</code> will be automatically converted to <code>data.table</code> . <code>dt_mutate</code> , <code>dt_mutate_if</code> , <code>dt_mutate_if</code> modify <code>data.table</code> object in-place.
<code>...</code>	List of variables or name-value pairs of summary/modifications functions. The name will be the name of the variable in the result. In the <code>mutate</code> function we can use <code>a = b</code> or <code>a := b</code> notation. Advantages of <code>:=</code> are multiassignment ( <code>c("a", "b") := list(1, 2)</code> ) and parametric assignment ( <code>((a) := 2)</code> ).
<code>by</code>	unquoted name of grouping variable or list of unquoted names of grouping variables. For details see <a href="#">data.table</a>



keyby	Same as by, but with an additional setkey() run on the by columns of the result, for convenience. It is common practice to use 'keyby=' routinely when you wish the result to be sorted. For details see <a href="#">data.table</a> .
fun	function which will be applied to all variables in dt_summarize and dt_summarize_all.
na.last	logical. FALSE by default. If TRUE, missing values in the data are put last; if FALSE, they are put first.

## Examples

```
# examples from 'dplyr'
# newly created variables are available immediately
mtcars %>%
  dt_mutate(
    cyl2 = cyl * 2,
    cyl4 = cyl2 * 2
  ) %>%
  head()

# you can also use dt_mutate() to remove variables and
# modify existing variables
mtcars %>%
  dt_mutate(
    mpg = NULL,
    disp = disp * 0.0163871 # convert to litres
  ) %>%
  head()

# window functions are useful for grouped mutates
mtcars %>%
  dt_mutate(
    rank = rank(-mpg, ties.method = "min"),
    keyby = cyl) %>%
  print()

# You can drop variables by setting them to NULL
mtcars %>% dt_mutate(cyl = NULL) %>% head()

# A summary applied without by returns a single row
mtcars %>%
  dt_summarise(mean = mean(disp), n = .N)

# Usually, you'll want to group first
mtcars %>%
  dt_summarise(mean = mean(disp), n = .N, by = cyl)

# Multiple 'by' - variables
mtcars %>%
  dt_summarise(cyl_n = .N, by = list(cyl, vs))
```

```

# Newly created summaries immediately
# doesn't overwrite existing variables
mtcars %>%
  dt_summarise(displ = mean(displ),
               sd = sd(displ),
               by = cyl)

# You can group by expressions:
mtcars %>%
  dt_summarise_all(mean, by = list(vsam = vs + am))

# filter by condition
mtcars %>%
  dt_filter(am==0)

# filter by compound condition
mtcars %>%
  dt_filter(am==0, mpg>mean(mpg))

# select
mtcars %>% dt_select(vs:carb, cyl)
mtcars %>% dt_select(-am, -cyl)

# regular expression pattern
dt_select(iris, "^Petal") # variables which start from 'Petal'
dt_select(iris, "Width$") # variables which end with 'Width'
# move Species variable to the front.
# pattern "^." matches all variables
dt_select(iris, Species, "^.")
# pattern "^.*i" means "contains 'i'"
dt_select(iris, "^.*i")
dt_select(iris, 1:4) # numeric indexing - all variables except Species

# sorting
dt_arrange(mtcars, cyl, displ)
dt_arrange(mtcars, -displ)

```

---

let\_if

---

*Modify, aggregate, select or filter data.frame/data.table*


---

### Description

- let adds new variables or modify existing variables. 'let\_if' make the same thing on the subset of rows.
- take/take\_if aggregate data or select subset of the data by rows or columns.
- let\_all applies expressions to all variables in the dataset. It is also possible to modify the subset of the variables.

- `take_all` aggregates all variables in the dataset. It is also possible to aggregate the subset of the variables.

All functions return `data.table`. Expression in the `'take_all'` and `'let_all'` can use predefined variables: `'.x'` is a value of current variable, `'.name'` is a name of the variable and `'.index'` is sequential number of the variable. `'.value'` is an alias to `'.x'`.

- Add new variables: `let(mtcars, new_var = 42, new_var2 = new_var*hp)`
- Filter data: `take_if(mtcars, am==0)`
- Select variables: `take(mtcars, am, vs, mpg)`
- Aggregate data: `take(mtcars, mean_mpg = mean(mpg), by = am)`
- Aggregate all non-grouping columns: `take_all(mtcars, mean = mean(.x), sd = sd(.x), n = .N, by = am)`
- Aggregate all numeric columns: `take_all(iris, if(is.numeric(.x)) mean(.x))`
- To modify all non-grouping variables:

```
iris %>%
  let_all(
    scaled = (.x - mean(.x))/sd(.x),
    by = Species) %>%
  head()
```

- Aggregate specific columns: `take_all(iris, if(startsWith(.name, "Sepal"))) mean(.x)`

