# Package 'dplyr'

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**Type** Package

**Version** 0.4.3

**Title** A Grammar of Data Manipulation

**Description** A fast, consistent tool for working with data frame like objects, both in memory and out of memory.

**URL** https://github.com/hadley/dplyr

**BugReports** https://github.com/hadley/dplyr/issues

**Depends** R (>= 3.1.2)

**Imports** assertthat, utils, R6, Rcpp, magrittr, lazyeval (>= 0.1.10), DBI (>= 0.3)

**Suggests** RSQLite (>= 1.0.0), RMySQL, RPostgreSQL, data.table, testthat, knitr, microbenchmark, ggplot2, mgcv, Lahman (>= 3.0-1), nycflights13, methods

**VignetteBuilder** knitr

**LazyData** yes

**LinkingTo** Rcpp (>= 0.12.0), BH (>= 1.58.0-1)

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**Collate** 'RcppExports.R' 'all-equal.r' 'bench-compare.r' 'chain.r' 'cluster.R' 'colwise.R' 'compute-collect.r' 'copy-to.r' 'data-lahman.r' 'data-nasa.r' 'data-nycflights13.r' 'data-temp.r' 'data.r' 'dataframe.R' 'dbi-s3.r' 'desc.r' 'distinct.R' 'do.r' 'dplyr.r' 'explain.r' 'failwith.r' 'frame-data.R' 'funs.R' 'glimpse.R' 'group-by.r' 'group-indices.R' 'group-size.r' 'grouped-df.r' 'grouped-dt.r' 'id.r' 'inline.r' 'join.r' 'lead-lag.R' 'location.R' 'manip.r' 'nth-value.R' 'order-by.R' 'over.R' 'partial-eval.r' 'progress.R' 'query.r' 'rank.R' 'rbind.r' 'rowwise.r' 'sample.R' 'select-utils.R' 'select-vars.R' 'sets.r' 'sql-escape.r' 'sql-star.r' 'src-local.r' 'src-mysql.r' 'src-postgres.r' 'src-sql.r' 'src-sqlite.r' 'src.r' 'tally.R' 'tbl-cube.r' 'tbl-df.r' 'tbl-dt.r' 'tbl-sql.r' 'tbl.r'

'top-n.R' 'translate-sql-helpers.r' 'translate-sql-base.r'
'translate-sql-window.r' 'translate-sql.r' 'type-sum.r'
'utils-dt.R' 'utils-format.r' 'utils.r' 'view.r' 'zzz.r'

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**Author** Hadley Wickham [aut, cre],
Romain Francois [aut],
RStudio [cph]

**Maintainer** Hadley Wickham <hadley@rstudio.com>

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# R topics documented:

---

`add_rownames` *Convert row names to an explicit variable.*

---

## Description

Convert row names to an explicit variable.

## Usage

```
add_rownames(df, var = "rowname")
```

## Arguments

| | |
|---|---|
| `df` | Input data frame with rownames. |
| `var` | Name of variable to use |

## Examples

```
mtcars %>% tbl_df()

mtcars %>% add_rownames()
```

---

`all.equal.tbl_df` *Provide a useful implementation of all.equal for data.frames.*

---

## Description

Provide a useful implementation of all.equal for data.frames.

## Usage

```
## S3 method for class 'tbl_df'
all.equal(target, current, ignore_col_order = TRUE,
  ignore_row_order = TRUE, convert = FALSE, ...)

## S3 method for class 'tbl_dt'
all.equal(target, current, ignore_col_order = TRUE,
  ignore_row_order = TRUE, convert = FALSE, ...)
```

## Arguments

| | |
|---|---|
| `target,current` | two data frames to compare |
| `ignore_col_order` | should order of columns be ignored? |
| `ignore_row_order` | should order of rows be ignored? |
| `convert` | Should similar classes be converted? Currently this will convert factor to character and integer to double. |
| `...` | Ignored. Needed for compatibility with the generic. |

## Value

`TRUE` if equal, otherwise a character vector describing the first reason why they're not equal. Use `isTRUE` if using the result in an `if` expression.

## Examples

```
scramble <- function(x) x[sample(nrow(x)), sample(ncol(x))]

# By default, ordering of rows and columns ignored
mtcars_df <- tbl_df(mtcars)
all.equal(mtcars_df, scramble(mtcars_df))

# But those can be overriden if desired
all.equal(mtcars_df, scramble(mtcars_df), ignore_col_order = FALSE)
all.equal(mtcars_df, scramble(mtcars_df), ignore_row_order = FALSE)
```

`arrange` *Arrange rows by variables.*

## Description

Use `desc` to sort a variable in descending order.

## Usage

```
arrange(.data, ...)

arrange_(.data, ..., .dots)
```

## Arguments

| | |
|---|---|
| `.data` | A tbl. All main verbs are S3 generics and provide methods for `tbl_df`, `tbl_dt` and `tbl_sql`. |
| `...` | Comma separated list of unquoted variable names. Use `desc` to sort a variable in descending order. |
| `.dots` | Used to work around non-standard evaluation. See `vignette("nse")` for details. |

## Value

An object of the same class as `.data`.

Data frame row names are silently dropped. To preserve, convert to an explicit variable.

## Locales

Note that for local data frames, the ordering is done in C++ code which does not have access to the local specific ordering usually done in R. This means that strings are ordered as if in the C locale.

## See Also

Other single.table.verbs: `filter`, `filter_`; `mutate`, `mutate_`, `transmute`, `transmute_`; `rename`, `rename_`, `select`, `select_`; `slice`, `slice_`; `summarise`, `summarise_`, `summarize`, `summarize_`

## Examples

```
arrange(mtcars, cyl, disp)
arrange(mtcars, desc(disp))
```

## as.tbl_cube — *Coerce an existing data structure into a* tbl_cube

### Description

Coerce an existing data structure into a tbl_cube

### Usage

```
as.tbl_cube(x, ...)

## S3 method for class 'array'
as.tbl_cube(x, met_name = deparse(substitute(x)),
  dim_names = names(dimnames(x)), ...)

## S3 method for class 'table'
as.tbl_cube(x, met_name = deparse(substitute(x)),
  dim_names = names(dimnames(x)), ...)

## S3 method for class 'matrix'
as.tbl_cube(x, met_name = deparse(substitute(x)),
  dim_names = names(dimnames(x)), ...)

## S3 method for class 'data.frame'
as.tbl_cube(x, dim_names, ...)
```

### Arguments

| | |
|---|---|
| x | an object to convert. Built in methods will convert arrays, tables and data frames. |
| ... | Passed on to individual methods; otherwise ignored. |
| met_name | a string to use as the name for the metric |
| dim_names | names of the dimesions. Defaults to the names of the dimnames. |

## as_data_frame — *Coerce a list to a data frame.*

### Description

as.data.frame is effectively a thin wrapper around data.frame, and hence is rather slow (because it calls data.frame on each element before cbinding together). as_data_frame just verifies that the list is structured correctly (i.e. named, and each element is same length) then sets class and row name attributes.

## Usage

```
as_data_frame(x)
```

## Arguments

| | |
|---|---|
| x | A list. Each element of the list must have the same length. |

## Examples

```
l <- list(x = 1:500, y = runif(500), z = 500:1)
df <- as_data_frame(l)

# Coercing to a data frame does not copy columns
changes(as_data_frame(l), as_data_frame(l))

# as_data_frame is considerably simpler/faster than as.data.frame
# making it more suitable for use when you have things that are
# lists
## Not run:
l2 <- replicate(26, sample(letters), simplify = FALSE)
names(l2) <- letters
microbenchmark::microbenchmark(
  as_data_frame(l2),
  as.data.frame(l2)
)

## End(Not run)
```

---

# bench_compare *Evaluate, compare, benchmark operations of a set of srcs.*

---

## Description

These functions support the comparison of results and timings across multiple sources.

## Usage

```
bench_tbls(tbls, op, ..., times = 10)

compare_tbls(tbls, op, ref = NULL, compare = equal_data_frame, ...)

eval_tbls(tbls, op)
```

## Arguments

| | |
|---|---|
| tbls | A list of tbls. |
| op | A function with a single argument, called often with each element of tbls. |
| times | For benchmarking, the number of times each operation is repeated. |

| | |
|---|---|
| ref | For checking, an data frame to test results against. If not supplied, defaults to the results from the first src. |
| compare | A function used to compare the results. Defaults to equal_data_frame which ignores the order of rows and columns. |
| ... | For compare_tbls: additional parameters passed on the compare function<br>For bench_tbls: additional benchmarks to run. |

## Value

eval_tbls: a list of data frames.

compare_tbls: an invisible TRUE on success, otherwise an error is thrown.

bench_tbls: an object of class microbenchmark

## See Also

src_local for working with local data

## Examples

```
## Not run:
if (require("microbenchmark") && has_lahman()) {
lahman_local <- lahman_srcs("df", "dt")
teams <- lapply(lahman_local, function(x) x %>% tbl("Teams"))

compare_tbls(teams, function(x) x %>% filter(yearID == 2010))
bench_tbls(teams, function(x) x %>% filter(yearID == 2010))

# You can also supply arbitrary additional arguments to bench_tbls
# if there are other operations you'd like to compare.
bench_tbls(teams, function(x) x %>% filter(yearID == 2010),
   base = subset(Lahman::Teams, yearID == 2010))

# A more complicated example using multiple tables
setup <- function(src) {
  list(
    src %>% tbl("Batting") %>% filter(stint == 1) %>% select(playerID:H),
    src %>% tbl("Master") %>% select(playerID, birthYear)
  )
}
two_tables <- lapply(lahman_local, setup)

op <- function(tbls) {
  semi_join(tbls[[1]], tbls[[2]], by = "playerID")
}
# compare_tbls(two_tables, op)
bench_tbls(two_tables, op, times = 2)

}

## End(Not run)
```

## between *Do values in a numeric vector fall in specified range?*

### Description

This is a shortcut for `x >= left & x <= right`, implemented efficiently in C++ for local values, and translated to the appropriate SQL for remote tables.

### Usage

```
between(x, left, right)
```

### Arguments

| | |
|---|---|
| `x` | A numeric vector of values |
| `left,right` | Boundary values |

### Examples

```
x <- rnorm(1e2)
x[between(x, -1, 1)]
```

## bind *Efficiently bind multiple data frames by row and column.*

### Description

This is an efficient implementation of the common pattern of `do.call(rbind, dfs)` or `do.call(cbind, dfs)` for binding many data frames into one. `combine()` acts like `c()` or `unlist()` but uses consistent dplyr coercion rules.

### Usage

```
bind_rows(..., .id = NULL)

bind_cols(x, ...)

combine(x, ...)
```

## Arguments

| | |
|---|---|
| `.id` | Data frames identifier.<br>When `.id` is supplied, a new column of identifiers is created to link each row to its original data frame. The labels are taken from the named arguments to `bind_rows()`. When a list of data frames is supplied, the labels are taken from the names of the list. If no names are found a numeric sequence is used instead. |
| `x,...` | Data frames to combine.<br>You can either supply one data frame per argument, or a list of data frames in the first argument.<br>When column-binding, rows are matched by position, not value so all data frames must have the same number of rows. To match by value, not position, see `left_join` etc. When row-binding, columns are matched by name, and any values that don't match will be filled with NA. |

## Value

`bind_rows` and `bind_cols` always return a `tbl_df`

## Deprecated functions

`rbind_list` and `rbind_all` have been deprecated. Instead use `bind_rows`.

