Data Transformation with dplyr Cheat Sheet



dplyr functions work with pipes and expect **tidy data**. In tidy data:



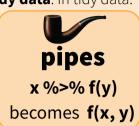
Each variable is











Summarise Cases

These apply **summary functions** to columns to create a new table. Summary functions take vectors as input and return one value (see back).





summarise (). summarise(mtcars, avg = mean(mpg))



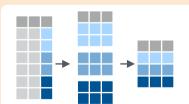
count(x, ..., wt = NULL, sort = FALSE) Count number of rows in each group defined by the variables in ... Also **tally()**. count(iris, Species)

Variations

- summarise_all() Apply funs to every column.
- summarise_at() Apply funs to specific columns.
- summarise if() Apply funs to all cols of one type.

Group Cases

Use **group_by()** to created a "grouped" copy of a table. dplyr functions will manipulate each "group" separately and then combine the results.



mtcars %>% group_by(cyl) %>% summarise(avg = mean(mpg))

group_by(.data, ..., add = FALSE) Returns copy of table grouped by ... g_iris <- group_by(iris, Species)</pre>

ungroup(x, ...)

Returns ungrouped copy of table. ungroup(g_iris)

Manipulate Cases

Extract Cases

Row functions return a subset of rows as a new table. Use a variant that ends in for non-standard evaluation friendly code.



filter(.data....)

Extract rows that meet logical criteria. Also **filter ()**. *filter(iris, Sepal.Length > 7)*



distinct(.data, ..., .keep all = FALSE) Remove rows with duplicate values. Also distinct_(). distinct(iris, Species)



sample frac(tbl, size = 1, replace = FALSE, weight = NULL, .env = parent.frame()) Randomly select fraction of rows. sample frac(iris, 0.5, replace = TRUE)

sample n(tbl, size, replace = FALSE, weight = NULL, .env = parent.frame()) Randomly select size rows. sample_n(iris, 10, replace = TRUE)



slice(.data, ...)

Select rows by position. Also **slice** (). *slice(iris, 10:15)*

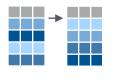
top_n(*x*, *n*, *wt***)**

Select and order top n entries (by group if grouped data). *top_n(iris, 5, Sepal.Width)*

Logical and boolean operators to use with filter()

is.na() <= %in% xor() >= !is.na() See ?base::logic and ?Comparison for help.

Arrange Cases



arrange(.data, ...)

Order rows by values of a column (low to high), use with **desc()** to order from high to low. arrange(mtcars, mpg) arrange(mtcars, desc(mpg))

Add Cases



add_row(.data, ..., .before = NULL, .after = NULL)

Add one or more rows to a table. add row(faithful, eruptions = 1, waiting = 1)

Manipulate Variables

Extract Variables

Column functions return a set of columns as a new table. Use a variant that ends in for non-standard evaluation friendly code.



select(.data....)

Extract columns by name. Also **select_if()** select(iris, Sepal.Length, Species)

Use these helpers with select(),

e.g. select(iris, starts with("Sepal"))

contains(match) ends with(match) matches(match)

num_range(prefix, range) one of(...)

:, e.g. mpg:cyl -, e.g, -Species everything()

Make New Variables

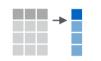
starts with(match)

These apply **vectorized functions** to columns. Vectorized funs take vectors as input and return vectors of the same length as output (see back).





Compute new column(s). mutate(mtcars, apm = 1/mpq)



transmute(.data, ...)

Compute new column(s), drop others. transmute(mtcars, qpm = 1/mpa)



mutate all(.tbl, .funs, ...)

Apply funs to every column. Use with funs(). mutate all(faithful, funs(log(.), log2(.)))



mutate_at(.tbl, .cols, .funs, ...)

Apply funs to specific columns. Use with funs(), vars() and the helper functions for

mutate at(iris, vars(-Species), funs(log(.)))

mutate_if(.tbl, .predicate, .funs, ...) Apply funs to all columns of one type. Use with funs().

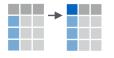
mutate if(iris, is.numeric, funs(log(.)))



add column(.data, ..., .before = NULL, .after = NULL)

Add new column(s).

add column(mtcars, new = 1:32)



rename(.data, ...)

Rename columns.

rename(iris, Length = Sepal.Length)

Vectorized Functions

to use with mutate()

mutate() and transmute() apply vectorized functions to columns to create new columns. Vectorized functions take vectors as input and return vectors of the same length as output.