## Usage

```
let_if(data, i, ..., by, keyby)
```

```
take_if(data, i, ..., by, keyby, .SDcols, autoname = TRUE, fun = NULL)
```

```
take(data, ..., by, keyby, .SDcols, autoname = TRUE, fun = NULL)
```

```
let(data, ..., by, keyby)
```

```
## S3 method for class 'data.frame'
```

```
let(data, ..., by, keyby, i)
```

```
sort_by(data, ..., na.last = FALSE)
```

```
let_all(data, ..., by, keyby, .SDcols, suffix = TRUE, sep = "_", i)
```

```
take_all(data, ..., by, keyby, .SDcols, suffix = TRUE, sep = "_", i)
```

**Arguments**

data	data.table/data.frame data.frame will be automatically converted to data.table. let modify data.table object in-place.
i	integer/logical vector. Supposed to use to subset/conditional modifications of data. For details see <a href="#">data.table</a>
...	List of variables or name-value pairs of summary/modifications functions. The name will be the name of the variable in the result. In the let and take functions we can use a = b or a := b notation. Advantages of := is parametric assignment, e. g. (a) := 2 create variable with name which are stored in a. In let := can be used for multiassignment (c("a", "b") := list(1, 2)). Expression in the 'take_all' and 'let_all' can use predefined variables: '.x' is a value of current variable, '.name' is a name of the variable and '.index' is sequential number of the variable. '.value' is is an alias to '.x'.
by	unquoted name of grouping variable of list of unquoted names of grouping variables. For details see <a href="#">data.table</a>
keyby	Same as by, but with an additional setkey() run on the by columns of the result, for convenience. It is common practice to use 'keyby=' routinely when you wish the result to be sorted. For details see <a href="#">data.table</a> .
.SDcols	Specifies the columns of x to be included in the special symbol .SD which stands for Subset of data.table. May be character column names or numeric positions. For details see <a href="#">data.table</a> .
autoname	logical. TRUE by default. Should we create names for unnamed expressions in take?
fun	Function which will be applied to all variables in take. If there are no variables in take then it will be applied to all non-grouping variables in the data.
na.last	logical. FALSE by default. If TRUE, missing values in the data are put last; if FALSE, they are put first.
suffix	logical TRUE by default. For 'let_all'/'take_all'. If TRUE than we append summary name to the end of the variable name. If FALSE summary name will be added at the beginning of the variable name.
sep	character. "_" by default. Separator between the old variables name and prefix or suffix for 'let_all' and 'take_all'.

**Value**

data.table. let returns its result invisibly.

**Examples**

```
# examples form 'dplyr' package
data(mtcars)

# Newly created variables are available immediately
mtcars %>%
  let(
    cyl2 = cyl * 2,
```

```
      cyl4 = cyl2 * 2
    ) %>% head()

# You can also use let() to remove variables and
# modify existing variables
mtcars %>%
  let(
    mpg = NULL,
    disp = disp * 0.0163871 # convert to litres
  ) %>% head()

# window functions are useful for grouped computations
mtcars %>%
  let(rank = rank(-mpg, ties.method = "min"),
    by = cyl) %>%
  head()

# You can drop variables by setting them to NULL
mtcars %>% let(cyl = NULL) %>% head()

# keeps all existing variables
mtcars %>%
  let(displ_l = disp / 61.0237) %>%
  head()

# keeps only the variables you create
mtcars %>%
  take(displ_l = disp / 61.0237)

# can refer to both contextual variables and variable names:
var = 100
mtcars %>%
  let(cyl = cyl * var) %>%
  head()

# filter by condition
mtcars %>%
  take_if(am==0)

# filter by compound condition
mtcars %>%
  take_if(am==0 & mpg>mean(mpg))

# A 'take' with summary functions applied without 'by' argument returns an aggregated data
mtcars %>%
  take(mean = mean(disp), n = .N)

# Usually, you'll want to group first
mtcars %>%
  take(mean = mean(disp), n = .N, by = cyl)
```

```

# You can group by expressions:
mtcars %>%
  take_all(mean, by = list(vsam = vs + am))

# modify all non-grouping variables in-place
mtcars %>%
  let_all((.x - mean(.x))/sd(.x), by = am) %>%
  head()

# modify all non-grouping variables to new variables
mtcars %>%
  let_all(scaled = (.x - mean(.x))/sd(.x), by = am) %>%
  head()

# conditionally modify all variables
iris %>%
  let_all(mean = if(is.numeric(.x)) mean(.x)) %>%
  head()

# modify all variables conditionally on name
iris %>%
  let_all(
    mean = if(startsWith(.name, "Sepal")) mean(.x),
    median = if(startsWith(.name, "Petal")) median(.x),
    by = Species
  ) %>%
  head()

# aggregation with 'take_all'
mtcars %>%
  take_all(mean = mean(.x), sd = sd(.x), n = .N, by = am)

# conditionally aggregate all variables
iris %>%
  take_all(mean = if(is.numeric(.x)) mean(.x))

# aggregate all variables conditionally on name
iris %>%
  take_all(
    mean = if(startsWith(.name, "Sepal")) mean(.x),
    median = if(startsWith(.name, "Petal")) median(.x),
    by = Species
  )

# parametric evaluation:
var = quote(mean(cyl))
mtcars %>%
  let(mean_cyl = eval(var)) %>%
  head()
take(mtcars, eval(var))

# all together