## Examples

```
one <- mtcars[1:4, ]
two <- mtcars[11:14, ]

# You can either supply data frames as arguments
bind_rows(one, two)
# Or a single argument containing a list of data frames
bind_rows(list(one, two))
bind_rows(split(mtcars, mtcars$cyl))

# When you supply a column name with the `.id` argument, a new
# column is created to link each row to its original data frame
bind_rows(list(one, two), .id = "id")
bind_rows(list(a = one, b = two), .id = "id")
bind_rows("group 1" = one, "group 2" = two, .id = "groups")

# Columns don't need to match when row-binding
bind_rows(data.frame(x = 1:3), data.frame(y = 1:4))
## Not run:
# Rows do need to match when column-binding
bind_cols(data.frame(x = 1), data.frame(y = 1:2))

## End(Not run)

bind_cols(one, two)
bind_cols(list(one, two))
```

```
# combine applies the same coercion rules
f1 <- factor("a")
f2 <- factor("b")
c(f1, f2)
unlist(list(f1, f2))

combine(f1, f2)
combine(list(f1, f2))
```

---

`build_sql` *Build a SQL string.*

---

## Description

This is a convenience function that should prevent sql injection attacks (which in the context of dplyr are most likely to be accidental not deliberate) by automatically escaping all expressions in the input, while treating bare strings as sql. This is unlikely to prevent any serious attack, but should make it unlikely that you produce invalid sql.

## Usage

```
build_sql(..., .env = parent.frame(), con = NULL)
```

## Arguments

| | |
|---|---|
| `...` | input to convert to SQL. Use `sql` to preserve user input as is (dangerous), and `ident` to label user input as sql identifiers (safe) |
| `.env` | the environment in which to evalute the arguments. Should not be needed in typical use. |
| `con` | database connection; used to select correct quoting characters. |

## Examples

```
build_sql("SELECT * FROM TABLE")
x <- "TABLE"
build_sql("SELECT * FROM ", x)
build_sql("SELECT * FROM ", ident(x))
build_sql("SELECT * FROM ", sql(x))

# http://xkcd.com/327/
name <- "Robert'); DROP TABLE Students;--"
build_sql("INSERT INTO Students (Name) VALUES (", name, ")")
```

`chain` *Chain together multiple operations.*

## Description

The downside of the functional nature of dplyr is that when you combine multiple data manipulation operations, you have to read from the inside out and the arguments may be very distant to the function call. These functions providing an alternative way of calling dplyr (and other data manipulation) functions that you read can from left to right.

## Usage

```
chain(..., env = parent.frame())

chain_q(calls, env = parent.frame())

lhs %.% rhs

lhs %>% rhs
```

## Arguments

| | |
|---|---|
| `...,calls` | A sequence of data transformations, starting with a dataset. The first argument of each call should be omitted - the value of the previous step will be substituted in automatically. Use `chain` and `...` when working interactive; use `chain_q` and `calls` when calling from another function. |
| `env` | Environment in which to evaluation expressions. In ordinary operation you should not need to set this parameter. |
| `lhs,rhs` | A dataset and function to apply to it |

## Details

The functions work via simple substitution so that `x %>% f(y)` is translated into `f(x, y)`.

## Deprecation

`chain` was deprecated in version 0.2, and will be removed in 0.3. `%.%` was deprecated in version 0.3, and defunct in 0.4. They wwere removed in the interest of making dplyr code more standardised and `%>%` is much more popular.

## See Also

`%>%` in the magrittr package for a more detailed explanation of the forward pipe semantics.

## Examples

```
# If you're performing many operations you can either do step by step
if (require("nycflights13")) {
a1 <- group_by(flights, year, month, day)
a2 <- select(a1, arr_delay, dep_delay)
a3 <- summarise(a2,
  arr = mean(arr_delay, na.rm = TRUE),
  dep = mean(dep_delay, na.rm = TRUE))
a4 <- filter(a3, arr > 30 | dep > 30)

# If you don't want to save the intermediate results, you need to
# wrap the functions:
filter(
  summarise(
    select(
      group_by(flights, year, month, day),
      arr_delay, dep_delay
    ),
    arr = mean(arr_delay, na.rm = TRUE),
    dep = mean(dep_delay, na.rm = TRUE)
  ),
  arr > 30 | dep > 30
)

# This is difficult to read because the order of the operations is from
# inside to out, and the arguments are a long way away from the function.
# Alternatively you can use %>% to sequence the operations
# linearly:

flights %>%
  group_by(year, month, day) %>%
  select(arr_delay, dep_delay) %>%
  summarise(
    arr = mean(arr_delay, na.rm = TRUE),
    dep = mean(dep_delay, na.rm = TRUE)
  ) %>%
  filter(arr > 30 | dep > 30)
}
```

---

`compute` *Compute a lazy tbl.*

---

## Description

`compute` forces computation of lazy tbls, leaving data in the remote source. `collect` also forces computation, but will bring data back into an R data.frame (stored in a `tbl_df`). `collapse` doesn't force computation, but collapses a complex tbl into a form that additional restrictions can be placed on.

## Usage

```
compute(x, name = random_table_name(), ...)

collect(x, ...)

collapse(x, ...)

## S3 method for class 'tbl_sql'
compute(x, name = random_table_name(), temporary = TRUE,
  ...)
```

## Arguments

| | |
|---|---|
| x | a data tbl |
| name | name of temporary table on database. |
| ... | other arguments passed on to methods |
| temporary | if TRUE, will create a temporary table that is local to this connection and will be automatically deleted when the connection expires |

## Grouping

compute and collect preserve grouping, collapse drops it.

## See Also

copy_to which is the conceptual opposite: it takes a local data frame and makes it available to the remote source.

## Examples

```
if (require("RSQLite") && has_lahman("sqlite")) {
  batting <- tbl(lahman_sqlite(), "Batting")
  remote <- select(filter(batting, yearID > 2010 && stint == 1), playerID:H)
  remote2 <- collapse(remote)
  cached <- compute(remote)
  local  <- collect(remote)
}
```

---

# copy_to *Copy a local data frame to a remote src.*

---

## Description

This uploads a local data frame into a remote data source, creating the table definition as needed. Wherever possible, the new object will be temporary, limited to the current connection to the source.

### Usage

```
copy_to(dest, df, name = deparse(substitute(df)), ...)
```

### Arguments

| | |
|---|---|
| dest | remote data source |
| df | local data frame |
| name | name for new remote table. |
| ... | other parameters passed to methods. |

### Value

a tbl object in the remote source

---

## copy_to.src_sql *Copy a local data frame to a sqlite src.*

---

### Description

This standard method works for all sql sources.

### Usage

```
## S3 method for class 'src_sql'
copy_to(dest, df, name = deparse(substitute(df)),
  types = NULL, temporary = TRUE, indexes = NULL, analyze = TRUE, ...)
```

### Arguments

| | |
|---|---|
| dest | remote data source |
| df | local data frame |
| name | name for new remote table. |
| types | a character vector giving variable types to use for the columns. See http://www.sqlite.org/datatype3.html for available types. |
| temporary | if TRUE, will create a temporary table that is local to this connection and will be automatically deleted when the connection expires |
| indexes | a list of character vectors. Each element of the list will create a new index. |
| analyze | if TRUE (the default), will automatically ANALYZE the new table so that the query optimiser has useful information. |
| ... | other parameters passed to methods. |

### Value

a sqlite tbl object

## Examples

```
if (requireNamespace("RSQLite")) {
db <- src_sqlite(tempfile(), create = TRUE)

iris2 <- copy_to(db, iris)
mtcars$model <- rownames(mtcars)
mtcars2 <- copy_to(db, mtcars, indexes = list("model"))

explain(filter(mtcars2, model == "Hornet 4 Drive"))

# Note that tables are temporary by default, so they're not
# visible from other connections to the same database.
src_tbls(db)
db2 <- src_sqlite(db$path)
src_tbls(db2)
}
```

---

# cumall *Cumulativate versions of any, all, and mean*

---

## Description

dplyr adds cumall, cumany, and cummean to complete R's set of cumulate functions to match the aggregation functions available in most databases

## Usage

```
cumall(x)

cumany(x)

cummean(x)
```

## Arguments

| | |
|---|---|
| x | For cumall & cumany, a logical vector; for cummean an integer or numeric vector |

---

# data_frame *Build a data frame.*

---

## Description

A trimmed down version of `data.frame` that:

1. Never coerces inputs (i.e. strings stay as strings!).
2. Never adds `row.names`.
3. Never munges column names.
4. Only recycles length 1 inputs.
5. Evaluates its arguments lazily and in order.
6. Adds `tbl_df` class to output.

## Usage

```
data_frame(...)

data_frame_(columns)
```

## Arguments

| | |
|---|---|
| `...` | A set of named arguments |
| `columns` | A `lazy_dots`. |

## See Also

`as_data_frame` to turn an existing list into a data frame.

## Examples

```
a <- 1:5
data_frame(a, b = a * 2)
data_frame(a, b = a * 2, c = 1)
data_frame(x = runif(10), y = x * 2)

# data_frame never coerces its inputs
str(data_frame(letters))
str(data_frame(x = list(diag(1), diag(2))))

# or munges column names
data_frame(`a + b` = 1:5)

# With the SE version, you give it a list of formulas/expressions
data_frame_(list(x = ~1:10, y = quote(x * 2)))
```

## desc *Descending order.*

### Description

Transform a vector into a format that will be sorted in descending order.

### Usage

```
desc(x)
```

### Arguments

| | |
|---|---|
| x | vector to transform |

### Examples

```
desc(1:10)
desc(factor(letters))
first_day <- seq(as.Date("1910/1/1"), as.Date("1920/1/1"), "years")
desc(first_day)
```

## distinct *Select distinct/unique rows.*

### Description

Retain only unique/distinct rows from an input tbl. This is an efficient version of unique. distinct() is best-suited for interactive use, distinct_() for calling from a function.

### Usage

```
distinct(.data, ...)

distinct_(.data, ..., .dots)
```

### Arguments

| | |
|---|---|
| .data | a tbl |
| ... | Variables to use when determining uniqueness. If there are multiple rows for a given combination of inputs, only the first row will be preserved. |
| .dots | Used to work around non-standard evaluation. See vignette("nse") for details. |

## Examples

```
df <- data.frame(
  x = sample(10, 100, rep = TRUE),
  y = sample(10, 100, rep = TRUE)
)
nrow(df)
nrow(distinct(df))
distinct(df, x)
distinct(df, y)

# You can also use distinct on computed variables
distinct(df, diff = abs(x - y))
```

---

do *Do arbitrary operations on a tbl.*

---

## Description

This is a general purpose complement to the specialised manipulation functions `filter`, `select`, `mutate`, `summarise` and `arrange`. You can use do to perform arbitrary computation, returning either a data frame or arbitrary objects which will be stored in a list. This is particularly useful when working with models: you can fit models per group with do and then flexibly extract components with either another do or `summarise`.

