

An Introduction to Tablet for PDF

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Motivation

Occasionally it is useful to generate a table of summary statistics for rows of a dataset, where such rows represent sampling units and columns may be categorical or continuous. The excellent R package [table1](#) does exactly this, and was the inspiration for **tablet**. **table1** however is optimized for html; **tablet** tries to provide a format-neutral implementation and relies on [kableExtra](#) to handle the rendering. Support for pdf (latex) is of particular interest, and is illustrated here. See the companion vignette for a proof-of-concept html implementation.

Software

To support our examples, we load some other packages and in particular locate the melanoma dataset from [boot](#). By the way, in the yaml header for the Rmd source file, we've added the header-includes as described on p. 4 of the [kableExtra documentation](#).

```
library(tidyr)
library(dplyr)
library(magrittr)
library(kableExtra)
library(boot)
library(yamlet)
library(tablet)
```

```
x <- melanoma
x %<>% select(-time, -year)
```

Simple Case

For starters, we'll just coerce two variables to factor to show that they are categorical, and then pass the whole thing to `tablet()`. Then we forward to `as_kable()` for rendering (calls `kableExtra::kbl` and adds some magic).

```
x %>%
  mutate(
    sex = factor(sex),
    ulcer = factor(ulcer)
  ) %>%
  tablet %>%
  as_kable
```

| All (N = 205) | |
|------------------|------------------|
| status | |
| Mean (SD) | 1.79 (0.551) |
| Median (range) | 2 (1, 3) |
| sex | |
| 0 | 126 (61.5%) |
| 1 | 79 (38.5%) |
| age | |
| Mean (SD) | 52.5 (16.7) |
| Median (range) | 54 (4, 95) |
| thickness | |
| Mean (SD) | 2.92 (2.96) |
| Median (range) | 1.94 (0.1, 17.4) |
| ulcer | |
| 0 | 115 (56.1%) |
| 1 | 90 (43.9%) |

With Metadata

Now we redefine the dataset, supplying metadata almost verbatim from `?melanoma`. This is fairly easy using package `yamlet`. Note that we reverse the authors' factor order of 1, 0 for ulcer and move status 'Alive' to first position.

```
x <- melanoma

x %<>% decorate('
time:      [ Survival Time Since Operation, day ]
status:
- End of Study Patient Status
-
- Alive: 2
- Melanoma Death: 1
- Unrelated Death: 3
sex:       [ Sex, [ Male: 1, Female: 0 ] ]
age:       [ Age at Time of Operation, year ]
year:      [ Year of Operation, year ]
thickness: [ Tumor Thickness, mm ]
ulcer:     [ Ulceration, [ Absent: 0, Present: 1 ] ]
')
x %<>% select(-time, -year)
x %<>% group_by(status)
x %<>% resolve
```

- `group_by(status)` causes statistics to be summarized in columns by group.
- `resolve()` disambiguates labels, units, and factor levels (actually creating factors where appropriate, such as for sex and ulcer).

Now we pass `x` to `tablet()` and `as_kable()` for a more informative result.

```
x %>% tablet %>% as_kable
```

| | Alive (N = 134) | Melanoma Death (N = 57) | Unrelated Death (N = 14) | All (N = 205) |
|--|--------------------|----------------------------|-----------------------------|------------------|
| Sex | | | | |
| Male | 43 (32.1%) | 29 (50.9%) | 7 (50%) | 79 (38.5%) |
| Female | 91 (67.9%) | 28 (49.1%) | 7 (50%) | 126 (61.5%) |
| Age at Time of Operation (year) | | | | |
| Mean (SD) | 50 (15.9) | 55.1 (17.9) | 65.3 (10.9) | 52.5 (16.7) |
| Median (range) | 52 (4, 84) | 56 (14, 95) | 65 (49, 86) | 54 (4, 95) |
| Tumor Thickness (mm) | | | | |
| Mean (SD) | 2.24 (2.33) | 4.31 (3.57) | 3.72 (3.63) | 2.92 (2.96) |
| Median (range) | 1.36 (0.1, 12.9) | 3.54 (0.32, 17.4) | 2.26 (0.16, 12.6) | 1.94 (0.1, 17.4) |
| Ulceration | | | | |
| Absent | 92 (68.7%) | 16 (28.1%) | 7 (50%) | 115 (56.1%) |
| Present | 42 (31.3%) | 41 (71.9%) | 7 (50%) | 90 (43.9%) |

Notice that:

- the order of variables down the left side is exactly their order in the dataset;
- the order of factor levels is exactly that in `x`;
- the order of groups across the top is exactly the levels (if any) of the grouping variable(s), and
- labels and titles (highest priority) are substituted for item names.