```

```

new_var = "mean_cyl"
mtcars %>%
  let((new_var) := eval(var)) %>%
  head()
take(mtcars, (new_var) := eval(var))

#####

# examples from data.table
dat = data.table(
  x=rep(c("b","a","c"), each=3),
  y=c(1,3,6),
  v=1:9
)

# basic row subset operations
take_if(dat, 2)           # 2nd row
take_if(dat, 3:2)        # 3rd and 2nd row
take_if(dat, order(x))   # no need for order(dat$x)
take_if(dat, y>2)        # all rows where dat$y > 2
take_if(dat, y>2 & v>5)  # compound logical expressions
take_if(dat, !2:4)       # all rows other than 2:4
take_if(dat, -(2:4))     # same

# select|compute columns
take(dat, v)              # v column (as data.table)
take(dat, sum(v))         # return data.table with sum of v (column autonamed 'sum(v)')
take(dat, sv = sum(v))    # same, but column named "sv"
take(dat, v, v*2)        # return two column data.table, v and v*2

# subset rows and select|compute
take_if(dat, 2:3, sum(v)) # sum(v) over rows 2 and 3
take_if(dat, 2:3, sv = sum(v)) # same, but return data.table with column sv

# grouping operations
take(dat, sum(v), by = x) # ad hoc by, order of groups preserved in result
take(dat, sum(v), keyby = x) # same, but order the result on by cols

# all together now
take_if(dat, x!="a", sum(v), by=x) # get sum(v) by "x" for each x != "a"

# more on special symbols, see also ?"data.table::special-symbols"
take_if(dat, .N)          # last row
take(dat, .N)             # total number of rows in DT
take(dat, .N, by=x)      # number of rows in each group

take(dat, .I[1], by=x)    # row number in DT corresponding to each group

# add/update/delete by reference
# [] at the end of expression is for autoprinting
let(dat, grp = .GRP, by=x)[] # add a group counter column

```

```

let(dat, z = 42L)[] # add new column by reference
let(dat, z = NULL)[] # remove column by reference
let_if(dat, x=="a", v = 42L)[] # subassign to existing v column by reference
let_if(dat, x=="b", v2 = 84L)[] # subassign to new column by reference (NA padded)

let(dat, m = mean(v), by=x)[] # add new column by reference by group

# advanced usage
dat = data.table(x=rep(c("b","a","c"), each=3),
                v=c(1,1,1,2,2,1,1,2,2),
                y=c(1,3,6),
                a=1:9,
                b=9:1)

take(dat, sum(v), by=list(y%%2)) # expressions in by
take(dat, sum(v), by=list(bool = y%%2)) # same, using a named list to change by column name
take_all(dat, sum, by=x) # sum of all (other) columns for each group
take(dat,
      MySum=sum(v),
      MyMin=min(v),
      MyMax=max(v),
      by = list(x, y%%2) # by 2 expressions
)

take(dat, seq = min(a):max(b), by=x) # j is not limited to just aggregations
dat %>%
  take(V1 = sum(v), by=x) %>%
  take_if(V1<20) # compound query

dat %>%
  take(V1 = sum(v), by=x) %>%
  sort_by(-V1) %>% # ordering results
  head()

```

---

maditr

*maditr: Pipe-Style Interface for 'data.table'*


---

## Description

Package provides pipe-style interface for `data.table`. It preserves all `data.table` features without significant impact on performance. `'let'` and `'take'` functions are simplified interfaces for most common data manipulation tasks.

- To select rows from data: `take_if(mtcars, am==0)`
- To select columns from data: `take(mtcars, am, vs, mpg)`
- To aggregate data: `take(mtcars, mean_mpg = mean(mpg), by = am)`
- To aggregate all non-grouping columns: `take_all(mtcars, mean, by = am)`



- To conditionally aggregate all non-grouping columns: `take_all(iris, if(is.numeric(.x)) mean(.x))`
- To aggregate several columns with one summary: `take(mtcars, mpg, hp, fun = mean, by = am)`
- To get total summary skip 'by' argument: `take_all(mtcars, mean)`
- Use magrittr pipe '`%>%`' to chain several operations:

```
mtcars %>%
  let(mpg_hp = mpg/hp) %>%
  take(mean(mpg_hp), by = am)
```

- To modify variables or add new variables:

```
mtcars %>%
  let(new_var = 42,
      new_var2 = new_var*hp) %>%
  head()
```

- To modify all non-grouping variables:

```
iris %>%
  let_all(
    scaled = (.x - mean(.x))/sd(.x),
    by = Species) %>%
  head()
```

- To drop variable assign NULL: `let(mtcars, am = NULL) %>% head()`
- To aggregate all variables conditionally on name:

```
iris %>%
  take_all(
    mean = if(startsWith(.name, "Sepal")) mean(.x),
    median = if(startsWith(.name, "Petal")) median(.x),
    by = Species
  )
```

- For parametric assignment use '`:=`':

```
new_var = "my_var"
old_var = "mpg"
mtcars %>%
  let((new_var) := get(old_var)*2) %>%
  head()
```

- For more sophisticated operations see '`query`'/'`query_if`': these functions translates its arguments one-to-one to '`[.data.table]`' method. Additionally there are some conveniences such as automatic '`data.frame`' conversion to '`data.table`'.

