1

```
rm(list = ls())
# Local github
codepath <- c("/media/larryleon/My Projects/GitHub/Forest-Search/R/")</pre>
source(paste0(codepath, "source_forestsearch_v0.R"))
source_fs_functions(file_loc = codepath)
library(kableExtra)
library(knitr)
library(ggplot2)
library(gridExtra)
library(cubature)
library(aVirtualTwins)
library(randomForest)
library(survival)
library(survminer)
library(grf)
library(policytree)
library(data.table)
library(plyr)
library(dplyr)
library(glmnet)
library(corrplot)
library(table1)
library(cli) # for colors in cat
```

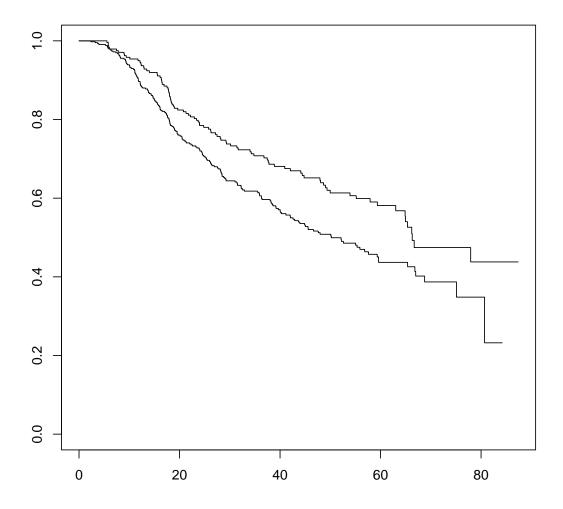
```
t.start.all <- proc.time()[3]

df.analysis <- gbsg
df.analysis <- within(df.analysis, {
    id <- as.numeric(c(1:nrow(df.analysis)))
    # time to months
    time_months <- rfstime/30.4375
    grade3 <- ifelse(grade == "3", 1, 0)
})

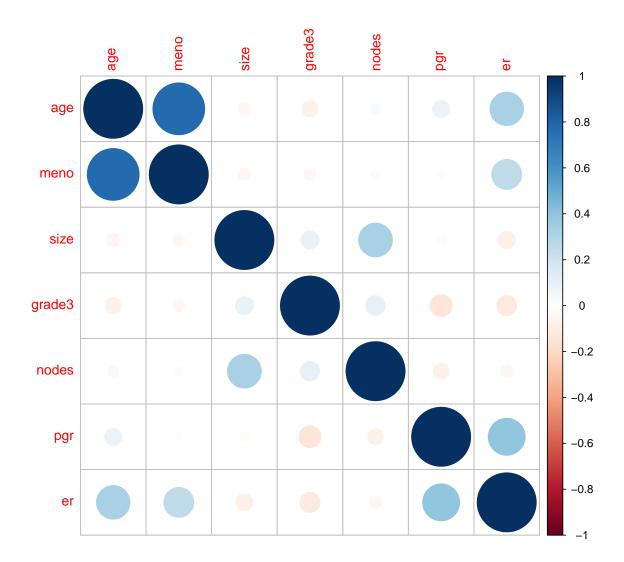
confounders.name <- c("age", "meno", "size", "grade3", "nodes", "pgr", "er")

outcome.name <- c("time_months")
event.name <- c("status")
id.name <- c("id")
treat.name <- c("hormon")

plot(survfit(Surv(time_months, status) ~ hormon, data = df.analysis))</pre>
```



Zm <- cor(as.matrix(df.analysis[, c(confounders.name)]))
corrplot(Zm)</pre>



```
# suppressWarnings(table1 (~ age + wtkg + karnof + cd40 + cd80 + hemo + homo +
# drugs + race + gender + oprior + symptom | treat, data=df.analysis))

hr.threshold <- 1.25  # Initital candidates
hr.consistency <- 1  # Candidates for many splits
pconsistency.threshold <- 0.9

stop.threshold <- 0.99

# NOTE: Allows for Age intervals since GRF cuts at 2 age levels
maxk <- 3

nmin.fs <- 60

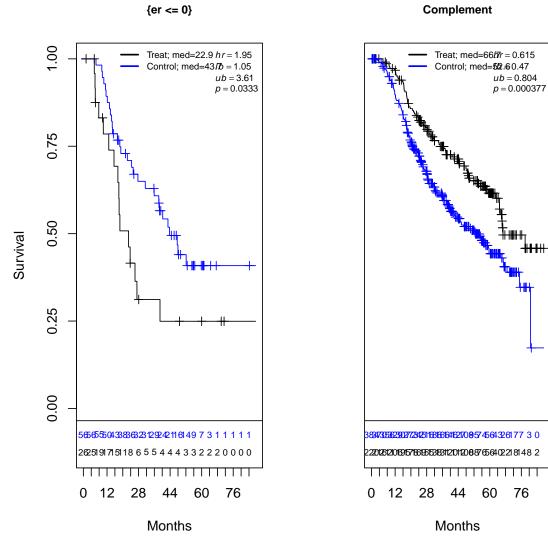
pstop_futile <- 0.5

# Limit timing for forestsearch</pre>
```

```
max.minutes <- 3</pre>
m1.threshold <- Inf # Turning this off (Default)</pre>
# pconsistency.threshold<-0.70 # Minimum threshold (will choose max among
# subgroups satisfying)
fs.splits <- 400 # How many times to split for consistency
# vi is % factor is selected in cross-validation --> higher more important
vi.grf.min <- 0.2</pre>
# Null, turns off grf screening
d.min <- 10 # Min number of events for both arms (d0.min=d1.min=d.min)
# default=5
use_lasso <- TRUE
use_grf <- TRUE
use_grf_only <- FALSE</pre>
# Now run with stop.threshold
fs.est <- forestsearch(df.analysis = df.analysis, Allconfounders.name = confounders.name,
   details = TRUE, use_lasso = use_lasso, use_grf = use_grf, use_grf_only = use_grf_only,