If you don't particularly care for some aspect of the presentation, you can jump in between `tablet()` and `as_kable()` to fix things up. For example, if you don't want the "All" column you can just say

- `x %>% tablet %>% select(-All) %>% as_kable.`

If you **only** want the the "All" column, you can just remove the group(s):

- `x %>% ungroup %>% select(-1) %>% tablet %>% as_kable.`

By the way, you can also pass `all = NULL` to suppress the 'All' column.

As Xtable

Some support is provided for 'xtable'. Currently, grouped columns (see next section) are not supported.

```
library(xtable)
x %>%
  filter(!(status == 'Alive' & sex == 'Male')) %>%
  tablet %>% as_xtable(format_value = function(x,...)x) %>%
  print(
    booktabs = TRUE,
    include.rownames = FALSE
  )
```

| | Alive (N = 91) | Melanoma Death (N = 57) | Unrelated Death (N = 14) | All (N = 162) |
|--|------------------|-------------------------|--------------------------|------------------|
| Age at Time of Operation (year) | | | | |
| Mean (SD) | 48.8 (15.4) | 55.1 (17.9) | 65.3 (10.9) | 52.4 (16.7) |
| Median (range) | 49 (4, 77) | 56 (14, 95) | 65 (49, 86) | 54 (4, 95) |
| Sex | | | | |
| Male | | 29 (50.9%) | 7 (50%) | 36 (22.2%) |
| Female | 91 (100%) | 28 (49.1%) | 7 (50%) | 126 (77.8%) |
| Tumor Thickness (mm) | | | | |
| Mean (SD) | 2.02 (2.22) | 4.31 (3.57) | 3.72 (3.63) | 2.97 (3.08) |
| Median (range) | 1.29 (0.1, 12.9) | 3.54 (0.32, 17.4) | 2.26 (0.16, 12.6) | 1.94 (0.1, 17.4) |
| Ulceration | | | | |
| Absent | 68 (74.7%) | 16 (28.1%) | 7 (50%) | 91 (56.2%) |
| Present | 23 (25.3%) | 41 (71.9%) | 7 (50%) | 71 (43.8%) |

Grouped Columns

In `tablet()`, most columns are the consequences of a grouping variable. Not surprisingly, grouped columns are just a consequence of nested grouping variables. To illustrate, we follow the [table1 vignette](#) by adding a grouping variable that groups the two kinds of death.

| | Alive | | Melanoma | | Unrelated | | All |
|--|-------------------|--------------------|-------------------|--------------------|-------------------|-------------------|------------------|
| | Male (N = 43) | Female (N = 91) | Male (N = 29) | Female (N = 28) | Male (N = 7) | Female (N = 7) | (N = 205) |
| Age at Time of Operation (year) | | | | | | | |
| Mean (SD) | 52.5 (16.9) | 48.8 (15.4) | 53.9 (19.7) | 56.4 (16.2) | 62.4 (11.2) | 68.1 (10.6) | 52.5 (16.7) |
| Median (range) | 55 (12, 84) | 49 (4, 77) | 52 (19, 95) | 58 (14, 89) | 64 (49, 76) | 66 (54, 86) | 54 (4, 95) |
| Tumor Thickness (mm) | | | | | | | |
| Mean (SD) | 2.73 (2.49) | 2.02 (2.22) | 4.63 (3.47) | 3.99 (3.71) | 4.83 (4.19) | 2.6 (2.84) | 2.92 (2.96) |
| Median (range) | 1.62 (0.16, 8.38) | 1.29 (0.1, 12.9) | 4.04 (0.81, 14.7) | 3.14 (0.32, 17.4) | 4.84 (0.65, 12.6) | 1.45 (0.16, 8.54) | 1.94 (0.1, 17.4) |
| Ulceration | | | | | | | |
| Absent | 24 (55.8%) | 68 (74.7%) | 8 (27.6%) | 8 (28.6%) | 4 (57.1%) | 3 (42.9%) | 115 (56.1%) |
| Present | 19 (44.2%) | 23 (25.3%) | 21 (72.4%) | 20 (71.4%) | 3 (42.9%) | 4 (57.1%) | 90 (43.9%) |

```
x %<>% mutate(class = status)           # copy the current group
x %<>% modify(class, label = 'class')     # change its label
levels(x$status) <- c('Alive', 'Melanoma', 'Unrelated') # tweak current group
levels(x$class) <- c(' ', 'Death', 'Death') # cluster groups
x %<>% group_by(class, status)           # nest groups
x %>% tablet %>% as_kable               # render
```