## Examples

```
# examples form 'dplyr' package
data(mtcars)

# Newly created variables are available immediately
mtcars %>%
  let(
    cyl2 = cyl * 2,
    cyl4 = cyl2 * 2
  ) %>% head()

# You can also use let() to remove variables and
# modify existing variables
mtcars %>%
  let(
    mpg = NULL,
    disp = disp * 0.0163871 # convert to litres
  ) %>% head()

# window functions are useful for grouped computations
mtcars %>%
  let(rank = rank(-mpg, ties.method = "min"),
    by = cyl) %>%
  head()

# You can drop variables by setting them to NULL
mtcars %>% let(cyl = NULL) %>% head()

# keeps all existing variables
mtcars %>%
  let(displ_l = disp / 61.0237) %>%
  head()

# keeps only the variables you create
mtcars %>%
  take(displ_l = disp / 61.0237)

# can refer to both contextual variables and variable names:
var = 100
mtcars %>%
  let(cyl = cyl * var) %>%
  head()

# filter by condition
mtcars %>%
  take_if(am==0)

# filter by compound condition
mtcars %>%
  take_if(am==0 & mpg>mean(mpg))
```

```
# A 'take' with summary functions applied without 'by' argument returns an aggregated data
mtcars %>%
  take(mean = mean(displ), n = .N)

# Usually, you'll want to group first
mtcars %>%
  take(mean = mean(displ), n = .N, by = cyl)

# You can group by expressions:
mtcars %>%
  take_all(mean, by = list(vsam = vs + am))

# modify all non-grouping variables in-place
mtcars %>%
  let_all((.x - mean(.x))/sd(.x), by = am) %>%
  head()

# modify all non-grouping variables to new variables
mtcars %>%
  let_all(scaled = (.x - mean(.x))/sd(.x), by = am) %>%
  head()

# conditionally modify all variables
iris %>%
  let_all(mean = if(is.numeric(.x)) mean(.x)) %>%
  head()

# modify all variables conditionally on name
iris %>%
  let_all(
    mean = if(startsWith(.name, "Sepal")) mean(.x),
    median = if(startsWith(.name, "Petal")) median(.x),
    by = Species
  ) %>%
  head()

# aggregation with 'take_all'
mtcars %>%
  take_all(mean = mean(.x), sd = sd(.x), n = .N, by = am)

# conditionally aggregate all variables
iris %>%
  take_all(mean = if(is.numeric(.x)) mean(.x))

# aggregate all variables conditionally on name
iris %>%
  take_all(
    mean = if(startsWith(.name, "Sepal")) mean(.x),
    median = if(startsWith(.name, "Petal")) median(.x),
    by = Species
  )
```

```

# parametric evaluation:
var = quote(mean(cyl))
mtcars %>%
  let(mean_cyl = eval(var)) %>%
  head()
take(mtcars, eval(var))

# all together
new_var = "mean_cyl"
mtcars %>%
  let((new_var) := eval(var)) %>%
  head()
take(mtcars, (new_var) := eval(var))

```

---

query\_if

*One-to-one interface for data.table 'f' method*


---

## Description

Quote from [data.table](#):

```

query(data, j, by) # + extra arguments
  |  |
  |  -----> grouped by what?
  -----> what to do?

```

or,

```

query_if(data, i, j, by) # + extra arguments
  |  |  |
  |  |  -----> grouped by what?
  |  -----> what to do?
  ---> on which rows?

```

If you don't need 'i' argument, use 'query'. In this case you can avoid printing leading comma inside brackets to denote empty 'i'.

## Usage

```

query_if(
  data,
  i,
  j,
  by,
  keyby,
  with = TRUE,

```

```

nomatch = getOption("datatable.nomatch"),
mult = "all",
roll = FALSE,
rollends = if (roll == "nearest") c(TRUE, TRUE) else if (roll >= 0) c(FALSE, TRUE)
  else c(TRUE, FALSE),
which = FALSE,
.SDcols,
verbose = getOption("datatable.verbose"),
allow.cartesian = getOption("datatable.allow.cartesian"),
drop = NULL,
on = NULL
)

query(
  data,
  j,
  by,
  keyby,
  with = TRUE,
  nomatch = getOption("datatable.nomatch"),
  mult = "all",
  roll = FALSE,
  rollends = if (roll == "nearest") c(TRUE, TRUE) else if (roll >= 0) c(FALSE, TRUE)
    else c(TRUE, FALSE),
  which = FALSE,
  .SDcols,
  verbose = getOption("datatable.verbose"),
  allow.cartesian = getOption("datatable.allow.cartesian"),
  drop = NULL,
  on = NULL
)

