Model Complete Cases

Martin Skarzynski 2018-04-20

```
library(here)
## here() starts at /Users/marskar/gdrive/nhanes
library(readr)
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(tidyr)
library(survey)
## Loading required package: grid
## Loading required package: methods
## Loading required package: Matrix
##
## Attaching package: 'Matrix'
## The following object is masked from 'package:tidyr':
##
##
       expand
## Loading required package: survival
##
## Attaching package: 'survey'
## The following object is masked from 'package:graphics':
##
##
       dotchart
library(purrr)
# this function takes in two integers as arguments
# the seed and the number of random variables
# and returns a dataframe
get_modelstats <- function(seed, n_random_vars){</pre>
    set.seed(seed)
    #Variables included in the model
    chosen_vars <- c(</pre>
                     "PERMTH_INT",
```

```
"canc_mort",
                  "SDPPSU6",
                  "SDPSTRA6",
                  "WTPFQX6"
remove_vars <- c(</pre>
                  "HAN9", #remove age variables
                  "HAQ7",
                  "HAT29",
                  "HAJO",
                  "WTPXRP2", #remove unneed weight variables
                  "WTPQRP21",
                  "WTPQRP27",
                  "WTPQRP43"
#remove variables,
#move chosen variables to the beginning, and
#sample from the variables to be randomized
read_rds(here('dat/3-clean-complete-cases.rds')) %>%
    select(-one_of(remove_vars)) %>%
    select(one_of(chosen_vars),
           everything()[sample(seq(ncol(.)),
                               n_random_vars)]) ->
dat
# create survey design object
svydesign(ids = ~SDPPSU6,
          strata = ~SDPSTRA6,
          weights = ~WTPFQX6,
          nest = TRUE,
          data = dat) ->
des
# create left sides of equations
form <- as.formula(Surv(PERMTH_INT, canc_mort) ~ x1)</pre>
# create right sides of equations
n_params <- ncol(dat) # the total number of parameters</pre>
# n_params should equal the number of chosen variables
# plus the number of random variables
r_cols <- (length(chosen_vars)+1):n_params</pre>
if(n_random_vars==1){
vrs <- as.name(names(dat)[n_params])</pre>
vrs2 <- as.name(names(dat)[n_params])</pre>
} else{
vrs <- as.name(paste(names(dat)[r_cols], collapse=' + '))</pre>
vrs2 <- as.name(paste(names(dat)[r_cols], collapse=', '))</pre>
}
set.seed(seed)
\#train \leftarrow sample(x = seq(nrow(dat)),
                 n_random_vars = round(nrow(dat)*.7))
```

```
# generate cox models without and with penalties
    cox <- try(svycoxph(update(form,</pre>
                            paste("~ ", vrs)),
                     design = des, data = dat))
    rid <- try(svycoxph(update(form,</pre>
                            paste("~ ridge(", vrs2, ')')),
                     design = des, data = dat))
    # define functions needed to create first table
    get_con <- function(x) {</pre>
    signif(summary(x)$concordance[1]*100, digits = 2)
    }
    get_HR <- function(x) {</pre>
    summary(x)$conf.int[,"exp(coef)"]
    get_HR_CI_lower <- function(x) {</pre>
    summary(x)$conf.int[,"lower .95"]
    get_HR_CI_upper <- function(x) {</pre>
    summary(x)$conf.int[,"upper .95"]
    }
    get_coef_pvalue <- function(x) {</pre>
        coefs <- summary(x)$coef</pre>
        coefs[,ncol(coefs)]
    model_list <- try(list(cox, rid))</pre>
    try(data_frame(seed = rep(seed,2),
               n_random_vars = n_random_vars,
                type = c('coxph', 'ridge'),
                aic = AIC(cox, rid)[,"AIC"],
                concordance = map_dbl(model_list,
                                        get_con),
               hazard_ratio = map(model_list,
                                    get_HR),
               HR_CI_lower = map(model_list,
                                    get_HR_CI_lower),
                               map(model_list,
               HR_CI_upper =
                                    get_HR_CI_upper),
                coef_pvalue = map(model_list,
                                   get_coef_pvalue)))
}
#save an object with 1000 models
map_seeds <- function(seed){</pre>
map2_dfr(.x = seed,
         y = seq(10), #max number of variables to randomize
         get_modelstats)
map_dfr(seq(50), map_seeds) %>%
```

write_rds(here(paste0("dat/4-model-complete-cases.rds")))

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## With (98) clusters.
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