

rtables - Reporting tables with R : : CHEAT SHEET



Basics

The **rtables** R package is designed to create and display complex tables with R.

Every rtable layout is constructed starting with **basic_table** and is rendered using **build_table**.

CODE

```
tbl_a <- basic_table() %>%
  split_cols_by("ARM") %>%
  split_rows_by("STRATA1") %>%
  analyze("AGE") %>%
  build_table(ads1)
```

TABLE OUTPUT

| | ARM X | ARM Y |
|------|-------|-------|
| A | | |
| Mean | 33.32 | 35.86 |
| B | | |
| Mean | 33.65 | 38.00 |

Layout & Tabulation

ANALYZE & SUMMARIZE FUNCTIONS

analyze()
analyze_colvars()
summarize_row_groups()

LAYOUT MODIFIERS

append_topleft()
add_colcounts()
add_overall_col()

CUSTOMIZED TABLE CODE

```
basic_table(show_colcounts = TRUE) %>%
  split_cols_by("ARM") %>%
  add_overall_col("TOTAL") %>%
  split_rows_by("BMRKR2",
    split_label = "Biomarker 2 Level",
    label_pos = "topleft") %>%
  summarize_row_groups() %>%
  analyze("AGE", var_labels = "Age (yrs)",
    afun = mean, format = "xx.x") %>%
  analyze("STRATA1", var_labels = "Stratif. Term",
    afun = function(x, .N_col) lapply(
      table(x),
      function(xi) rcell(
        xi * c(1, 1 / .N_col),
        format = "xx (xx.xx%)"
      ))) %>%
  append_topleft("Attribute") %>%
  build_table(ads1)
```

CUSTOMIZED TABLE OUTPUT

| Biomarker 2 Level Attribute | ARM X (N=42) | ARM Y (N=40) | TOTAL (N=82) |
|--------------------------------|-----------------|-----------------|-----------------|
| LOW | 22 (52.4%) | 25 (62.5%) | 47 (57.3%) |
| Age (yrs) | | | |
| mean | 33.5 | 36.4 | 35.1 |
| Stratif. Term | | | |
| A | 9 (21.43%) | 12 (30.00%) | 21 (25.61%) |
| B | 13 (30.95%) | 13 (32.50%) | 26 (31.71%) |
| HIGH | 20 (47.6%) | 15 (37.5%) | 35 (42.7%) |
| Age (yrs) | | | |
| mean | 33.5 | 37.7 | 35.3 |
| Stratif. Term | | | |
| A | 10 (23.81%) | 9 (22.50%) | 19 (23.17%) |
| B | 10 (23.81%) | 6 (15.00%) | 16 (19.51%) |

Customization Options

ANALYZE & SUMMARIZE FUNCTIONS

| Argument | Input | Effect on Table |
|--------------------|-------------------------------------|---|
| afun/cfun | Analysis function | The function is used to calculate cell values |
| var_labels | Labels for variables being analyzed | Labels are printed in the leftmost column |
| format | Format string or function | Format is applied to render cell values |
| na_str | String to represent NA values | String is printed in place of missing values |
| inclNAs | TRUE or FALSE | Changes whether records with NA are included in analysis |
| show_labels | "default", "visible", or "hidden" | var_labels are printed or hidden in the table |
| indent_mod | Number of spaces to indent by | Current analysis rows are indented |
| section_div | String to divide split sections by | String is printed between groups defined by current split |

Simple Tabulation

Quick tables with **qtable** – an extension of **base::table** for exploratory work & data summarization.

CODE

```
qtable(
  ads1,
  row_vars = c(
    "STRATA1", "STRATA2"
  ),
  col_vars = c("ARM"),
  avar = "AGE",
  afun = mean
)
```

TABLE OUTPUT

| AGE - mean | ARM X (N=42) | ARM Y (N=40) |
|------------|-----------------|-----------------|
| A | | |
| X | 33.00 | 33.44 |
| Z | 34.33 | 39.25 |
| Y | 32.75 | 34.50 |
| B | | |
| X | 34.29 | 34.50 |
| Z | 26.25 | 47.50 |
| Y | 35.75 | 36.57 |

Titles & Footers

CODE

```
main_title(tbl_a) <- "My Title"
subtitles(tbl_a) <- c("A subtitle")
main_footer(tbl_a) <- c("A footnote")
prov_footer(tbl_a) <- c("A provenance footer")
fnotes_at_path(tbl_a,
  rowpath = c("STRATA1", "A", "AGE", "Mean"),
  colpath = c("ARM", "ARM X")
) <- "Mean age for arm X"
```

TABLE OUTPUT

| | | |
|--------------------------|-----------|-------|
| My Title | | |
| A subtitle | | |
| | ARM X | ARM Y |
| A | | |
| Mean | 33.32 {1} | 35.86 |
| B | | |
| Mean | 33.65 | 38.00 |
| {1} - Mean age for arm X | | |
| A footnote | | |
| A provenance footer | | |



Split Functions

Split functions are used to **add, remove, or transform** the levels of the variable used in a split.