```

## Arguments

- |      |  |
|------|--|
| data | data.table/data.frame data.frame will be automatically converted to data.table.  |
| i    | Integer, logical or character vector, single column numeric matrix, expression of column names, list, data.frame or data.table. integer and logical vectors work the same way they do in [.data.frame except logical NAs are treated as FALSE. expression is evaluated within the frame of the data.table (i.e. it sees column names as if they are variables) and can evaluate to any of the other types. For details see <a href="#">data.table</a>  |
| j    | When with=TRUE (default), j is evaluated within the frame of the data.table; i.e., it sees column names as if they are variables. This allows to not just select columns in j, but also compute on them e.g., x[, a] and x[, sum(a)] returns x\$a and sum(x\$a) as a vector respectively. x[, .(a,b)] and x[, .(sa=sum(a), sb=sum(b))] returns a two column data.table each, the first simply selecting columns a, b and the second computing their sums. For details see <a href="#">data.table</a> . |

by	unquoted name of grouping variable or list of unquoted names of grouping variables. For details see <a href="#">data.table</a>
keyby	Same as by, but with an additional <code>setkey()</code> run on the by columns of the result, for convenience. It is common practice to use 'keyby=' routinely when you wish the result to be sorted. For details see <a href="#">data.table</a>
with	logical. For details see <a href="#">data.table</a> .
nomatch	Same as nomatch in match. For details see <a href="#">data.table</a> .
mult	For details see <a href="#">data.table</a> .
roll	For details see <a href="#">data.table</a> .
rollends	For details see <a href="#">data.table</a> .
which	For details see <a href="#">data.table</a> .
.SDcols	Specifies the columns of x to be included in the special symbol .SD which stands for Subset of data.table. May be character column names or numeric positions. For details see <a href="#">data.table</a> .
verbose	logical. For details see <a href="#">data.table</a> .
allow.cartesian	For details see <a href="#">data.table</a> .
drop	For details see <a href="#">data.table</a> .
on	For details see <a href="#">data.table</a> .

## Value

It depends. For details see [data.table](#).

## Examples

```
# examples from data.table
dat = data.table(x=rep(c("b","a","c"),each=3), y=c(1,3,6), v=1:9)
dat
# basic row subset operations
query_if(dat, 2)                # 2nd row
query_if(dat, 3:2)              # 3rd and 2nd row
query_if(dat, order(x))        # no need for order(dat$x)
query_if(dat, y>2)              # all rows where dat$y > 2
query_if(dat, y>2 & v>5)       # compound logical expressions
query_if(dat, !2:4)            # all rows other than 2:4
query_if(dat, -(2:4))          # same

# select|compute columns data.table way
query(dat, v)                   # v column (as vector)
query(dat, list(v))             # v column (as data.table)
query(dat, sum(v))              # sum of column v, returned as vector
query(dat, list(sum(v)))        # same, but return data.table (column autonamed V1)
query(dat, list(v, v*2))        # return two column data.table, v and v*2

# subset rows and select|compute data.table way
query_if(dat, 2:3, sum(v))      # sum(v) over rows 2 and 3, return vector
```

```

query_if(dat, 2:3, list(sum(v)))           # same, but return data.table with column V1
query_if(dat, 2:3, list(sv=sum(v)))       # same, but return data.table with column sv
query_if(dat, 2:5, cat(v, "\n"))          # just for j's side effect

# select columns the data.frame way
query(dat, 2, with=FALSE)                  # 2nd column, returns a data.table always
colNum = 2
query(dat, colNum, with=FALSE)             # same, equivalent to DT[, .SD, .SDcols=colNum]

# grouping operations - j and by
query(dat, sum(v), by=x)                   # ad hoc by, order of groups preserved in result
query(dat, sum(v), keyby=x)                # same, but order the result on by cols
query(dat, sum(v), by=x) %>%
  query_if(order(x))                       # same but by chaining expressions together

# fast ad hoc row subsets (subsets as joins)
# same as x == "a" but uses binary search (fast)
query_if(dat, "a", on="x")
# same, for convenience, no need to quote every column
query_if(dat, "a", on=list(x))
query_if(dat, .("a"), on="x")              # same
# same, single "==" internally optimised to use binary search (fast)
query_if(dat, x=="a")
# not yet optimized, currently vector scan subset
query_if(dat, x!="b" | y!=3)
# join on columns x,y of 'dat'; uses binary search (fast)
query_if(dat, .("b", 3), on=c("x", "y"))
query_if(dat, .("b", 3), on=list(x, y))    # same, but using on=list()
query_if(dat, .("b", 1:2), on=c("x", "y")) # no match returns NA
query_if(dat, .("b", 1:2), on=(x, y), nomatch=0) # no match row is not returned
# locf, nomatch row gets rolled by previous row
query_if(dat, .("b", 1:2), on=c("x", "y"), roll=Inf)
query_if(dat, .("b", 1:2), on=(x, y), roll=-Inf) # nocb, nomatch row gets rolled by next row
# on rows where dat$x=="b", calculate sum(v*y)
query_if(dat, "b", sum(v*y), on="x")

# all together now
query_if(dat, x!="a", sum(v), by=x)         # get sum(v) by "x" for each i != "a"
query_if(dat, !"a", sum(v), by=.EACHI, on="x") # same, but using subsets-as-joins
query_if(dat, c("b", "c"), sum(v), by=.EACHI, on="x") # same
query_if(dat, c("b", "c"), sum(v), by=.EACHI, on=(x)) # same, using on=(.)

# joins as subsets
X = data.table(x=c("c", "b"), v=8:7, foo=c(4,2))
X

query_if(dat, X, on="x")                    # right join
query_if(X, dat, on="x")                   # left join
query_if(dat, X, on="x", nomatch=0)        # inner join
query_if(dat, !X, on="x")                  # not join
# join using column "y" of 'dat' with column "v" of X
query_if(dat, X, on=c(y="v"))
query_if(dat, X, on="y=v")                 # same as above (v1.9.8+)