## Usage

```
do(.data, ...)

do_(.data, ..., .dots)

## S3 method for class 'tbl_sql'
do_(.data, ..., .dots, .chunk_size = 10000L)
```

## Arguments

| | |
|---|---|
| `.data` | a tbl |
| `...` | Expressions to apply to each group. If named, results will be stored in a new column. If unnamed, should return a data frame. You can use `.` to refer to the current group. You can not mix named and unnamed arguments. |
| `.dots` | Used to work around non-standard evaluation. See `vignette("nse")` for details. |
| `.chunk_size` | The size of each chunk to pull into R. If this number is too big, the process will be slow because R has to allocate and free a lot of memory. If it's too small, it will be slow, because of the overhead of talking to the database. |

## Value

do always returns a data frame. The first columns in the data frame will be the labels, the others will be computed from .... Named arguments become list-columns, with one element for each group; unnamed elements must be data frames and labels will be duplicated accordingly.

Groups are preserved for a single unnamed input. This is different to summarise because do generally does not reduce the complexity of the data, it just expresses it in a special way. For multiple named inputs, the output is grouped by row with rowwise. This allows other verbs to work in an intuitive way.

## Connection to plyr

If you're familiar with plyr, do with named arguments is basically eqvuivalent to dlply, and do with a single unnamed argument is basically equivalent to ldply. However, instead of storing labels in a separate attribute, the result is always a data frame. This means that summarise applied to the result of do can act like ldply.

## Examples

```
by_cyl <- group_by(mtcars, cyl)
do(by_cyl, head(., 2))

models <- by_cyl %>% do(mod = lm(mpg ~ disp, data = .))
models

summarise(models, rsq = summary(mod)$r.squared)
models %>% do(data.frame(coef = coef(.$mod)))
models %>% do(data.frame(
  var = names(coef(.$mod)),
  coef(summary(.$mod)))
)

models <- by_cyl %>% do(
  mod_linear = lm(mpg ~ disp, data = .),
  mod_quad = lm(mpg ~ poly(disp, 2), data = .)
)
models
compare <- models %>% do(aov = anova(.$mod_linear, .$mod_quad))
# compare %>% summarise(p.value = aov$`Pr(>F)`)

if (require("nycflights13")) {
# You can use it to do any arbitrary computation, like fitting a linear
# model. Let's explore how carrier departure delays vary over the time
carriers <- group_by(flights, carrier)
group_size(carriers)

mods <- do(carriers, mod = lm(arr_delay ~ dep_time, data = .))
mods %>% do(as.data.frame(coef(.$mod)))
mods %>% summarise(rsq = summary(mod)$r.squared)

## Not run:
# This longer example shows the progress bar in action
```

```
by_dest <- flights %>% group_by(dest) %>% filter(n() > 100)
library(mgcv)
by_dest %>% do(smooth = gam(arr_delay ~ s(dep_time) + month, data = .))

## End(Not run)
}
```

---

`dplyr` *dplyr: a grammar of data manipulation*

---

## Description

dplyr provides a flexible grammar of data manipulation. It's the next iteration of plyr, focused on tools for working with data frames (hence the *d* in the name).

## Details

It has three main goals:

- Identify the most important data manipulation verbs and make them easy to use from R.
- Provide blazing fast performance for in-memory data by writing key pieces in C++ (using Rcpp)
- Use the same interface to work with data no matter where it's stored, whether in a data frame, a data table or database.

To learn more about dplyr, start with the vignettes: `browseVignettes(package = "dplyr")`

---

`explain` *Explain details of a tbl.*

---

## Description

This is a generic function which gives more details about an object than `print`, and is more focussed on human readable output than `str`.

## Usage

```
explain(x, ...)

show_query(x)
```

## Arguments

| | |
|---|---|
| `x` | An object to explain |
| `...` | Other parameters possibly used by generic |

### Databases

Explaining a `tbl_sql` will run the SQL `EXPLAIN` command which will describe the query plan. This requires a little bit of knowledge about how `EXPLAIN` works for your database, but is very useful for diagnosing performance problems.

### Examples

```
if (require("RSQLite") && has_lahman("sqlite")) {

lahman_s <- lahman_sqlite()
batting <- tbl(lahman_s, "Batting")
batting %>% show_query()
batting %>% explain()

# The batting database has indices on all ID variables:
# SQLite automatically picks the most restrictive index
batting %>% filter(lgID == "NL" & yearID == 2000L) %>% explain()

# OR's will use multiple indexes
batting %>% filter(lgID == "NL" | yearID == 2000) %>% explain()

# Joins will use indexes in both tables
teams <- tbl(lahman_s, "Teams")
batting %>% left_join(teams, c("yearID", "teamID")) %>% explain()
}
```

---

## `failwith` *Fail with specified value.*

---

### Description

Modify a function so that it returns a default value when there is an error.

### Usage

```
failwith(default = NULL, f, quiet = FALSE)
```

### Arguments

| | |
|---|---|
| `default` | default value |
| `f` | function |
| `quiet` | all error messages be suppressed? |

### Value

a function

## See Also

try_default

## Examples

```
f <- function(x) if (x == 1) stop("Error!") else 1
## Not run:
f(1)
f(2)

## End(Not run)

safef <- failwith(NULL, f)
safef(1)
safef(2)
```

---

filter *Return rows with matching conditions.*

---

## Description

Return rows with matching conditions.

## Usage

```
filter(.data, ...)

filter_(.data, ..., .dots)
```

## Arguments

| | |
|---|---|
| .data | A tbl. All main verbs are S3 generics and provide methods for tbl_df, tbl_dt and tbl_sql. |
| ... | Logical predicates. Multiple conditions are combined with &. |
| .dots | Used to work around non-standard evaluation. See vignette("nse") for details. |

## Value

An object of the same class as .data.

Data frame row names are silently dropped. To preserve, convert to an explicit variable.

## See Also

Other single.table.verbs: arrange, arrange_; mutate, mutate_, transmute, transmute_; rename, rename_, select, select_; slice, slice_; summarise, summarise_, summarize, summarize_

## Examples

```
filter(mtcars, cyl == 8)
filter(mtcars, cyl < 6)

# Multiple criteria
filter(mtcars, cyl < 6 & vs == 1)
filter(mtcars, cyl < 6 | vs == 1)

# Multiple arguments are equivalent to and
filter(mtcars, cyl < 6, vs == 1)
```

---

# frame_data *Row-wise data_frame creation*

---

## Description

Create a row-wise data_frame.

## Usage

```
frame_data(...)

tibble(...)
```

## Arguments

| | |
|---|---|
| ... | Arguments specifying the structure of a data_frame. |

## Examples

```
frame_data(
  ~colA, ~colB,
  "a",   1,
  "b",   2
)
```

---

# funs *Create a list of functions calls.*

---

## Description

funs provides a flexible way to generate a named list of functions for input to other functions like summarise_each.

### Usage

```
funs(...)

funs_(dots)
```

### Arguments

| | |
|---|---|
| dots,... | A list of functions specified by:<br>• Their name, "mean"<br>• The function itself, mean<br>• A call to the function with . as a dummy parameter, mean(., na.rm = TRUE) |

### Examples

```
funs(mean, "mean", mean(., na.rm = TRUE))

# Overide default names
funs(m1 = mean, m2 = "mean", m3 = mean(., na.rm = TRUE))

# If you have function names in a vector, use funs_
fs <- c("min", "max")
funs_(fs)
```

---

## glimpse *Get a glimpse of your data.*

---

### Description

This is like a transposed version of print: columns run down the page, and data runs across. This makes it possible to see every column in a data frame. It's a little like str applied to a data frame but it tries to show you as much data as possible. (And it always shows the underlying data, even when applied to a remote data source.)

### Usage

```
glimpse(tbl, width = getOption("width"))
```

### Arguments

| | |
|---|---|
| tbl | A data table |
| width | Width of output: defaults to the width of the console. |

## Examples

```
glimpse(mtcars)


if (require("RSQLite") && has_lahman("sqlite")) {
  batting <- tbl(lahman_sqlite(), "Batting")
  glimpse(batting)
}
```

---

grouped_dt *A grouped data table.*

---

## Description

The easiest way to create a grouped data table is to call the group_by method on a data table or tbl: this will take care of capturing the unevalated expressions for you.

## Usage

```
grouped_dt(data, vars, copy = TRUE)

is.grouped_dt(x)
```

## Arguments

| | |
|---|---|
| data | a tbl or data frame. |
| vars | a list of quoted variables. |
| copy | If TRUE, will make copy of input. |
| x | an object to check |

## Examples

```
if (require("data.table") && require("nycflights13")) {
flights_dt <- tbl_dt(flights)
group_size(group_by(flights_dt, year, month, day))
group_size(group_by(flights_dt, dest))

monthly <- group_by(flights_dt, month)
summarise(monthly, n = n(), delay = mean(arr_delay))
}
```

## groups — *Get/set the grouping variables for tbl.*

### Description

These functions do not perform non-standard evaluation, and so are useful when programming against `tbl` objects. `ungroup` is a convenient inline way of removing existing grouping.

### Usage

```
groups(x)

ungroup(x)
```

### Arguments

| | |
|---|---|
| `x` | data `tbl` |

### Examples

```
grouped <- group_by(mtcars, cyl)
groups(grouped)
groups(ungroup(grouped))
```

## group_by — *Group a tbl by one or more variables.*

### Description

Most data operations are useful done on groups defined by variables in the the dataset. The `group_by` function takes an existing tbl and converts it into a grouped tbl where operations are performed "by group".

### Usage

```
group_by(.data, ..., add = FALSE)

group_by_(.data, ..., .dots, add = FALSE)
```

### Arguments

| | |
|---|---|
| `.data` | a tbl |
| `...` | variables to group by. All tbls accept variable names, some will also accept functions of variables. Duplicated groups will be silently dropped. |
| `add` | By default, when `add = FALSE`, `group_by` will override existing groups. To instead add to the existing groups, use `add = TRUE` |
| `.dots` | Used to work around non-standard evaluation. See `vignette("nse")` for details. |

### Tbl types

group_by is an S3 generic with methods for the three built-in tbls. See the help for the corresponding classes and their manip methods for more details:

- data.frame: grouped_df
- data.table: grouped_dt
- SQLite: src_sqlite
- PostgreSQL: src_postgres
- MySQL: src_mysql

### See Also

ungroup for the inverse operation, groups for accessors that don't do special evaluation.

### Examples

```
by_cyl <- group_by(mtcars, cyl)
summarise(by_cyl, mean(disp), mean(hp))
filter(by_cyl, disp == max(disp))

# summarise peels off a single layer of grouping
by_vs_am <- group_by(mtcars, vs, am)
by_vs <- summarise(by_vs_am, n = n())
by_vs
summarise(by_vs, n = sum(n))
# use ungroup() to remove if not wanted
summarise(ungroup(by_vs), n = sum(n))

# You can group by expressions: this is just short-hand for
# a mutate/rename followed by a simple group_by
group_by(mtcars, vsam = vs + am)
group_by(mtcars, vs2 = vs)

# You can also group by a constant, but it's not very useful
group_by(mtcars, "vs")

# By default, group_by sets groups. Use add = TRUE to add groups
groups(group_by(by_cyl, vs, am))
groups(group_by(by_cyl, vs, am, add = TRUE))

# Duplicate groups are silently dropped
groups(group_by(by_cyl, cyl, cyl))
```

## group_indices *Group id.*

### Description

Generate a unique id for each group

### Usage

```
group_indices(.data, ...)

group_indices_(.data, ..., .dots, add = FALSE)
```

### Arguments

| | |
|---|---|
| .data | a tbl |
| ... | variables to group by. All tbls accept variable names, some will also accept functions of variables. Duplicated groups will be silently dropped. |
| .dots | Used to work around non-standard evaluation. See vignette("nse") for details. |
| add | By default, when add = FALSE, group_by will override existing groups. To instead add to the existing groups, use add = TRUE |

### See Also

group_by

### Examples

```
group_indices(mtcars, cyl)
```

## group_size *Calculate group sizes.*

### Description

Calculate group sizes.

