Test

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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
x <- read_sas('adsl.sas7bdat')</pre>
m <- read_yamlet('adsl.yaml')</pre>
# https://github.com/haozhu233/kableExtra/issues/703
names(m$race$guide)[[3]] <- 'Oriental'</pre>
# fortify to mimic app.R
have <- names(x)
need <- names(m)</pre>
make <- setdiff(need, have)</pre>
for(col in make) x[[col]] <- rep(NA_integer_, nrow(x))</pre>
# ensure positive nrow
if(nrow(x) == 0) x \leftarrow x['', drop = FALSE]
# drop unspecified
x %<>% select(!!!names(m))
# apply meta
x <- redecorate(x, m)
# # Promote NA to a level of the factor
# x %<>% resolve(exclude = NULL)
x %<>% 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$set(out.format = 'latex')
# debug(tablet:::widgets.devalued)
#x %>% group_by(trt01a, trt01aa) %>% select(race) %>% tablet
# x$trt01a[] <- NA
# x$trt01aa[] <- NA
# debug(categoricals)
x <- x %>%
 filter(saffl == 'Y') %>%
  group_by(trt01a, trt01aa) %>%
  select(
  # age, agegr, sex, weight, bmi,
```

```
race, bmi
  ) %>%
  tablet(
   all levels = TRUE,
   fun = list(
     sum ~ sum(x, na.rm = TRUE),
     ave ~ signif(digits = 3, mean(x, na.rm = TRUE)),
     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 make
codelist <- attr(x$`_tablet_name`, 'codelist')</pre>
x$`_tablet_original` <- unlist(codelist[x$`_tablet_name`])</pre>
# very elegant, but blows away attributes
# x %<>% mutate(
     .cols = -starts_with('_tablet_'),
#
      .fns = ~ ifelse(`_tablet_original` %in% names(conf$imputed), '-', .x)
#
# )
targets <- seq_along(x)[!(grepl('_tablet_', names(x)))]</pre>
imputed <- x$`_tablet_original` %in% make</pre>
if(length(imputed) & length(targets)) x[imputed, targets] <- '-'</pre>
x$`_tablet_original` <- NULL
x %>%
 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} $	$ \begin{array}{c} \text{TRT Total} \\ (N = 4) \end{array} $		All (N = 12)
Race				
White	-	-	-	
Black	-	-	-	-
Oriental	-	-	-	-
Other	-	-	-	-
Missing	-	_	_	-
Body Mass Index				
n	-	-	-	-
Mean (SD)	-	_	_	-
Median	-	-	-	-
Min, Max	-	_	_	-

a something b something