ROW SPLITS

`split_rows_by()`
`split_rows_by_multivar()`
`split_rows_by_cuts()`
`split_rows_by_cutfun()`
`split_rows_by_quartiles()`

COLUMN SPLITS

`split_cols_by()`
`split_cols_by_multivar()`
`split_cols_by_cuts()`
`split_cols_by_cutfun()`
`split_cols_by_quartiles()`

SPLIT FUNCTIONS

`remove_split_levels()` `add_overall_level()` `trim_levels_in_group()`
`keep_split_levels()` `add_combo_levels()` `trim_levels_to_map()`
`drop_split_levels()` `reorder_split_levels()`
`drop_and_remove_levels()`

CODE

```
basic_table() %>%  
  split_cols_by(  
    "ARM",  
    split_fun = remove_split_levels(c("ARM Y"))  
  ) %>%  
  split_rows_by(  
    "STRATA1",  
    split_fun = reorder_split_levels(c("B", "A"))  
  ) %>%  
  analyze("AGE") %>%  
  build_table(ads1)
```

TABLE OUTPUT

| | | ARM X |
|---|------|-------|
| B | Mean | 33.65 |
| A | Mean | 33.32 |

For information on custom split functions, see [?custom_split_funs](#)

Sorting & Pruning

Sorting functions are used to **reorder table rows** according to a given criteria function. Pruning functions are used to **remove table rows** according to a given criteria.

SORTING

`sort_at_path()`
`cont_n_allcols()`
`cont_n_onecol()`

PRUNING

`prune_table()`
`all_zero_or_na()`
`all_zero()`
`content_all_zeros_nas()`

`prune_empty_level()`
`prune_zeros_only()`
`low_obs_pruner()`

```
tbl <- basic_table() %>%  
  split_cols_by("ARM") %>%  
  split_rows_by("STRATA2") %>%  
  summarize_row_groups() %>%  
  build_table(ads1)
```

| | ARM X | ARM Y |
|---|------------|------------|
| X | 19 (45.2%) | 19 (47.5%) |
| Y | 23 (54.8%) | 21 (52.5%) |
| Z | 0 (0.0%) | 0 (0.0%) |

```
tbl %>%  
  sort_at_path(  
    "STRATA2",  
    scorefun = cont_n_allcols  
  ) %>%  
  prune_table()
```

| | ARM X | ARM Y |
|---|------------|------------|
| Y | 23 (54.8%) | 21 (52.5%) |
| X | 19 (45.2%) | 19 (47.5%) |

Access & Modify

ACCESSORS

`head(tbl, n)` `cell_values(tbl, rowpath, colpath)`
`tail(tbl, n)` `value_at(tbl, rowpath, colpath)`
`tbl[x, y]` `top_left(tbl)`

MODIFIERS

`tbl[x, y] <- rcell(...)` `rbind(tbl_1, tbl_2)`
`top_left(tbl) <- "XXX"` `cbind_rtables(tbl_1, tbl_2)`

`head(tbl_a, 2)`

| | ARM X | ARM Y |
|---|-------|-------|
| A | Mean | 33.32 |
| | | 35.86 |

`tail(tbl_a, 2)`

| | ARM X | ARM Y |
|---|-------|-------|
| B | Mean | 33.65 |
| | | 38.00 |

`tbl_a[3:4, 1]`

| | ARM X |
|---|-------|
| B | Mean |
| | 33.65 |

```
cell_values(  
  tbl_a, rowpath = c(  
    "STRATA1", "A", "AGE", "Mean"  
  ),  
  colpath = c("ARM", "ARM Y")  
)
```

`$`ARM Y`
[1] 35.86`

Rendering

rtables prints output in ASCII format in the R console.

rtable objects can also be paginated or converted to different output types in the console, or exported to various file types.

R SESSION OUTPUT

`Viewer(tbl)`
`toString(tbl)`
`as_html(tbl)`
`tt_to_flextable(tbl)`

PAGINATION

```
paginate_table(  
  tbl,  
  page_type = "letter",  
  font_family = "Courier",  
  font_size = 8,  
  landscape = FALSE  
)
```

EXPORT



`export_as_docx(tbl, "tbl.docx")`
`export_as_pdf(tbl, "tbl.pdf")`
`export_as_rtf(tbl, "tbl.rtf")`
`export_as_tsv(tbl, "tbl.tsv")`
`export_as_txt(tbl, "tbl.txt")`