### Usage

```
group_size(x)

n_groups(x)
```

### Arguments

| | |
|---|---|
| x | a grouped tbl |

### Examples

```
if (require("nycflights13")) {

by_day <- flights %>% group_by(year, month, day)
n_groups(by_day)
group_size(by_day)

by_dest <- flights %>% group_by(dest)
n_groups(by_dest)
group_size(by_dest)
}
```

---

## join *Join two tbls together.*

---

### Description

These are generic functions that dispatch to individual tbl methods - see the method documentation for details of individual data sources. x and y should usually be from the same data source, but if copy is TRUE, y will automatically be copied to the same source as x - this may be an expensive operation.

### Usage

```
inner_join(x, y, by = NULL, copy = FALSE, ...)

left_join(x, y, by = NULL, copy = FALSE, ...)

right_join(x, y, by = NULL, copy = FALSE, ...)

full_join(x, y, by = NULL, copy = FALSE, ...)

semi_join(x, y, by = NULL, copy = FALSE, ...)

anti_join(x, y, by = NULL, copy = FALSE, ...)
```

### Arguments

| | |
|---|---|
| x,y | tbls to join |
| 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.<br>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. |

copy — If x and y are not from the same data source, and copy is TRUE, then y will be copied into the same src as x. This allows you to join tables across srcs, but it is a potentially expensive operation so you must opt into it.

... — other parameters passed onto methods

### Join types

Currently dplyr supports four join types:

inner_join return all rows from x where there are matching values in x, and all columns from x and y. If there are multiple matches between x and y, all combination of the matches are returned.

left_join return all rows from x, and all columns from x and y. Rows in x with no match in y will have NA values in the new columns. If there are multiple matches between x and y, all combinations of the matches are returned.

right_join return all rows from y, and all columns from x and y. Rows in y with no match in x will have NA values in the new columns. If there are multiple matches between x and y, all combinations of the matches are returned.

semi_join return all rows from x where there are matching values in y, keeping just columns from x.

A semi join differs from an inner join because an inner join will return one row of x for each matching row of y, where a semi join will never duplicate rows of x.

anti_join return all rows from x where there are not matching values in y, keeping just columns from x.

full_join return all rows and all columns from both x and y. Where there are not matching values, returns NA for the one missing.

### Grouping

Groups are ignored for the purpose of joining, but the result preserves the grouping of x.

---

## join.tbl_df *Join data frame tbls.*

---

### Description

See join for a description of the general purpose of the functions.

### Usage

```
## S3 method for class 'tbl_df'
inner_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'tbl_df'
left_join(x, y, by = NULL, copy = FALSE, ...)
```

```
## S3 method for class 'tbl_df'
right_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'tbl_df'
full_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'tbl_df'
semi_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'tbl_df'
anti_join(x, y, by = NULL, copy = FALSE, ...)
```

## Arguments

| | |
|---|---|
| x,y | tbls to join |
| 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, supply a character vector. |
| copy | If y is not a data frame or tbl_df and copy is TRUE, y will be converted into a data frame |
| ... | included for compatibility with the generic; otherwise ignored. |

## Examples

```
if (require("Lahman")) {
batting_df <- tbl_df(Batting)
person_df <- tbl_df(Master)

uperson_df <- tbl_df(Master[!duplicated(Master$playerID), ])

# Inner join: match batting and person data
inner_join(batting_df, person_df)
inner_join(batting_df, uperson_df)

# Left join: match, but preserve batting data
left_join(batting_df, uperson_df)

# Anti join: find batters without person data
anti_join(batting_df, person_df)
# or people who didn't bat
anti_join(person_df, batting_df)
}
```

---

# join.tbl_dt *Join data table tbls.*

---

## Description

See join for a description of the general purpose of the functions.

## Usage

```
## S3 method for class 'data.table'
inner_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'data.table'
left_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'data.table'
semi_join(x, y, by = NULL, copy = FALSE, ...)

## S3 method for class 'data.table'
anti_join(x, y, by = NULL, copy = FALSE, ...)
```

## Arguments

| | |
|---|---|
| x,y | tbls to join |
| 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.<br>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. |
| copy | If x and y are not from the same data source, and copy is TRUE, then y will be copied into the same src as x. This allows you to join tables across srcs, but it is a potentially expensive operation so you must opt into it. |
| ... | Included for compatibility with generic; otherwise ignored. |

## Examples

```
if (require("data.table") && require("Lahman")) {
batting_dt <- tbl_dt(Batting)
person_dt <- tbl_dt(Master)

# Inner join: match batting and person data
inner_join(batting_dt, person_dt)

# Left join: keep batting data even if person missing
left_join(batting_dt, person_dt)

# Semi-join: find batting data for top 4 teams, 2010:2012
grid <- expand.grid(
  teamID = c("WAS", "ATL", "PHI", "NYA"),
  yearID = 2010:2012)
top4 <- semi_join(batting_dt, grid, copy = TRUE)

# Anti-join: find batting data with out player data
```

```
anti_join(batting_dt, person_dt)
}
```

---

`join.tbl_sql` *Join sql tbls.*

---

## Description

See `join` for a description of the general purpose of the functions.

## Usage

```
## S3 method for class 'tbl_sql'
inner_join(x, y, by = NULL, copy = FALSE,
  auto_index = FALSE, ...)

## S3 method for class 'tbl_sql'
left_join(x, y, by = NULL, copy = FALSE,
  auto_index = FALSE, ...)

## S3 method for class 'tbl_sql'
semi_join(x, y, by = NULL, copy = FALSE,
  auto_index = FALSE, ...)

## S3 method for class 'tbl_sql'
anti_join(x, y, by = NULL, copy = FALSE,
  auto_index = FALSE, ...)
```

## Arguments

| | |
|---|---|
| `x,y` | tbls to join |
| `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.<br>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`. |
| `copy` | If `x` and `y` are not from the same data source, and `copy` is `TRUE`, then `y` will be copied into a temporary table in same database as `x`. `join` will automatically run `ANALYZE` on the created table in the hope that this will make you queries as efficient as possible by giving more data to the query planner.<br>This allows you to join tables across srcs, but it's potentially expensive operation so you must opt into it. |
| `auto_index` | if `copy` is `TRUE`, automatically create indices for the variables in `by`. This may speed up the join if there are matching indexes in `x`. |
| `...` | other parameters passed onto methods |

### Implementation notes

Semi-joins are implemented using `WHERE EXISTS`, and anti-joins with `WHERE NOT EXISTS`. Support for semi-joins is somewhat partial: you can only create semi joins where the `x` and `y` columns are compared with `=` not with more general operators.

### Examples

```
## Not run:
if (require("RSQLite") && has_lahman("sqlite")) {

# Left joins ------------------------------------------------------------------
lahman_s <- lahman_sqlite()
batting <- tbl(lahman_s, "Batting")
team_info <- select(tbl(lahman_s, "Teams"), yearID, lgID, teamID, G, R:H)

# Combine player and whole team statistics
first_stint <- select(filter(batting, stint == 1), playerID:H)
both <- left_join(first_stint, team_info, type = "inner", by = c("yearID", "teamID", "lgID"))
head(both)
explain(both)

# Join with a local data frame
grid <- expand.grid(
  teamID = c("WAS", "ATL", "PHI", "NYA"),
  yearID = 2010:2012)
top4a <- left_join(batting, grid, copy = TRUE)
explain(top4a)

# Indices don't really help here because there's no matching index on
# batting
top4b <- left_join(batting, grid, copy = TRUE, auto_index = TRUE)
explain(top4b)

# Semi-joins ------------------------------------------------------------------

people <- tbl(lahman_s, "Master")

# All people in half of fame
hof <- tbl(lahman_s, "HallOfFame")
semi_join(people, hof)

# All people not in the hall of fame
anti_join(people, hof)

# Find all managers
manager <- tbl(lahman_s, "Managers")
semi_join(people, manager)

# Find all managers in hall of fame
famous_manager <- semi_join(semi_join(people, manager), hof)
famous_manager
explain(famous_manager)
```

```
# Anti-joins ------------------------------------------------------------

# batters without person covariates
anti_join(batting, people)
}

## End(Not run)
```

---

lead-lag *Lead and lag.*

---

## Description

Lead and lag are useful for comparing values offset by a constant (e.g. the previous or next value)

## Usage

```
lead(x, n = 1L, default = NA, order_by = NULL, ...)

lag(x, n = 1L, default = NA, order_by = NULL, ...)
```

## Arguments

| | |
|---|---|
| x | a vector of values |
| n | a postive integer of length 1, giving the number of positions to lead or lag by |
| default | value used for non-existant rows. Defaults to NA. |
| order_by | override the default ordering to use another vector |
| ... | Needed for compatibility with lag generic. |

## Examples

```
lead(1:10, 1)
lead(1:10, 2)

lag(1:10, 1)
lead(1:10, 1)

x <- runif(5)
cbind(ahead = lead(x), x, behind = lag(x))

# Use order_by if data not already ordered
df <- data.frame(year = 2000:2005, value = (0:5) ^ 2)
scrambled <- df[sample(nrow(df)), ]

wrong <- mutate(scrambled, prev = lag(value))
arrange(wrong, year)

right <- mutate(scrambled, prev = lag(value, order_by = year))
arrange(right, year)
```

## location *Print the location in memory of a data frame*

### Description

This is useful for understand how and when dplyr makes copies of data frames

### Usage

```
location(df)

changes(x, y)
```

### Arguments

| | |
|---|---|
| df, | a data frame |
| x,y | two data frames to compare |

### Examples

```
location(mtcars)

mtcars2 <- mutate(mtcars, cyl2 = cyl * 2)
location(mtcars2)

changes(mtcars, mtcars)
changes(mtcars, mtcars2)
```

## mutate *Add new variables.*

### Description

Mutate adds new variables and preserves existing; transmute drops existing variables.

### Usage

```
mutate(.data, ...)

mutate_(.data, ..., .dots)

transmute(.data, ...)

transmute_(.data, ..., .dots)
```

### Arguments

| | |
|---|---|
| .data | A tbl. All main verbs are S3 generics and provide methods for tbl_df, tbl_dt and tbl_sql. |
| ... | Name-value pairs of expressions. Use NULL to drop a variable. |
| .dots | Used to work around non-standard evaluation. See vignette("nse") for details. |

### Value

An object of the same class as .data.

Data frame row names are silently dropped. To preserve, convert to an explicit variable.