```

```

query_if(dat, X, on = .(y<=foo))           # NEW non-equi join (v1.9.8+)
query_if(dat, X, on="y<=foo")             # same as above
query_if(dat, X, on=c("y<=foo"))         # same as above
query_if(dat, X, on=.(y>=foo))           # NEW non-equi join (v1.9.8+)
query_if(dat, X, on=.(x, y<=foo))         # NEW non-equi join (v1.9.8+)
query_if(dat, X, .(x,y,x.y,v), on=.(x, y>=foo)) # Select x's join columns as well

query_if(dat, X, on="x", mult="first")    # first row of each group
query_if(dat, X, on="x", mult="last")     # last row of each group
query_if(dat, X, sum(v), by=.EACHI, on="x") # join and eval j for each row in i
query_if(dat, X, sum(v)*foo, by=.EACHI, on="x") # join inherited scope
query_if(dat, X, sum(v)*i.v, by=.EACHI, on="x") # 'i,v' refers to X's v column
query_if(dat, X, on=.(x, v>=v), sum(y)*foo, by=.EACHI) # NEW non-equi join with by=.EACHI (v1.9.8+)

# more on special symbols, see also ?"special-symbols"
query_if(dat, .N)                         # last row
query(dat, .N)                            # total number of rows in DT
query(dat, .N, by=x)                      # number of rows in each group
query(dat, .SD, .SDcols=x:y)              # select columns 'x' and 'y'
query(dat, .SD[1])                        # first row of all columns
query(dat, .SD[1], by=x)                  # first row of 'y' and 'v' for each group in 'x'
query(dat, c(.N, lapply(.SD, sum)), by=x) # get rows *and* sum columns 'v' and 'y' by group
query(dat, .I[1], by=x)                  # row number in DT corresponding to each group
query(dat, grp := .GRP, by=x) %>% head()  # add a group counter column
query(X, query_if(dat, .BY, y, on="x"), by=x) # join within each group

# add/update/delete by reference (see ?assign)
query(dat, z:=42L) %>% head()             # add new column by reference
query(dat, z:=NULL) %>% head()           # remove column by reference
query_if(dat, "a", v:=42L, on="x") %>% head() # subassign to existing v column by reference
query_if(dat, "b", v2:=84L, on="x") %>% head() # subassign to new column by reference (NA padded)

# NB: postfix [] is shortcut to print()
query(dat, m:=mean(v), by=x)[]           # add new column by reference by group

# advanced usage
dat = data.table(x=rep(c("b","a","c"),each=3),
                 v=c(1,1,1,2,2,1,1,2,2),
                 y=c(1,3,6),
                 a=1:9,
                 b=9:1)

dat
query(dat, sum(v), by=.(y%%2))           # expressions in by
query(dat, sum(v), by=.(bool = y%%2))   # same, using a named list to change by column name
query(dat, .SD[2], by=x)                 # get 2nd row of each group
query(dat, tail(.SD,2), by=x)            # last 2 rows of each group
query(dat, lapply(.SD, sum), by=x)       # sum of all (other) columns for each group
query(dat, .SD[which.min(v)], by=x)      # nested query by group

query(dat, list(MySum=sum(v),
                MyMin=min(v)),

```



```

        MyMax=max(v)),
    by=(x, y%%2)
) # by 2 expressions

query(dat, .(a = .(a), b = .(b)), by=x) # list columns
query(dat, .(seq = min(a):max(b)), by=x) # j is not limited to just aggregations
query(dat, sum(v), by=x) %>%
  query_if(V1<20) # compound query
query(dat, sum(v), by=x) %>%
  setorder(-V1) %>%
  head() # ordering results
query(dat, c(.N, lapply(.SD,sum)), by=x) # get number of observations and sum per group

# anonymous lambda in 'j', j accepts any valid
# expression. TO REMEMBER: every element of
# the list becomes a column in result.
query(dat,
  {tmp = mean(y);
  .(a = a-tmp, b = b-tmp)
  },
  by=x)

## Not run:
pdf("new.pdf")
query(dat, plot(a,b), by=x) # can also plot in 'j'
dev.off()

## End(Not run)
# using rleid, get max(y) and min of all cols in .SDcols for each consecutive run of 'v'
query(dat,
  c(.(y=max(y)), lapply(.SD, min)),
  by=rleid(v),
  .SDcols=v:b
)