Offsets

dplyr::lag() - Offset elements by 1 dplyr::lead() - Offset elements by -1

Cumulative Aggregates

dplyr::cumall() - Cumulative all() dplyr::cumany() - Cumulative any() cummax() - Cumulative max() dplyr::cummean() - Cumulative mean() cummin() - Cumulative min() cumprod() - Cumulative prod() cumsum() - Cumulative sum()

Rankings

dplyr::cume_dist() - Proportion of all values <=</pre> dplyr::dense rank() - rank with ties = min, no dplyr::min_rank() - rank with ties = min dplyr::ntile() - bins into n bins dplyr::percent_rank() - min_rank scaled to [0,1] dplyr::row number() - rank with ties = "first"

Math

+, -, *, /, ^, %/%, %% - arithmetic ops log(), log2(), log10() - logs <, <=, >, >=, !=, == - logical comparisons

Misc

dplyr::between() - x >= left & x <= right</pre> dplyr::case_when() - multi-case if_else() dplyr::coalesce() - first non-NA values by element across a set of vectors dplyr::if else() - element-wise if() + else() dplyr::na_if() - replace specific values with NA pmax() - element-wise max() pmin() - element-wise min() dplyr::recode() - Vectorized switch() dplyr::recode_factor() - Vectorized switch() for factors

Summary Functions

to use with summarise()

summarise() applies summary functions to columns to create a new table. Summary functions take vectors as input and return single values as output.



Counts

dplyr::n() - number of values/rows :n distinct() - # of uniques sum(!is.na()) - # of non-NA's

Location

mean() - mean. also mean(!is.na()) median() - median

Logicals

mean() - Proportion of TRUE's sum() - # of TRUE's

Position/Order

dplyr::first() - first value :last() - last value

dplyr::nth() - value in nth location of vector

Rank

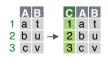
quantile() - nth quantile min() - minimum value max() - maximum value

Spread

IOR() - Inter-Quartile Range mad() - mean absolute deviation sd() - standard deviation var() - variance

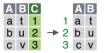
Row names

Tidy data does not use rownames, which store a variable outside of the columns. To work with the rownames, first move them into a column.



CAB rownames_to_column()

Move row names into col. a <- rownames_to_column(iris, *var* = "C")



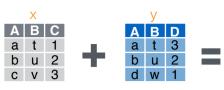
AB column_to_rownames()

Move col in row names. column_to_rownames(a, var = "C"

Also has_rownames(), remove_rownames()

Combine Tables

Combine Variables



Use **bind** cols() to paste tables beside each other as they are.



bind cols(...)

Returns tables placed side by side as a single table. BE SURE THAT ROWS ALIGN.

Use a "Mutating Join" to join one table to columns from another, matching values with the rows that they correspond to. Each join retains a different combination of values from the tables.



left_join(x, y, by = NULL, copy=FALSE, suffix=c(".x",".y"),...) Join matching values from v to x.



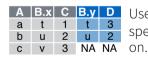
FALSE, suffix=c(".x",".y"),...) Join matching values from x to y.



A B C D inner_join(x, y, by = NULL, copy = FALSE, suffix=c(".x",".y"),...) Join data. Retain only rows with matches.

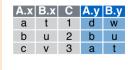


full_join(x, y, by = NULL, copy=FALSE, suffix=c(".x",".y"),...) Join data. Retain all values, all rows.



A B.x C B.y D Use by = c("col1", "col2") to a t 1 t 3 specify the column(s) to match specify the column(s) to match

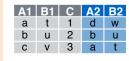
 $left_join(x, y, by = "A")$



A.x B.x C A.y B.y Use a named vector, by =

a t 1 d w c("col1" = "col2") to max c("col1" = "col2"), to match on columns with different names in each data set.

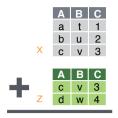
 $left_join(x, y, by = c("C" = "D"))$



A1 B1 C A2 B2 Use suffix to specify suffix to give to duplicate column names.

c v 3 a t left_join(x, y, by = c("C" = "D"), suffix = c("1", "2"))

Combine Cases



Use **bind rows()** to paste tables below each other as they are.



bind rows(...,.id = NULL)

Returns tables one on top of the other as a single table. Set .id to a column name to add a column of the original table names (as pictured)



intersect(x, y, ...)

Rows that appear in both x and z.



setdiff(x, y, ...)

Rows that appear in x but not z

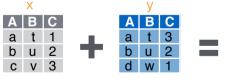


union(x, y, ...)

Rows that appear in x or z. (Duplicates removed). union_all() retains duplicates.

Use **setequal()** to test whether two data sets contain the exact same rows (in any order).

Extract Rows



Use a "Filtering Join" to filter one table against the rows of another.



semi join(x, y, by = NULL, ...)

Return rows of x that have a match in v. USEFUL TO SEE WHAT WILL BE JOINED



anti_join(x, y, by = NULL, ...)

Return rows of x that do not have a match in y. USEFUL TO SEE WHAT WILL NOT BE JOINED.