### See Also

Other single.table.verbs: arrange, arrange_; filter, filter_; rename, rename_, select, select_; slice, slice_; summarise, summarise_, summarize, summarize_

### Examples

```
mutate(mtcars, displ_l = disp / 61.0237)
transmute(mtcars, displ_l = disp / 61.0237)

mutate(mtcars, cyl = NULL)
```

---

## n *The number of observations in the current group.*

---

### Description

This function is implemented special for each data source and can only be used from within summarise, mutate and filter

### Usage

```
n()
```

### Examples

```
if (require("nycflights13")) {
carriers <- group_by(flights, carrier)
summarise(carriers, n())
mutate(carriers, n = n())
filter(carriers, n() < 100)
}
```

nasa *NASA spatio-temporal data*

## Description

This data comes from the ASA 2007 data expo, http://stat-computing.org/dataexpo/2006/. The data are geographic and atmospheric measures on a very coarse 24 by 24 grid covering Central America. The variables are: temperature (surface and air), ozone, air pressure, and cloud cover (low, mid, and high). All variables are monthly averages, with observations for Jan 1995 to Dec 2000. These data were obtained from the NASA Langley Research Center Atmospheric Sciences Data Center (with permission; see important copyright terms below).

## Usage

```
nasa
```

## Format

A tbl_cube with 41,472 observations.

## Dimensions

- lat, long: latitude and longitude
- year, month: month and year

## Measures

- cloudlow, cloudmed, cloudhigh: cloud cover at three heights
- ozone
- surftemp and temperature
- pressure

## Examples

```
nasa
```

nth *Extract the first, last or nth value from a vector.*

## Description

These are straightforward wrappers around `[[`. The main advantage is that you can provide an optional secondary vector that defines the ordering, and provide a default value to use when the input is shorter than expected.

## Usage

```
nth(x, n, order_by = NULL, default = default_missing(x))

first(x, order_by = NULL, default = default_missing(x))

last(x, order_by = NULL, default = default_missing(x))
```

## Arguments

| | |
|---|---|
| `x` | A vector |
| `n` | For `nth_value`, a single integer specifying the position. If a numeric is supplied, it will be silently truncated. |
| `order_by` | An optional vector used to determine the order |
| `default` | A default value to use if the position does not exist in the input. This is guessed by default for atomic vectors, where a missing value of the appropriate type is return, and for lists, where a `NULL` is return. For more complicated objects, you'll need to supply this value. |

## Value

A single value. `[[` is used to do the subsetting.

## Examples

```
x <- 1:10
y <- 10:1

last(x)
last(x, y)
```

## n_distinct *Efficiently count the number of unique values in a vector.*

### Description

This is a faster and more concise equivalent of `length(unique(x))`

### Usage

```
n_distinct(x, na_rm = FALSE)
```

### Arguments

| | |
|---|---|
| `x` | a vector of values |
| `na_rm` | if `TRUE` missing values don't count |

### Examples

```
x <- sample(1:10, 1e5, rep = TRUE)
length(unique(x))
n_distinct(x)
```

## order_by *A helper function for ordering window function output.*

### Description

This is a useful function to control the order of window functions in R that don't have a specific ordering parameter. When translated to SQL it will modify the order clause of the OVER function.

### Usage

```
order_by(order_by, call)
```

### Arguments

| | |
|---|---|
| `order_by` | a vector to order_by |
| `call` | a function call to a window function, where the first argument is the vector being operated on |

### Details

This function works by changing the `call` to instead call `with_order` with the appropriate arguments.

## Examples

```
order_by(10:1, cumsum(1:10))
x <- 10:1
y <- 1:10
order_by(x, cumsum(y))

df <- data.frame(year = 2000:2005, value = (0:5) ^ 2)
scrambled <- df[sample(nrow(df)), ]

wrong <- mutate(scrambled, running = cumsum(value))
arrange(wrong, year)

right <- mutate(scrambled, running = order_by(year, cumsum(value)))
arrange(right, year)
```

---

# ranking *Windowed rank functions.*

---

## Description

Six variations on ranking functions, mimicing the ranking functions described in SQL2003. They are currently implemented using the built in `rank` function, and are provided mainly as a convenience when converting between R and SQL. All ranking functions map smallest inputs to smallest outputs. Use `desc` to reverse the direction..

## Usage

```
row_number(x)

ntile(x, n)

min_rank(x)

dense_rank(x)

percent_rank(x)

cume_dist(x)
```

## Arguments

| | |
|---|---|
| x | a vector of values to rank. Missing values are left as is. If you want to treat them as the smallest or largest values, replace with Inf or -Inf before ranking. |
| n | number of groups to split up into. |

### Details

- row_number: equivalent to rank(ties.method = "first")
- min_rank: equivalent to rank(ties.method = "min")
- dense_rank: like min_rank, but with no gaps between ranks
- percent_rank: a number between 0 and 1 computed by rescaling min_rank to [0, 1]
- cume_dist: a cumulative distribution function. Proportion of all values less than or equal to the current rank.
- ntile: a rough rank, which breaks the input vector into n buckets.

### Examples

```
x <- c(5, 1, 3, 2, 2, NA)
row_number(x)
min_rank(x)
dense_rank(x)
percent_rank(x)
cume_dist(x)

ntile(x, 2)
ntile(runif(100), 10)
```

---

## rowwise *Group input by rows*

---

### Description

rowwise is used for the results of do when you create list-variables. It is also useful to support arbitrary complex operations that need to be applied to each row.

### Usage

```
rowwise(data)
```

### Arguments

| | |
|---|---|
| data | Input data frame. |

### Details

Currently rowwise grouping only works with data frames. Its main impact is to allow you to work with list-variables in summarise and mutate without having to use [[1]]. This makes summarise() on a rowwise tbl effectively equivalent to plyr's ldply.

### Examples

```
df <- expand.grid(x = 1:3, y = 3:1)
df %>% rowwise() %>% do(i = seq(.$x, .$y))
.Last.value %>% summarise(n = length(i))
```

sample *Sample n rows from a table.*

## Description

This is a wrapper around sample.int to make it easy to select random rows from a table. It currently only works for local tbls.

## Usage

```
sample_n(tbl, size, replace = FALSE, weight = NULL, .env = parent.frame())

sample_frac(tbl, size = 1, replace = FALSE, weight = NULL,
  .env = parent.frame())
```

## Arguments

| | |
|---|---|
| tbl | tbl of data. |
| size | For sample_n, the number of rows to select. For sample_frac, the fraction of rows to select. If tbl is grouped, size applies to each group. |
| replace | Sample with or without replacement? |
| weight | Sampling weights. This expression is evaluated in the context of the data frame. It must return a vector of non-negative numbers the same length as the input. Weights are automatically standardised to sum to 1. |
| .env | Environment in which to look for non-data names used in weight. Non-default settings for experts only. |

## Examples

```
by_cyl <- mtcars %>% group_by(cyl)

# Sample fixed number per group
sample_n(mtcars, 10)
sample_n(mtcars, 50, replace = TRUE)
sample_n(mtcars, 10, weight = mpg)

sample_n(by_cyl, 3)
sample_n(by_cyl, 10, replace = TRUE)
sample_n(by_cyl, 3, weight = mpg / mean(mpg))

# Sample fixed fraction per group
# Default is to sample all data = randomly resample rows
sample_frac(mtcars)

sample_frac(mtcars, 0.1)
sample_frac(mtcars, 1.5, replace = TRUE)
sample_frac(mtcars, 0.1, weight = 1 / mpg)
```

```
sample_frac(by_cyl, 0.2)
sample_frac(by_cyl, 1, replace = TRUE)
```

---

`select` *Select/rename variables by name.*

---

## Description

`select()` keeps only the variables you mention; `rename()` keeps all variables.

## Usage

```
select(.data, ...)

select_(.data, ..., .dots)

rename(.data, ...)

rename_(.data, ..., .dots)
```

## Arguments

| | |
|---|---|
| `.data` | A tbl. All main verbs are S3 generics and provide methods for `tbl_df`, `tbl_dt` and `tbl_sql`. |
| `...` | Comma separated list of unquoted expressions. You can treat variable names like they are positions. Use positive values to select variables; use negative values to drop variables. |
| `.dots` | Use `select_()` to do standard evaluation. See `vignette("nse")` for details |

## Value

An object of the same class as `.data`.

Data frame row names are silently dropped. To preserve, convert to an explicit variable.

## Special functions

As well as using existing functions like `:` and `c`, there are a number of special functions that only work inside `select`

- `starts_with(x, ignore.case = TRUE)`: names starts with `x`
- `ends_with(x, ignore.case = TRUE)`: names ends in `x`
- `contains(x, ignore.case = TRUE)`: selects all variables whose name contains `x`
- `matches(x, ignore.case = TRUE)`: selects all variables whose name matches the regular expression `x`
- `num_range("x", 1:5, width = 2)`: selects all variables (numerically) from x01 to x05.

- `one_of("x", "y", "z")`: selects variables provided in a character vector.
- `everything()`: selects all variables.

To drop variables, use `-`. You can rename variables with named arguments.

## See Also

Other single.table.verbs: `arrange`, `arrange_`; `filter`, `filter_`; `mutate`, `mutate_`, `transmute`, `transmute_`; `slice`, `slice_`; `summarise`, `summarise_`, `summarize`, `summarize_`

## Examples

```
iris <- tbl_df(iris) # so it prints a little nicer
select(iris, starts_with("Petal"))
select(iris, ends_with("Width"))
select(iris, contains("etal"))
select(iris, matches(".t."))
select(iris, Petal.Length, Petal.Width)
vars <- c("Petal.Length", "Petal.Width")
select(iris, one_of(vars))

df <- as.data.frame(matrix(runif(100), nrow = 10))
df <- tbl_df(df[c(3, 4, 7, 1, 9, 8, 5, 2, 6, 10)])
select(df, V4:V6)
select(df, num_range("V", 4:6))

# Drop variables
select(iris, -starts_with("Petal"))
select(iris, -ends_with("Width"))
select(iris, -contains("etal"))
select(iris, -matches(".t."))
select(iris, -Petal.Length, -Petal.Width)

# Rename variables:
# * select() keeps only the variables you specify
select(iris, petal_length = Petal.Length)
# Renaming multiple variables uses a prefix:
select(iris, petal = starts_with("Petal"))

# Reorder variables: keep the variable "Species" in the front
select(iris, Species, everything())

# * rename() keeps all variables
rename(iris, petal_length = Petal.Length)

# Programming with select ----------------------------------------------------
select_(iris, ~Petal.Length)
select_(iris, "Petal.Length")
select_(iris, lazyeval::interp(~matches(x), x = ".t."))
select_(iris, quote(-Petal.Length), quote(-Petal.Width))
select_(iris, .dots = list(quote(-Petal.Length), quote(-Petal.Width)))
```

## setops *Set operations.*

### Description

These functions override the set functions provided in base to make them generic so that efficient versions for data frames and other tables can be provided. The default methods call the base versions.

### Usage

```
intersect(x, y, ...)

union(x, y, ...)

setdiff(x, y, ...)

setequal(x, y, ...)
```

### Arguments

| | |
|---|---|
| x,y | objects to compare (ignoring order) |
| ... | other arguments passed on to methods |

### Examples

```
mtcars$model <- rownames(mtcars)
first <- mtcars[1:20, ]
second <- mtcars[10:32, ]

intersect(first, second)
union(first, second)
setdiff(first, second)
setdiff(second, first)

setequal(mtcars, mtcars[32:1, ])
```

## slice *Select rows by position.*

### Description

Slice does not work with relational databases because they have no intrinsic notion of row order. If you want to perform the equivalent operation, use `filter()` and `row_number()`.