```

---

to\_list

*Apply an expression to each element of a list or vector*


---

### Description

- to\_list always returns a list, each element of which is the result of expression expr on the elements of data. By default, NULL's will be removed from the result. You can change this behavior with skip\_null argument.
- to\_vec is the same as to\_list but tries to convert its result to vector via [unlist](#).
- to\_df and to\_dfr try to combine its results to data.table by rows.
- to\_dfc tries to combine its result to data.table by columns.

Expression can use predefined variables: '.x' is a value of current list element, '.name' is a name of the element and '.index' is sequential number of the element. '.value' is an alias to '.x'.

**Usage**

```
to_list(  
  data,  
  expr = NULL,  
  ...,  
  skip_null = TRUE,  
  trace = FALSE,  
  trace_step = 1L  
)
```

```
to_vec(  
  data,  
  expr = NULL,  
  ...,  
  skip_null = TRUE,  
  trace = FALSE,  
  trace_step = 1L,  
  recursive = TRUE,  
  use.names = TRUE  
)
```

```
to_df(  
  data,  
  expr = NULL,  
  ...,  
  trace = FALSE,  
  trace_step = 1L,  
  idvalue = NULL,  
  idname = "item_id"  
)
```

```
to_dfr(  
  data,  
  expr = NULL,  
  ...,  
  trace = FALSE,  
  trace_step = 1L,  
  idvalue = NULL,  
  idname = "item_id"  
)
```

```
to_dfc(data, expr = NULL, ..., trace = FALSE, trace_step = 1)
```

**Arguments**

data	data.frame/list/vector
expr	expression or function. Expression can use predefined variables: '.x' is a value of current list element, '.name' is a name of the element and '.index' is sequen-

	tial number of the element.
...	further arguments provided if 'expr' is function.
skip_null	logical Should we skip NULL's from result? Default is TRUE
trace	FALSE by default. Should we report progress during execution? Possible values are TRUE, FALSE, "pb" (progress bar) or custom expression in 'quote', e.g. 'quote(print(.x))'. Expression can contain '.x', '.name', '.res' and '.index' variables.
trace_step	integer. 1 by default. Step for reporting progress. Ignored if 'trace' argument is equal to FALSE.
recursive	logical. Should unlisting be applied to list components of x? For details see <a href="#">unlist</a> .
use.names	logical. TRUE by default. Should names of source list be preserved? Setting it to FALSE in some cases can greatly increase performance. For details see <a href="#">unlist</a> .
idvalue	expression for calculation id column. Usually it is just unquoted symbols: one of the '.name', '.index' or '.x'.
idname	character, 'item_id' by default. Name for the id column.

### Value

'to\_list' returns list, 'to\_vec' tries to return vector and other functions return data.table

### Examples

```
1:5 %>%
  to_list(rnorm(n = 3, .x))

# or in 'lapply' style
1:5 %>%
  to_list(rnorm, n = 3) %>%
  to_vec(mean)

# or use an anonymous function
1:5 %>%
  to_list(function(x) rnorm(3, x))

# Use to_vec() to reduce output to a vector instead
# of a list:
# filtering - return only even numbers
to_vec(1:10, if(.x %% 2 == 0) .x)

# filtering - mean only on the numeric columns
to_vec(iris, if(is.numeric(.x)) mean(.x))

# mean for numerics, number of distincts for others
to_vec(iris, if(is.numeric(.x)) mean(.x) else uniqueN(.x))

# means for Sepal
to_vec(iris, if(startsWith(.name, "Sepal")) mean(.x))
```

```

# A more realistic example: split a data frame into pieces, fit a
# model to each piece, summarise and extract R^2
mtcars %>%
  split(.$cyl) %>%
  to_list(summary(lm(mpg ~ wt, data = .x))) %>%
  to_vec(.$r.squared)

# If each element of the output is a data frame, use
# to_df to row-bind them together:
mtcars %>%
  split(.$cyl) %>%
  to_list(lm(mpg ~ wt, data = .x)) %>%
  to_df(c(cyl = .name, coef(.x)))

## Not run:
# read all csv files in "data" to data.frame
all_files = dir("data", pattern = "csv$", full.names = TRUE) %>%
  to_df(fread,
        idvalue = basename(.x),
        idname = "filename",
        trace = "pb"
        )

## End(Not run)

```

---

vlookup

*Look up values in dictionary.*


---

## Description

vlookup function is inspired by VLOOKUP spreadsheet function. It looks for a `lookup_value` in the `lookup_column` of the `dict`, and then returns values in the same rows from `result_column`. `xlookup` is simplified version of `vlookup`. It searches for a `lookup_value` in the `lookup_vector` and return values in the same position from the `result_vector`.

