Test

Tim Bergsma

2022-04-16

Contents

```
library(tablet)
library(haven)
library(yamlet)
##
## Attaching package: 'yamlet'
## The following object is masked from 'package:stats':
##
##
       filter
library(magrittr)
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
library(kableExtra)
##
## Attaching package: 'kableExtra'
## The following object is masked from 'package:dplyr':
##
##
       group_rows
## The following object is masked from 'package:yamlet':
##
##
       footnote
```

```
library(knitr)
# make adsl with imputed bmi, imputed race, and two-row footnote
d <- read_sas('adsl.sas7bdat')</pre>
m <- read_yamlet('adsl.yaml')</pre>
# fortify to mimic app.R
have <- names(d)
need <- names(m)</pre>
make <- setdiff(need, have)</pre>
for(col in make) d[[col]] <- rep(NA_integer_, nrow(d))</pre>
# ensure positive nrow
if(nrow(d) == 0) d <- d['',,drop = FALSE]</pre>
# drop unspecified
d %<>% select(!!!names(m))
# apply meta
d <- redecorate(d, m)</pre>
# # Promote NA to a level of the factor
# d %<>% resolve(exclude = NULL)
d %<>% resolve()
## Warning in match.fun(test)(val, data = data[[i]], ...): data has values not in
## Placebo, TRT 10 mg, TRT 20 mg: e.g.
## Warning in match.fun(test)(val, data = data[[i]], ...): data has values not in
## Placebo, TRT 10 mg, TRT 20 mg, TRT Total: e.g.
## Warning in match.fun(test)(val, data = data[[i]], ...): data has values not in
## White, Black, Asian, Other, : e.g. NA
foot <-
'a clinicaltrial.gov
b some other comment'
options(knitr.kable.NA = 0)
#opts_knit$set(out.format = 'latex')
# debug(tablet:::widgets.devalued)
d %>% group_by(trt01a, trt01aa) %>% select(race) %>% tablet
## Adding missing grouping variables: 'trt01a', 'trt01aa'
## [1] _tablet_name _tablet_level _tablet_sort _tablet_stat All
## <0 rows> (or 0-length row.names)
t <- d %>%
  filter(saffl == 'Y') %>%
  group_by(trt01a, trt01aa) %>%
  select(
```

```
# age, agegr, sex, weight, bmi,
   race
  ) %>%
  tablet(
   all levels = TRUE,
   fun = list(
     sum ~ sum(x, na.rm = TRUE),
     mean(x, na.rm = TRUE)),
     ave ~ signif(digits = 3,
     std ~ signif(digits = 3,
                                sd(x, na.rm = TRUE)),
     med ~ signif(digits = 3, median(x, na.rm = TRUE)),
     min ~ signif(digits = 3, min(x, na.rm = TRUE)),
     max ~ signif(digits = 3,
                               max(x, na.rm = TRUE)),
     smn ~ sum(!is.na(x))
   ),
   num = list(
     n ~ smn,
     `Mean (SD)` ~ ave + ' (' + std + ')',
    Median ~ paste(med),
     `Min, Max` ~ min + ', ' + max
   ),
   fac = list(
     ` ~ ifelse(sum == 0, '0', sum + ' (' + pct + '\", ' + ')')
   )
  )
## Adding missing grouping variables: 'trt01a', 'trt01aa'
## Joining, by = c("trt01a", "trt01aa", "_tablet_N", "_tablet_n", "_tablet_name", "_tablet_level")
## Joining, by = c("_tablet_N", "_tablet_n", "_tablet_name", "_tablet_level")
# reverse lookup on
imputed <- sapply(select(d, !!!make), attr, 'label')</pre>
#to substitute '-' for all imputeds
# t \leftarrow t[names(t)[!duplicated(names(t))]]
# t %<>% mutate(
     .cols = -starts_with('_tablet_'),
     .fns = ~ ifelse(`_tablet_name` %in% imputed, '-', .x)
#
# )
t %>%
 as kable %>%
 footnote(
   general = # escape_latex(
     c('a something','b something')
     # )
   fixed_small_size = TRUE,
   general_title = " ",
```

threeparttable = TRUE) # %>% as.character %>% writeLines

	Placebo	TRT 10 mg	TRT 20 mg	
		$ \begin{array}{c} \text{TRT Total} \\ (N = 4) \end{array} $	$ \begin{array}{c} \text{TRT Total} \\ (N = 4) \end{array} $	All (N = 12)
Race				
White	0	0	0	0
Black	0	0	0	0
Asian	0	0	0	0
Other	0	0	0	0
Missing	0	0	0	0

a something b something