## Usage

```
slice(.data, ...)

slice_(.data, ..., .dots)
```

## Arguments

| | |
|---|---|
| .data | A tbl. All main verbs are S3 generics and provide methods for tbl_df, tbl_dt and tbl_sql. |
| ... | Integer row values |
| .dots | Used to work around non-standard evaluation. See vignette("nse") for details. |

## See Also

Other single.table.verbs: arrange, arrange_; filter, filter_; mutate, mutate_, transmute, transmute_; rename, rename_, select, select_; summarise, summarise_, summarize, summarize_

## Examples

```
slice(mtcars, 1L)
slice(mtcars, n())
slice(mtcars, 5:n())

by_cyl <- group_by(mtcars, cyl)
slice(by_cyl, 1:2)

# Equivalent code using filter that will also work with databases,
# but won't be as fast for in-memory data. For many databases, you'll
# need to supply an explicit variable to use to compute the row number.
filter(mtcars, row_number() == 1L)
filter(mtcars, row_number() == n())
filter(mtcars, between(row_number(), 5, n()))
```

---

# src_mysql *Connect to mysql/mariadb.*

---

## Description

Use src_mysql to connect to an existing mysql or mariadb database, and tbl to connect to tables within that database. If you are running a local mysqlql database, leave all parameters set as their defaults to connect. If you're connecting to a remote database, ask your database administrator for the values of these variables.

## Usage

```
src_mysql(dbname, host = NULL, port = 0L, user = "root", password = "",
  ...)

## S3 method for class 'src_mysql'
tbl(src, from, ...)
```

## Arguments

| | |
|---|---|
| dbname | Database name |
| host,port | Host name and port number of database |
| user,password | User name and password. Rather than supplying a username and password here, it's better to save them in my.cnf, as described in MySQL. In that case, supply NULL to both user and password. |
| ... | for the src, other arguments passed on to the underlying database connector, dbConnect. For the tbl, included for compatibility with the generic, but otherwise ignored. |
| src | a mysql src created with src_mysql. |
| from | Either a string giving the name of table in database, or sql described a derived table or compound join. |

## Debugging

To see exactly what SQL is being sent to the database, you see show_query and explain.

## Grouping

Typically you will create a grouped data table is to call the group_by method on a mysql tbl: this will take care of capturing the unevalated expressions for you.

For best performance, the database should have an index on the variables that you are grouping by. Use explain to check that the database is using the indexes that you expect.

## Output

All data manipulation on SQL tbls are lazy: they will not actually run the query or retrieve the data unless you ask for it: they all return a new tbl_sql object. Use compute to run the query and save the results in a temporary in the database, or use collect to retrieve the results to R.

Note that do is not lazy since it must pull the data into R. It returns a tbl_df or grouped_df, with one column for each grouping variable, and one list column that contains the results of the operation. do never simplifies its output.

## Query principles

This section attempts to lay out the principles governing the generation of SQL queries from the manipulation verbs. The basic principle is that a sequence of operations should return the same value (modulo class) regardless of where the data is stored.

- `arrange(arrange(df, x), y)` should be equivalent to `arrange(df, y, x)`
- `select(select(df, a:x), n:o)` should be equivalent to `select(df, n:o)`
- `mutate(mutate(df, x2 = x * 2), y2 = y * 2)` should be equivalent to `mutate(df, x2 = x * 2, y2 = y * 2)`
- `filter(filter(df, x == 1), y == 2)` should be equivalent to `filter(df, x == 1, y == 2)`
- `summarise` should return the summarised output with one level of grouping peeled off.

## Examples

```
## Not run:
# Connection basics ---------------------------------------------------------
# To connect to a database first create a src:
my_db <- src_mysql(host = "blah.com", user = "hadley",
  password = "pass")
# Then reference a tbl within that src
my_tbl <- tbl(my_db, "my_table")

## End(Not run)

# Here we'll use the Lahman database: to create your own local copy,
# create a local database called "lahman", or tell lahman_mysql() how to
# a database that you can write to

if (!has_lahman("postgres") && has_lahman("mysql")) {
lahman_m <- lahman_mysql()
# Methods -------------------------------------------------------------------
batting <- tbl(lahman_m, "Batting")
dim(batting)
colnames(batting)
head(batting)

# Data manipulation verbs ---------------------------------------------------
filter(batting, yearID > 2005, G > 130)
select(batting, playerID:lgID)
arrange(batting, playerID, desc(yearID))
summarise(batting, G = mean(G), n = n())
mutate(batting, rbi2 = 1.0 * R / AB)

# note that all operations are lazy: they don't do anything until you
# request the data, either by `print()`ing it (which shows the first ten
# rows), by looking at the `head()`, or `collect()` the results locally.

system.time(recent <- filter(batting, yearID > 2010))
system.time(collect(recent))

# Group by operations -------------------------------------------------------
# To perform operations by group, create a grouped object with group_by
players <- group_by(batting, playerID)
group_size(players)

# MySQL doesn't support windowed functions, which means that only
# grouped summaries are really useful:
```

```
summarise(players, mean_g = mean(G), best_ab = max(AB))

# When you group by multiple level, each summarise peels off one level
per_year <- group_by(batting, playerID, yearID)
stints <- summarise(per_year, stints = max(stint))
filter(ungroup(stints), stints > 3)
summarise(stints, max(stints))

# Joins ------------------------------------------------------------------
player_info <- select(tbl(lahman_m, "Master"), playerID,
  birthYear)
hof <- select(filter(tbl(lahman_m, "HallOfFame"), inducted == "Y"),
 playerID, votedBy, category)

# Match players and their hall of fame data
inner_join(player_info, hof)
# Keep all players, match hof data where available
left_join(player_info, hof)
# Find only players in hof
semi_join(player_info, hof)
# Find players not in hof
anti_join(player_info, hof)

# Arbitrary SQL ----------------------------------------------------------
# You can also provide sql as is, using the sql function:
batting2008 <- tbl(lahman_m,
  sql("SELECT * FROM Batting WHERE YearID = 2008"))
batting2008
}
```

---

`src_postgres` *Connect to postgresql.*

---

## Description

Use `src_postgres` to connect to an existing postgresql database, and `tbl` to connect to tables within that database. If you are running a local postgresql database, leave all parameters set as their defaults to connect. If you're connecting to a remote database, ask your database administrator for the values of these variables.

## Usage

```
src_postgres(dbname = NULL, host = NULL, port = NULL, user = NULL,
  password = NULL, ...)

## S3 method for class 'src_postgres'
tbl(src, from, ...)
```

## Arguments

| | |
|---|---|
| dbname | Database name |
| host,port | Host name and port number of database |
| user,password | User name and password (if needed) |
| ... | for the src, other arguments passed on to the underlying database connector, dbConnect. For the tbl, included for compatibility with the generic, but otherwise ignored. |
| src | a postgres src created with src_postgres. |
| from | Either a string giving the name of table in database, or sql described a derived table or compound join. |

## Debugging

To see exactly what SQL is being sent to the database, you see show_query and explain.

## Grouping

Typically you will create a grouped data table is to call the group_by method on a mysql tbl: this will take care of capturing the unevalated expressions for you.

For best performance, the database should have an index on the variables that you are grouping by. Use explain to check that the database is using the indexes that you expect.

## Output

All data manipulation on SQL tbls are lazy: they will not actually run the query or retrieve the data unless you ask for it: they all return a new tbl_sql object. Use compute to run the query and save the results in a temporary in the database, or use collect to retrieve the results to R.

Note that do is not lazy since it must pull the data into R. It returns a tbl_df or grouped_df, with one column for each grouping variable, and one list column that contains the results of the operation. do never simplifies its output.

## Query principles

This section attempts to lay out the principles governing the generation of SQL queries from the manipulation verbs. The basic principle is that a sequence of operations should return the same value (modulo class) regardless of where the data is stored.

- `arrange(arrange(df, x), y)` should be equivalent to `arrange(df, y, x)`
- `select(select(df, a:x), n:o)` should be equivalent to `select(df, n:o)`
- `mutate(mutate(df, x2 = x * 2), y2 = y * 2)` should be equivalent to `mutate(df, x2 = x * 2, y2 = y * 2)`
- `filter(filter(df, x == 1), y == 2)` should be equivalent to `filter(df, x == 1, y == 2)`
- `summarise` should return the summarised output with one level of grouping peeled off.

## Examples

```
## Not run:
# Connection basics ---------------------------------------------------------
# To connect to a database first create a src:
my_db <- src_postgres(host = "blah.com", user = "hadley",
  password = "pass")
# Then reference a tbl within that src
my_tbl <- tbl(my_db, "my_table")

## End(Not run)

# Here we'll use the Lahman database: to create your own local copy,
# create a local database called "lahman", or tell lahman_postgres() how to
# access a database that you can write to

if (has_lahman("postgres")) {
lahman_p <- lahman_postgres()
# Methods -------------------------------------------------------------------
batting <- tbl(lahman_p, "Batting")
dim(batting)
colnames(batting)
head(batting)

# Data manipulation verbs ---------------------------------------------------
filter(batting, yearID > 2005, G > 130)
select(batting, playerID:lgID)
arrange(batting, playerID, desc(yearID))
summarise(batting, G = mean(G), n = n())
mutate(batting, rbi2 = if(is.null(AB)) 1.0 * R / AB else 0)

# note that all operations are lazy: they don't do anything until you
# request the data, either by `print()`ing it (which shows the first ten
# rows), by looking at the `head()`, or `collect()` the results locally.

system.time(recent <- filter(batting, yearID > 2010))
system.time(collect(recent))

# Group by operations -------------------------------------------------------
# To perform operations by group, create a grouped object with group_by
players <- group_by(batting, playerID)
group_size(players)

summarise(players, mean_g = mean(G), best_ab = max(AB))
best_year <- filter(players, AB == max(AB) | G == max(G))
progress <- mutate(players,
  cyear = yearID - min(yearID) + 1,
  ab_rank = rank(desc(AB)),
  cumulative_ab = order_by(yearID, cumsum(AB)))

# When you group by multiple level, each summarise peels off one level
per_year <- group_by(batting, playerID, yearID)
stints <- summarise(per_year, stints = max(stint))
```

```
filter(stints, stints > 3)
summarise(stints, max(stints))
mutate(stints, order_by(yearID, cumsum(stints)))

# Joins ------------------------------------------------------------------------
player_info <- select(tbl(lahman_p, "Master"), playerID, birthYear)
hof <- select(filter(tbl(lahman_p, "HallOfFame"), inducted == "Y"),
 playerID, votedBy, category)

# Match players and their hall of fame data
inner_join(player_info, hof)
# Keep all players, match hof data where available
left_join(player_info, hof)
# Find only players in hof
semi_join(player_info, hof)
# Find players not in hof
anti_join(player_info, hof)

# Arbitrary SQL ----------------------------------------------------------------
# You can also provide sql as is, using the sql function:
batting2008 <- tbl(lahman_p,
  sql('SELECT * FROM "Batting" WHERE "yearID" = 2008'))
batting2008
}
```

---

`src_sqlite` *Connect to a sqlite database.*

---

## Description

Use `src_sqlite` to connect to an existing sqlite database, and `tbl` to connect to tables within that database. If you are running a local sqliteql database, leave all parameters set as their defaults to connect. If you're connecting to a remote database, ask your database administrator for the values of these variables.