| | Alive (N = 134) | Death | | All (N = 205) |
|--|--------------------|----------------------|-----------------------|------------------|
| | | Melanoma (N = 57) | Unrelated (N = 14) | |
| Sex | | | | |
| Male | 43 (32.1%) | 29 (50.9%) | 7 (50%) | 79 (38.5%) |
| Female | 91 (67.9%) | 28 (49.1%) | 7 (50%) | 126 (61.5%) |
| Age at Time of Operation (year) | | | | |
| Mean (SD) | 50 (15.9) | 55.1 (17.9) | 65.3 (10.9) | 52.5 (16.7) |
| Median (range) | 52 (4, 84) | 56 (14, 95) | 65 (49, 86) | 54 (4, 95) |
| Tumor Thickness (mm) | | | | |
| Mean (SD) | 2.24 (2.33) | 4.31 (3.57) | 3.72 (3.63) | 2.92 (2.96) |
| Median (range) | 1.36 (0.1, 12.9) | 3.54 (0.32, 17.4) | 2.26 (0.16, 12.6) | 1.94 (0.1, 17.4) |
| Ulceration | | | | |
| Absent | 92 (68.7%) | 16 (28.1%) | 7 (50%) | 115 (56.1%) |
| Present | 42 (31.3%) | 41 (71.9%) | 7 (50%) | 90 (43.9%) |

Transposed Groups

Categorical observations (in principle) and grouping variables are all factors, and are thus transposable. To illustrate, we drop the column group above and instead nest sex within status ...

```
x %<>% group_by(status, sex)
x %<>% select(-class)
x %>%
  tablet %>%
  as_kable %>%
  kable_styling(latex_options = 'scale_down')
```

... or nest ulceration within status ...

| | Alive | | Melanoma | | Unrelated | | All (N = 205) |
|--|--------------------|---------------------|--------------------|---------------------|-------------------|--------------------|------------------|
| | Absent (N = 92) | Present (N = 42) | Absent (N = 16) | Present (N = 41) | Absent (N = 7) | Present (N = 7) | |
| Sex | | | | | | | |
| Male | 24 (26.1%) | 19 (45.2%) | 8 (50%) | 21 (51.2%) | 4 (57.1%) | 3 (42.9%) | 79 (38.5%) |
| Female | 68 (73.9%) | 23 (54.8%) | 8 (50%) | 20 (48.8%) | 3 (42.9%) | 4 (57.1%) | 126 (61.5%) |
| Age at Time of Operation (year) | | | | | | | |
| Mean (SD) | 49.3 (15.4) | 51.6 (17.1) | 54.9 (19.9) | 55.1 (17.4) | 58.4 (8.66) | 72.1 (8.53) | 52.5 (16.7) |
| Median (range) | 50 (4, 83) | 54.5 (12, 84) | 59 (16, 83) | 56 (14, 95) | 56 (49, 71) | 72 (60, 86) | 54 (4, 95) |
| Tumor Thickness (mm) | | | | | | | |
| Mean (SD) | 1.63 (1.93) | 3.58 (2.58) | 2.7 (3.35) | 4.94 (3.5) | 2.1 (1.93) | 5.34 (4.33) | 2.92 (2.96) |
| Median (range) | 1.13 (0.1, 12.9) | 3.06 (0.32, 12.2) | 1.94 (0.32, 14.7) | 4.04 (0.97, 17.4) | 1.45 (0.65, 6.12) | 4.84 (0.16, 12.6) | 1.94 (0.1, 17.4) |

