Writing a simple R package in S3.

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Figure 1: S3 OOP in R

1 Introduction

Ok! It time to step up my R programming to S3. But how to start?

I'll begin by reading the relevant chapters in Advanced R (add reference).

Other references:

Introduction to Scientific Programming and Simulation using R. Jomes. Maillardet, Robinson.

A Simple Guide to S3 Methods | R-bloggers

Why your S3 method isn't working | R-bloggers

Dealing with S3 methods in R with a simple example \mid R-bloggers

Video on S3 Classes in R by Dr Andrew Robinson | R-bloggers

Unexported S3 Methods and R Packages | R-bloggers

2 Appendix zz.table1.c

```
zz.table1.c = function(df, form, pv=TRUE, totl=TRUE, grps=TRUE) {
if (!require("pacman")) install.packages("pacman", repo="cran.rstudio.com")
pacman::p_load(janitor )
prep = function(df, form) {
dfr = df \%
    ungroup %>%
    sel(all.vars(form[[3]]))
df_list = dfr %>% split( df_grp) %>%
list_merge(., "Total" = dfr) %>%
purrr::transpose()
process1 = function(x){
pv_chr = data.frame(x[["Total"]],df_grp) %>%
pvalue_chr
ll = x[[length(x)]]\%\% as.factor \%\% levels
ll_indent = paste("\\hspace{5mm}
                                  ",11)
sum_chr = x %>%
    lapply(function(x) factor(x, levels=11)) %>%
    map(categ)
                %>% as_tibble %>%
    cbind(variable=ll_indent, ., 'p-value'=NA) %>%
    mut(variable=as.character(variable)) %>%
    rbind(NA, .)
sum_chr[1,ncol(sum_chr)]=pv_chr
# browser()
return(sum_chr)
process2 = function(x){
pv_num = data.frame(x[["Total"]],df_grp) %>%
   pvalue_num
sum_num = x \%>\%
  map_chr(contin) %>%
   bind_rows %>%
   cbind(variable=NA,., 'p-value'=pv_num)
                                             %>%
```

```
mut(variable=as.character(variable))
return(sum_num)
}
contin= function(x) {
s1 = zz.sum.min(x)
pasteO(s1['Mean'],"$\\pm$", s1['SD'], " ({\\scriptsize $",s1['N'],"$})") }
categ = function(x) {
prps = table(x) %>% prop.table %>% round(2)*100
cnts_prps = table(x)%>%
paste0(.," ({\\scriptsize $",prps,"$})")
pvalue_num = function(df) {
tidy(anova(lm(as.formula(paste(names(df), collapse="~")), data = df)))$p.value[1]
pvalue_chr = function(df) {
tab = table(df[,1], df[,2])
ifelse((nrow(tab) >=2 & ncol(tab) >=2),
       stats::fisher.test(tab,simulate.p.value=T)$p.value, NA)
fieldclass =sapply(df, class)%>% enframe %>%
slice(match(all.vars(form[[3]]),name))
groupclass =sapply(df, class)%>% enframe %>%
slice(match(all.vars(form[[2]]),name))
df_grp<- df %>% pull(groupclass$name)
df2 = prep(df, form)
out = df2 \% > \%
 map_if(fieldclass$value=="numeric" | fieldclass$value=="integer", function(x){process2(x)}
       map_if(fieldclass$value=="character", function(x){process1(x)})
imap(function(x,y) {
         y2 = ifelse(fieldclass$value[fieldclass$name == y]=="character",
             paste(y, "-- {\\scriptsize no. (\\%)}"), y)
     x[1,1]=y2
# browser()
     X
       }) %>%
   bind_rows()
on= names(out)
nn = tabyl(df_grp)%>%
    adorn_totals() %>%
```

```
pull(n)
names(out) = paste(rep("{\\bf",length(on)),on, c("",paste0("\\scriptsize(n=",nn,")"),""),rep
if (!grps) out = out %>% sel(contains("variable"),contains("Total"),contains("p-value"))
if (!pv) {
   out = out %>% sel(-contains("p-value"))}
if (!totl) out = out %>% sel(-contains("Total"))
return(out)
}
```