## Usage

```
src_sqlite(path, create = FALSE)

## S3 method for class 'src_sqlite'
tbl(src, from, ...)
```

## Arguments

| | |
|---|---|
| `path` | Path to SQLite database |
| `create` | if `FALSE`, `path` must already exist. If `TRUE`, will create a new SQlite3 database at `path`. |
| `src` | a sqlite src created with `src_sqlite`. |

| | |
|---|---|
| from | Either a string giving the name of table in database, or sql described a derived table or compound join. |
| ... | Included for compatibility with the generic, but otherwise ignored. |

## Debugging

To see exactly what SQL is being sent to the database, you see show_query and explain.

## Grouping

Typically you will create a grouped data table is to call the group_by method on a mysql tbl: this will take care of capturing the unevalated expressions for you.

For best performance, the database should have an index on the variables that you are grouping by. Use explain to check that the database is using the indexes that you expect.

## Output

All data manipulation on SQL tbls are lazy: they will not actually run the query or retrieve the data unless you ask for it: they all return a new tbl_sql object. Use compute to run the query and save the results in a temporary in the database, or use collect to retrieve the results to R.

Note that do is not lazy since it must pull the data into R. It returns a tbl_df or grouped_df, with one column for each grouping variable, and one list column that contains the results of the operation. do never simplifies its output.

## Query principles

This section attempts to lay out the principles governing the generation of SQL queries from the manipulation verbs. The basic principle is that a sequence of operations should return the same value (modulo class) regardless of where the data is stored.

- `arrange(arrange(df, x), y)` should be equivalent to `arrange(df, y, x)`
- `select(select(df, a:x), n:o)` should be equivalent to `select(df, n:o)`
- `mutate(mutate(df, x2 = x * 2), y2 = y * 2)` should be equivalent to `mutate(df, x2 = x * 2, y2 = y * 2)`
- `filter(filter(df, x == 1), y == 2)` should be equivalent to `filter(df, x == 1, y == 2)`
- `summarise` should return the summarised output with one level of grouping peeled off.

## Examples

```
## Not run:
# Connection basics ---------------------------------------------------------
# To connect to a database first create a src:
my_db <- src_sqlite(path = tempfile(), create = TRUE)
# Then reference a tbl within that src
my_tbl <- tbl(my_db, "my_table")

## End(Not run)

# Here we'll use the Lahman database: to create your own local copy,
```

```
# run lahman_sqlite()

## Not run:
if (requireNamespace("RSQLite") && has_lahman("sqlite")) {
lahman_s <- lahman_sqlite()
# Methods -------------------------------------------------------------------
batting <- tbl(lahman_s, "Batting")
dim(batting)
colnames(batting)
head(batting)

# Data manipulation verbs ---------------------------------------------------
filter(batting, yearID > 2005, G > 130)
select(batting, playerID:lgID)
arrange(batting, playerID, desc(yearID))
summarise(batting, G = mean(G), n = n())
mutate(batting, rbi2 = 1.0 * R / AB)

# note that all operations are lazy: they don't do anything until you
# request the data, either by `print()`ing it (which shows the first ten
# rows), by looking at the `head()`, or `collect()` the results locally.

system.time(recent <- filter(batting, yearID > 2010))
system.time(collect(recent))

# Group by operations -------------------------------------------------------
# To perform operations by group, create a grouped object with group_by
players <- group_by(batting, playerID)
group_size(players)

# sqlite doesn't support windowed functions, which means that only
# grouped summaries are really useful:
summarise(players, mean_g = mean(G), best_ab = max(AB))

# When you group by multiple level, each summarise peels off one level
per_year <- group_by(batting, playerID, yearID)
stints <- summarise(per_year, stints = max(stint))
filter(ungroup(stints), stints > 3)
summarise(stints, max(stints))

# Joins ---------------------------------------------------------------------
player_info <- select(tbl(lahman_s, "Master"), playerID, birthYear)
hof <- select(filter(tbl(lahman_s, "HallOfFame"), inducted == "Y"),
 playerID, votedBy, category)

# Match players and their hall of fame data
inner_join(player_info, hof)
# Keep all players, match hof data where available
left_join(player_info, hof)
# Find only players in hof
semi_join(player_info, hof)
# Find players not in hof
anti_join(player_info, hof)
```

```
# Arbitrary SQL ------------------------------------------------------------
# You can also provide sql as is, using the sql function:
batting2008 <- tbl(lahman_s,
  sql("SELECT * FROM Batting WHERE YearID = 2008"))
batting2008
}

## End(Not run)
```

---

`src_tbls` *List all tbls provided by a source.*

---

## Description

This is a generic method which individual src's will provide methods for. Most methods will not be documented because it's usually pretty obvious what possible results will be.

## Usage

```
src_tbls(x)
```

## Arguments

| | |
|---|---|
| `x` | a data src. |

---

`summarise` *Summarise multiple values to a single value.*

---

## Description

Summarise multiple values to a single value.

## Usage

```
summarise(.data, ...)

summarise_(.data, ..., .dots)

summarize(.data, ...)

summarize_(.data, ..., .dots)
```

## Arguments

| | |
|---|---|
| .data | A tbl. All main verbs are S3 generics and provide methods for tbl_df, tbl_dt and tbl_sql. |
| ... | Name-value pairs of summary functions like min(), mean(), max() etc. |
| .dots | Used to work around non-standard evaluation. See vignette("nse") for details. |

## Value

An object of the same class as .data. One grouping level will be dropped.

Data frame row names are silently dropped. To preserve, convert to an explicit variable.

## Backend variations

Data frames are the only backend that supports creating a variable and using it in the same summary. See examples for more details.

## See Also

Other single.table.verbs: arrange, arrange_; filter, filter_; mutate, mutate_, transmute, transmute_; rename, rename_, select, select_; slice, slice_

## Examples

```
summarise(mtcars, mean(disp))
summarise(group_by(mtcars, cyl), mean(disp))
summarise(group_by(mtcars, cyl), m = mean(disp), sd = sd(disp))

# With data frames, you can create and immediately use summaries
by_cyl <- mtcars %>% group_by(cyl)
by_cyl %>% summarise(a = n(), b = a + 1)

## Not run:
# You can't with data tables or databases
by_cyl_dt <- mtcars %>% tbl_dt() %>% group_by(cyl)
by_cyl_dt %>% summarise(a = n(), b = a + 1)

by_cyl_db <- src_sqlite(":memory:", create = TRUE) %>%
  copy_to(mtcars) %>% group_by(cyl)
by_cyl_db %>% summarise(a = n(), b = a + 1)

## End(Not run)
```

summarise_each *Summarise and mutate multiple columns.*

## Description

Apply one or more functions to one or more columns. Grouping variables are always excluded from modification.

## Usage

```
summarise_each(tbl, funs, ...)

summarise_each_(tbl, funs, vars)

summarize_each(tbl, funs, ...)

summarize_each_(tbl, funs, vars)

mutate_each(tbl, funs, ...)

mutate_each_(tbl, funs, vars)
```

## Arguments

| | |
|---|---|
| tbl | a tbl |
| funs | List of function calls, generated by funs, or a character vector of function names. |
| vars,... | Variables to include/exclude in mutate/summarise. You can use same specifications as in select. If missing, defaults to all non-grouping variables.<br>For standard evaluation versions (ending in _) these can be either a list of expressions or a character vector. |

## Examples

```
# One function
by_species <- iris %>% group_by(Species)
by_species %>% summarise_each(funs(length))
by_species %>% summarise_each(funs(mean))
by_species %>% summarise_each(funs(mean), Petal.Width)
by_species %>% summarise_each(funs(mean), matches("Width"))

by_species %>% mutate_each(funs(half = . / 2))
by_species %>% mutate_each(funs(min_rank))

# Two functions
by_species %>% summarise_each(funs(min, max))
by_species %>% summarise_each(funs(min, max), Petal.Width, Sepal.Width)
by_species %>% summarise_each(funs(min, max), matches("Width"))
```

```
# Alternative function specification
iris %>% summarise_each(funs(ul = length(unique(.))))
by_species %>% summarise_each(funs(ul = length(unique(.))))

by_species %>% summarise_each(c("min", "max"))

# Alternative variable specification
summarise_each_(iris, funs(max), names(iris)[-5])
summarise_each_(iris, funs(max), list(quote(-Species)))
```

---

`tally` *Counts/tally observations by group.*

---

## Description

`tally` is a convenient wrapper for summarise that will either call `n` or `sum(n)` depending on whether you're tallying for the first time, or re-tallying. `count()` is similar, but also does the `group_by` for you.

## Usage

```
tally(x, wt, sort = FALSE)

count(x, ..., wt = NULL, sort = FALSE)

count_(x, vars, wt = NULL, sort = FALSE)
```

## Arguments

| | |
|---|---|
| `x` | a `tbl` to tally/count. |
| `wt` | (Optional) If not specified, will tally the number of rows. If specified, will perform a "weighted" tally but summing over the specified variable. |
| `sort` | if `TRUE` will sort output in descending order of `n` |
| `...,vars` | Variables to group by. |

## Examples

```
if (require("Lahman")) {
batting_tbl <- tbl_df(Batting)
tally(group_by(batting_tbl, yearID))
tally(group_by(batting_tbl, yearID), sort = TRUE)

# Multiple tallys progressively roll up the groups
plays_by_year <- tally(group_by(batting_tbl, playerID, stint), sort = TRUE)
tally(plays_by_year, sort = TRUE)
tally(tally(plays_by_year))
```

```
# This looks a little nicer if you use the infix %>% operator
batting_tbl %>% group_by(playerID) %>% tally(sort = TRUE)

# count is even more succinct - it also does the grouping for you
batting_tbl %>% count(playerID)
batting_tbl %>% count(playerID, wt = G)
batting_tbl %>% count(playerID, wt = G, sort = TRUE)
}
```

---

## tbl *Create a table from a data source*

---

### Description

This is a generic method that dispatches based on the first argument.

### Usage

```
tbl(src, ...)

is.tbl(x)

as.tbl(x, ...)
```

### Arguments

| | |
|---|---|
| src | A data source |
| ... | Other arguments passed on to the individual methods |
| x | an object to coerce to a tbl |

---

## tbl_cube *A data cube tbl.*

---

### Description

An cube tbl stores data in a compact array format where dimension names are not needlessly repeated. They are particularly appropriate for experimental data where all combinations of factors are tried (e.g. complete factorial designs), or for storing the result of aggregations. Compared to data frames, they will occupy much less memory when variables are crossed, not nested.

### Usage

```
tbl_cube(dimensions, measures)
```

## Arguments

| | |
|---|---|
| `dimensions` | A named list of vectors. A dimension is a variable whose values are known before the experiement is conducted; they are fixed by design (in **reshape2** they are known as id variables). `tbl_cubes` are dense which means that almost every combination of the dimensions should have associated measurements: missing values require an explicit NA, so if the variables are nested, not crossed, the majority of the data structure will be empty. Dimensions are typically, but not always, categorical variables. |
| `measures` | A named list of arrays. A measure is something that is actually measured, and is not known in advance. The dimension of each array should be the same as the length of the dimensions. Measures are typically, but not always, continuous values. |

## Details

`tbl_cube` support is currently experimental and little performance optimisation has been done, but you may find them useful if your data already comes in this form, or you struggle with the memory overhead of the sparse/crossed of data frames. There is no supported for hierarchical indices (although I think that would be a relatively straightforward extension to storing data frames for indices rather than vectors).