## Usage

```

vlookup(
  lookup_value,
  dict,
  result_column = 2,
  lookup_column = 1,
  no_match = NA
)

```

```

xlookup(lookup_value, lookup_vector, result_vector, no_match = NA)

```

**Arguments**

lookup_value	Vector of looked up values
dict	data.frame. Dictionary.
result_column	numeric or character. Resulting columns in the dict. Default value for result_column is 2 - for frequent case of dictionary with keys in the first column and results in the second column.
lookup_column	Column of dict in which lookup value will be searched. By default, it is the first column of the dict.
no_match	vector of length one. NA by default. Where a valid match is not found, return the 'no_match' value you supply.
lookup_vector	vector in which 'lookup_value' will be searched during 'xlookup'.
result_vector	vector with resulting values for 'xlookup'.

**Value**

xlookup always return vector, vlookup returns vector if the result\_column is single value. In the opposite case data.frame will be returned.

**Examples**

```
# with data.frame
dict = data.frame(num=1:26, small=letters, cap=LETTERS)
vlookup(1:3, dict)
vlookup(c(45,1:3,58), dict, result_column='cap')
vlookup(c(45,1:3,58), dict, result_column='cap', no_match = "Not found")

# the same with xlookup
xlookup(1:3, dict$num, dict$small)
xlookup(c(45,1:3,58), dict$num, dict$cap)
xlookup(c(45,1:3,58), dict$num, dict$cap, no_match = "Not found")

# example from base 'merge'
authors = data.table(
  surname = c("Tukey", "Venables", "Tierney", "Ripley", "McNeil"),
  nationality = c("US", "Australia", "US", "UK", "Australia"),
  deceased = c("yes", rep("no", 4))
)

books = data.table(
  surname = c("Tukey", "Venables", "Tierney",
             "Ripley", "Ripley", "McNeil", "R Core"),
  title = c("Exploratory Data Analysis",
            "Modern Applied Statistics ...",
            "LISP-STAT",
            "Spatial Statistics", "Stochastic Simulation",
            "Interactive Data Analysis",
            "An Introduction to R")
)
```

```

let(books,
  c("author_nationality", "author_deceased") := vlookup(surname,
    dict = authors,
    result_column = 2:3
  )
)[]

# Just for fun. Examples borrowed from Microsoft Excel.
# It is not the R way of doing things.

# Example 2

ex2 = fread("
  Item_ID Item Cost Markup
  ST-340 Stroller 145.67 0.30
  BI-567 Bib 3.56 0.40
  DI-328 Diapers 21.45 0.35
  WI-989 Wipes 5.12 0.40
  AS-469 Aspirator 2.56 0.45
")

# Calculates the retail price of diapers by adding the markup percentage to the cost.
vlookup("DI-328", ex2, 3) * (1 + vlookup("DI-328", ex2, 4)) # 28.9575

# Calculates the sale price of wipes by subtracting a specified discount from
# the retail price.
(vlookup("WI-989", ex2, "Cost") * (1 + vlookup("WI-989", ex2, "Markup"))) * (1 - 0.2) # 5.7344

A2 = ex2[["Item_ID"]][1]
A3 = ex2[["Item_ID"]][2]

# If the cost of an item is greater than or equal to $20.00, displays the string
# "Markup is nn%"; otherwise, displays the string "Cost is under $20.00".
ifelse(vlookup(A2, ex2, "Cost") >= 20,
  paste0("Markup is ", 100 * vlookup(A2, ex2, "Markup"), "%"),
  "Cost is under $20.00") # Markup is 30%

# If the cost of an item is greater than or equal to $20.00, displays the string
# Markup is nn%"; otherwise, displays the string "Cost is $n.nn".
ifelse(vlookup(A3, ex2, "Cost") >= 20,
  paste0("Markup is: ", 100 * vlookup(A3, ex2, "Markup") , "%"),
  paste0("Cost is $", vlookup(A3, ex2, "Cost"))) #Cost is $3.56

# Example 3

ex3 = fread('
  ID Last_name First_name Title Birth_date
  1 Davis Sara "Sales Rep." 12/8/1968
  2 Fontana Olivier "V.P. of Sales" 2/19/1952
  3 Leal Karina "Sales Rep." 8/30/1963
')
```

```
4 Patten Michael "Sales Rep." 9/19/1958
5 Burke Brian "Sales Mgr." 3/4/1955
6 Sousa Luis "Sales Rep." 7/2/1963
')

# If there is an employee with an ID of 5, displays the employee's last name;
# otherwise, displays the message "Employee not found".
vlookup(5, ex3, "Last_name", no_match = "Employee not found") # Burke

# Many employees
vlookup(1:10, ex3, "Last_name", no_match = "Employee not found")

# For the employee with an ID of 4, concatenates the values of three cells into
# a complete sentence.
paste0(vlookup(4, ex3, "First_name"), " ",
       vlookup(4, ex3, "Last_name"), " is a ",
       vlookup(4, ex3, "Title")) # Michael Patten is a Sales Rep.
```

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