| | Alive | | | | Melanoma | | | | Unrelated | | | | All (N = 205) | |
|---------------------------------|-------------------|--------------------|-------------------|--------------------|-------------------|-------------------|-------------------|--------------------|-------------------|-------------------|-------------------|-------------------|------------------|--|
| | Absent | | Present | | Absent | | Present | | Absent | | Present | | | |
| | Male (N = 24) | Female (N = 68) | Male (N = 19) | Female (N = 23) | Male (N = 8) | Female (N = 8) | Male (N = 21) | Female (N = 20) | Male (N = 4) | Female (N = 3) | Male (N = 3) | Female (N = 4) | | |
| | | | | | | | | | | | | | | |
| Age at Time of Operation (year) | | | | | | | | | | | | | | |
| Mean (SD) | 50.4 (17) | 48.9 (14.9) | 55.3 (16.9) | 48.7 (17) | 55.2 (22.2) | 54.6 (18.8) | 53.3 (19.2) | 57 (15.5) | 54.5 (7.14) | 63.7 (8.74) | 73 (2.65) | 71.5 (11.8) | 52.5 (16.7) | |
| Median (range) | 54 (15, 83) | 49 (4, 77) | 56 (12, 84) | 48 (19, 75) | 56 (27, 83) | 59 (16, 77) | 52 (19, 95) | 58 (14, 89) | 52.5 (49, 64) | 66 (54, 71) | 72 (71, 76) | 70 (60, 86) | 54 (4, 95) | |
| Tumor Thickness (mm) | | | | | | | | | | | | | | |
| Mean (SD) | 1.47 (1.72) | 1.69 (2) | 4.32 (2.42) | 2.97 (2.59) | 3.27 (4.68) | 2.14 (1.18) | 5.14 (2.86) | 4.72 (4.13) | 2.42 (2.5) | 1.67 (1.14) | 8.05 (4.02) | 3.3 (3.71) | 2.92 (2.96) | |
| Median (range) | 0.97 (0.16, 7.09) | 1.29 (0.1, 12.9) | 3.87 (0.81, 8.38) | 1.94 (0.32, 12.2) | 1.78 (0.81, 14.7) | 2.02 (0.32, 3.56) | 4.83 (1.62, 12.9) | 3.54 (0.97, 17.4) | 1.46 (0.65, 6.12) | 1.45 (0.65, 2.9) | 6.76 (4.84, 12.6) | 2.26 (0.16, 8.54) | 1.94 (0.1, 17.4) | |

```
x %<>% group_by(status, ulcer)
x %>%
  tablet %>%
  as_kable %>%
  kable_styling(latex_options = 'scale_down')
```

... or where it makes sense, use multiple levels of nesting.

```
x %<>% group_by(status, ulcer, sex)
x %>%
  tablet %>%
  as_kable %>%
  kable_styling(latex_options = 'scale_down') # %>% landscape ?
```

Aesthetics

`tablet` tries to give rather exhaustive control over formatting. Much can be achieved by replacing elements of ‘fun’, ‘fac’, ‘num’, and ‘lab’ (see `?tablet.data.frame`). For finer control, you can replace these entirely. In this example, we will ...

- ignore categoricals (other than groups) by replacing ‘fac’ with something of length zero,
- drop the ‘N =’ header material by substituting in ‘lab’, and
- switch to ‘(min - max)’ instead of ‘(min, max)’.

```
x %<>% group_by(status)
x %>%
  tablet(
    fac = NULL,
    lab ~ name,
    `Median (range)` ~ med + ' (' + min + ' - ' + max + ')'
  ) %>%
  as_kable
```

| | Alive | Melanoma | Unrelated | All |
|--|-------------------|--------------------|--------------------|-------------------|
| Age at Time of Operation (year) | | | | |
| Mean (SD) | 50 (15.9) | 55.1 (17.9) | 65.3 (10.9) | 52.5 (16.7) |
| Median (range) | 52 (4 - 84) | 56 (14 - 95) | 65 (49 - 86) | 54 (4 - 95) |
| Tumor Thickness (mm) | | | | |
| Mean (SD) | 2.24 (2.33) | 4.31 (3.57) | 3.72 (3.63) | 2.92 (2.96) |
| Median (range) | 1.36 (0.1 - 12.9) | 3.54 (0.32 - 17.4) | 2.26 (0.16 - 12.6) | 1.94 (0.1 - 17.4) |

Note

The default presentation includes “N =” under the header, but also has percent characters in the table. Considerable gymnastics are required to make this work! If you change the defaults you may want to consider the arguments to `?as_kable.tablet`.

- ‘linebreaker’ anticipates that ‘lab’ has inserted newlines before “N =”.
- ‘linebreak’ invokes the eponymous `kableExtra` function internally.
- ‘escape’ is turned off to protect resulting latex markup.
- ‘escape_latex’ is supplied to restore the usual escaping of percent characters.

Conclusion

`tablet` gives a flexible way of summarizing tables of observations. It reacts to numeric columns, factors, and grouping variables. Display order derives from the order of columns and factor levels in the data. Result columns can be grouped arbitrarily deep by supplying extra groups. Column labels and titles are respected. Rendering is largely the responsibility of `kableExtra` and can be extended. Further customization is possible by manipulating data after calling `tablet()` but before calling `as_kable()`. Powerful results are possible with very little code.