## Implementation

Manipulation functions:

- `select` (M)
- `summarise` (M), corresponds to roll-up, but rather more limited since there are no hierarchies.
- `filter` (D), corresponds to slice/dice.
- `mutate` (M) is not implemented, but should be relatively straightforward given the implementation of `summarise`.
- `arrange` (D?) Not implemented: not obvious how much sense it would make

Joins: not implemented. See `vignettes/joins.graffle` for ideas. Probably straightforward if you get the indexes right, and that's probably some straightforward array/tensor operation.

## See Also

`as.tbl_cube` for ways of coercing existing data structures into a `tbl_cube`.

## Examples

```
# The built in nasa dataset records meterological data (temperature,
# cloud cover, ozone etc) for a 4d spatio-temporal dataset (lat, long,
# month and year)
nasa
head(as.data.frame(nasa))

titanic <- as.tbl_cube(Titanic)
```

```
head(as.data.frame(titanic))

admit <- as.tbl_cube(UCBAdmissions)
head(as.data.frame(admit))

as.tbl_cube(esoph, dim_names = 1:3)

# Some manipulation examples with the NASA dataset --------------------------

# select() operates only on measures: it doesn't affect dimensions in any way
select(nasa, cloudhigh:cloudmid)
select(nasa, matches("temp"))

# filter() operates only on dimensions
filter(nasa, lat > 0, year == 2000)
# Each component can only refer to one dimensions, ensuring that you always
# create a rectangular subset
## Not run: filter(nasa, lat > long)

# Arrange is meaningless for tbl_cubes

by_loc <- group_by(nasa, lat, long)
summarise(by_loc, pressure = max(pressure), temp = mean(temperature))
```

---

`tbl_df` *Create a data frame tbl.*

---

## Description

A data frame tbl wraps a local data frame. The main advantage to using a `tbl_df` over a regular data frame is the printing: tbl objects only print a few rows and all the columns that fit on one screen, describing the rest of it as text.

## Usage

```
tbl_df(data)
```

## Arguments

| | |
|---|---|
| `data` | a data frame |

## Methods

`tbl_df` implements two important base methods:

**print** Only prints the first 10 rows, and the columns that fit on screen

`[` Never simplifies (drops), so always returns data.frame

## Examples

```
ds <- tbl_df(mtcars)
ds
as.data.frame(ds)

if (require("Lahman") && packageVersion("Lahman") >= "3.0.1") {
batting <- tbl_df(Batting)
dim(batting)
colnames(batting)
head(batting)

# Data manipulation verbs ---------------------------------------------------
filter(batting, yearID > 2005, G > 130)
select(batting, playerID:lgID)
arrange(batting, playerID, desc(yearID))
summarise(batting, G = mean(G), n = n())
mutate(batting, rbi2 = if(is.null(AB)) 1.0 * R / AB else 0)

# Group by operations -------------------------------------------------------
# To perform operations by group, create a grouped object with group_by
players <- group_by(batting, playerID)
head(group_size(players), 100)

summarise(players, mean_g = mean(G), best_ab = max(AB))
best_year <- filter(players, AB == max(AB) | G == max(G))
progress <- mutate(players, cyear = yearID - min(yearID) + 1,
 rank(desc(AB)), cumsum(AB))

# When you group by multiple level, each summarise peels off one level

per_year <- group_by(batting, playerID, yearID)
stints <- summarise(per_year, stints = max(stint))
filter(stints, stints > 3)
summarise(stints, max(stints))
mutate(stints, cumsum(stints))


# Joins ---------------------------------------------------------------------
player_info <- select(tbl_df(Master), playerID, birthYear)
hof <- select(filter(tbl_df(HallOfFame), inducted == "Y"),
 playerID, votedBy, category)

# Match players and their hall of fame data
inner_join(player_info, hof)
# Keep all players, match hof data where available
left_join(player_info, hof)
# Find only players in hof
semi_join(player_info, hof)
# Find players not in hof
anti_join(player_info, hof)
}
```

tbl_dt *Create a data table tbl.*

## Description

A data table tbl wraps a local data table.

## Usage

```
tbl_dt(data, copy = TRUE)
```

## Arguments

| | |
|---|---|
| data | a data table |
| copy | If the input is a data.table, copy it? |

## Examples

```
if (require("data.table")) {
ds <- tbl_dt(mtcars)
ds
as.data.table(ds)
as.tbl(mtcars)
}

if (require("data.table") && require("nycflights13")) {
flights2 <- tbl_dt(flights)
flights2 %>% filter(month == 1, day == 1, dest == "DFW")
flights2 %>% select(year:day)
flights2 %>% rename(Year = year)
flights2 %>%
  summarise(
    delay = mean(arr_delay, na.rm = TRUE),
    n = length(arr_delay)
  )
flights2 %>%
  mutate(gained = arr_delay - dep_delay) %>%
  select(ends_with("delay"), gained)
flights2 %>%
  arrange(dest, desc(arr_delay))

by_dest <- group_by(flights2, dest)

filter(by_dest, arr_delay == max(arr_delay, na.rm = TRUE))
summarise(by_dest, arr = mean(arr_delay, na.rm = TRUE))

# Normalise arrival and departure delays by airport
by_dest %>%
  mutate(arr_z = scale(arr_delay), dep_z = scale(dep_delay)) %>%
  select(starts_with("arr"), starts_with("dep"))
```

```
arrange(by_dest, desc(arr_delay))
select(by_dest, -(day:tailnum))
rename(by_dest, Year = year)

# All manip functions preserve grouping structure, except for summarise
# which removes a grouping level
by_day <- group_by(flights2, year, month, day)
by_month <- summarise(by_day, delayed = sum(arr_delay > 0, na.rm = TRUE))
by_month
summarise(by_month, delayed = sum(delayed))

# You can also manually ungroup:
ungroup(by_day)
}
```

---

## tbl_vars *List variables provided by a tbl.*

---

### Description

List variables provided by a tbl.

### Usage

```
tbl_vars(x)
```

### Arguments

| | |
|---|---|
| x | A tbl object |

---

## top_n *Select top n rows (by value).*

---

### Description

This is a convenient wrapper that uses `filter` and `min_rank` to select the top n entries in each group, ordered by `wt`.

### Usage

```
top_n(x, n, wt)
```

### Arguments

| | |
|---|---|
| x | a tbl to filter |
| n | number of rows to return. If x is grouped, this is the number of rows per group. May include more than n if there are ties. |
| wt | (Optional). The variable to use for ordering. If not specified, defaults to the last variable in the tbl. |

### Examples

```
# Find 10 players with most games
if (require("Lahman")) {
players <- group_by(tbl_df(Batting), playerID)
games <- tally(players, G)
top_n(games, 10, n)

# A little nicer with %>%
tbl_df(Batting) %>%
  group_by(playerID) %>%
  tally(G) %>%
  top_n(10)

# Find year with most games for each player
tbl_df(Batting) %>% group_by(playerID) %>% top_n(1, G)
}
```

---

## translate_sql *Translate an expression to sql.*

---

### Description

Translate an expression to sql.

### Usage

```
translate_sql(..., tbl = NULL, env = parent.frame(), variant = NULL,
  window = FALSE)

translate_sql_q(expr, tbl = NULL, env = parent.frame(), variant = NULL,
  window = FALSE)
```

### Arguments

| | |
|---|---|
| ... | unevaluated expression to translate |
| tbl | An optional tbl. If supplied, will be used to automatically figure out the SQL variant to use. |
| env | environment in which to evaluate expression. |
| variant | used to override default variant provided by source useful for testing/examples |

| | |
|---|---|
| window | If variant not supplied, used to determine whether the variant is window based or not. |
| expr | list of quoted objects to translate |

### Base translation

The base translator, base_sql, provides custom mappings for ! (to NOT), && and & to AND, || and | to OR, ^ to POWER, %>% to %, ceiling to CEIL, mean to AVG, var to VARIANCE, tolower to LOWER, toupper to UPPER and nchar to length.

c and : keep their usual R behaviour so you can easily create vectors that are passed to sql.

All other functions will be preserved as is. R's infix functions (e.g. %like%) will be converted to their sql equivalents (e.g. LIKE). You can use this to access SQL string concatenation: || is mapped to OR, but %||% is mapped to ||. To suppress this behaviour, and force errors immediately when dplyr doesn't know how to translate a function it encounters, using set the dplyr.strict_sql option to TRUE.

You can also use sql to insert a raw sql string.

### SQLite translation

The SQLite variant currently only adds one additional function: a mapping from sd to the SQL aggregation function stdev.

### Examples

```
# Regular maths is translated in a very straightforward way
translate_sql(x + 1)
translate_sql(sin(x) + tan(y))

# Logical operators are converted to their sql equivalents
translate_sql(x < 5 & !(y >= 5))

# If is translated into select case
translate_sql(if (x > 5) "big" else "small")

# Infix functions are passed onto SQL with % removed
translate_sql(first %like% "Had*")
translate_sql(first %is% NULL)
translate_sql(first %in% c("John", "Roger", "Robert"))

# Note that variable names will be escaped if needed
translate_sql(like == 7)

# And be careful if you really want integers
translate_sql(x == 1)
translate_sql(x == 1L)

# If you have an already quoted object, use translate_sql_q:
x <- quote(y + 1 / sin(t))
translate_sql(x)
translate_sql_q(list(x))
```

```
# Translation with data source -------------------------------------------
## Not run:
flights <- tbl(nycflights13_sqlite(), "flights")
# Note distinction between integers and reals
translate_sql(month == 1, tbl = flights)
translate_sql(month == 1L, tbl = flights)

# Know how to translate most simple mathematical expressions
translate_sql(month %in% 1:3, tbl = flights)
translate_sql(month >= 1L & month <= 3L, tbl = flights)
translate_sql((month >= 1L & month <= 3L) | carrier == "AA", tbl = flights)

# Some R functions don't have equivalents in SQL: where possible they
# will be translated to the equivalent
translate_sql(xor(month <= 3L, carrier == "AA"), tbl = flights)

# Local variables will be automatically inserted into the SQL
x <- 5L
translate_sql(month == x, tbl = flights)

# By default all computation will happen in sql
translate_sql(month < 1 + 1, source = flights)
# Use local to force local evaluation
translate_sql(month < local(1 + 1), source = flights)

# This is also needed if you call a local function:
inc <- function(x) x + 1
translate_sql(month == inc(x), source = flights)
translate_sql(month == local(inc(x)), source = flights)

# Windowed translation ---------------------------------------------
planes <- arrange(group_by(flights, tailnum), desc(DepTime))

translate_sql(dep_time > mean(dep_time), tbl = planes, window = TRUE)
translate_sql(dep_time == min(dep_time), tbl = planes, window = TRUE)

translate_sql(rank(), tbl = planes, window = TRUE)
translate_sql(rank(dep_time), tbl = planes, window = TRUE)
translate_sql(ntile(dep_time, 2L), tbl = planes, window = TRUE)
translate_sql(lead(dep_time, 2L), tbl = planes, window = TRUE)
translate_sql(cumsum(dep_time), tbl = planes, window = TRUE)
translate_sql(order_by(dep_time, cumsum(dep_time)), tbl = planes, window = TRUE)

## End(Not run)
```

# Index