

ec_report

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# reference: https://cran.r-project.org/web/packages/qwraps2/vignettes/summary-statistics.html
library(dplyr)
library(qwraps2)
options(qwraps2_markup = "markdown")
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```
dig = read.csv('dig.csv')
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```
our_summary1 <-
  list("CHARACTERISTIC" =
    list("Age (yr) - mean  $\pm$  SD" = ~ qwraps2::mean_sd(AGE, digits = getOption("qwraps2_fmt_digits", 1), na_rm = TRUE),
          "Ejection fraction - mean  $\pm$  SD" = ~ qwraps2::mean_sd(EJF_PER, digits = getOption("qwraps2_fmt_digits", 1), na_rm = TRUE),
          "Median duration of CHF - mo" = ~ median(CHFDUR, digits=1, na.rm = TRUE),
          "Female sex" = ~ perc_n(SEX == 2, digits=1, na_rm = TRUE),
          "Nonwhite race" = ~ perc_n(RACE == 2, digits=1, na_rm = TRUE),
          "Age>70 yr" = ~ perc_n(AGE > 70, digits=1, na_rm = TRUE)),
    "Method of assessing ejection fraction" =
    list("Radionuclide ventriculography" = ~ perc_n(EJFMETH == 1, digits=1, na_rm = TRUE),
          "Two-dimensional echocardiography" = ~ perc_n(EJFMETH == 3, digits=1, na_rm = TRUE),
          "Contrast angiography" = ~ perc_n(EJFMETH == 2, digits=1, na_rm = TRUE),
          "Cardiothoracic ratio>0.55" = ~ perc_n(CHESTX>0.55, digits=1, na_rm = TRUE)),
    "NYHA class" =
    list("I" = ~ perc_n(FUNCTCLS == 1, digits=1, na_rm = TRUE),
          "II" = ~ perc_n(FUNCTCLS == 2, digits=1, na_rm = TRUE),
          "III" = ~ perc_n(FUNCTCLS == 3, digits=1, na_rm = TRUE),
          "IV" = ~ perc_n(FUNCTCLS == 4, digits=1, na_rm = TRUE)),
    "No. of signs or symptoms of CHF" =
    list("0" = ~ perc_n(NSYM == 0, digits=1, na_rm = TRUE),
          "1" = ~ perc_n(NSYM == 1, digits=1, na_rm = TRUE),
          "2" = ~ perc_n(NSYM == 2, digits=1, na_rm = TRUE),
          "3" = ~ perc_n(NSYM == 3, digits=1, na_rm = TRUE),
          "4" = ~ perc_n(NSYM >= 4, digits=1, na_rm = TRUE)),
    "Medical history" =
    list("Previous myocardial infarction" = ~ perc_n(PREVM1 == 1, digits=1, na_rm = TRUE),
          "Current angina" = ~ perc_n(ANGINA == 1, digits=1, na_rm = TRUE),
          "Diabetes" = ~ perc_n(DIABETES == 1, digits=1, na_rm = TRUE),
          "Hypertension" = ~ perc_n(HYPERTEN == 1, digits=1, na_rm = TRUE),
          "Previous digoxin use" = ~ perc_n(DIGUSE == 1, digits=1, na_rm = TRUE)),
    "Primary cause of CHF" =
    list("Ischemic" = ~ perc_n(CHFETIOL == 1, digits=1, na_rm = TRUE),
          "Nonischemic" = ~ perc_n(CHFETIOL != 1, digits=1, na_rm = TRUE),
          "Idiopathic" = ~ perc_n(CHFETIOL == 4, digits=1, na_rm = TRUE),
```

```
# whole <- summary_table(dig, our_summary1, by = c("TRTMT"))
dig$TRTMT = relevel(as.factor(dig$TRTMT), ref = '1')
levels(dig$TRTMT) <- list("DIGOXIN"="1", "PLACEBO"="0")
levels(dig$TRTMT)
```

```
whole <- summary_table(dplyr::group_by(dig, TRTMT), our_summary1)
whole
```

2