   dmin.grf = 12, frac.tau = 1, maxk = maxk, max_n_confounders = 11, sg_focus = "Nsg_only",
   stop.threshold = stop.threshold, grf_depth = 2, outcome.name = outcome.name,
   treat.name = treat.name, event.name = event.name, id.name = id.name, n.min = nmin.fs,
   hr.threshold = hr.threshold, hr.consistency = hr.consistency, fs.splits = fs.splits,
   d0.min = d.min, d1.min = d.min, pstop_futile = pstop_futile, pconsistency.threshold = pconsisten
   max.minutes = max.minutes, by.risk = 4, plot.sg = TRUE, vi.grf.min = vi.grf.min)
## tau, maxdepth= 77.93018 2
## leaf.node control.mean control.size control.se treated.mean treated.size
          2 5.698218 82.000000 6.409425 -5.698218
                                                                 82.000000
            3
## 2
                 -8.273804 604.000000 2.134322
                                                     8.273804 604.000000
## 11
            4 -19.921490 112.000000 5.374664 19.921490 112.000000
## 21
            5
                 8.189949 177.000000 3.777446 -8.189949 177.000000
            7 -11.509826 356.000000 2.718385 11.509826
                                                                 356.000000
## treated.se diff Nsg depth
     6.409425 11.39644 82
## 1
     2.134322 -16.54761 604
## 11    5.374664   -39.84298   112
## 21  3.777446  16.37990  177
## 4 2.718385 -23.01965 356
## leaf.node control.mean control.size control.se treated.mean treated.size
## 21
         5
                 8.189949 177.000000 3.777446 -8.189949 177.000000
## treated.se
                 diff Nsg depth
## 21 3.777446 16.3799 177
## Subgroup found
## [1] "age <= 50" "age <= 43" "er <= 0"
## [1] "age <= 43"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.01843119
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                   s0
## age
## meno
```

```
## size 0.005433435
## grade3 0.178139021
## nodes 0.049670523
## pgr
         -0.001812895
## Cox-LASSO selected: size grade3 nodes pgr
## Cox-LASSO not selected: age meno er
## Median cuts after Lasso: size nodes pgr
## Categorical after Lasso: grade3
## Factors per GRF: age <= 50 age <= 43 er <= 0
## Medians prior to removing if also in GRF: size nodes pgr
## Factors after removing any duplicates also in GRF: size nodes pgr
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 50 age <= 43 er <= 0 \,
## Factors included per GRF (not in lasso) age <= 50 age <= 43 er <= 0
## # of candidate subgroup factors= 7
## [1] "size <= median(size)" "nodes <= median(nodes)" "pgr <= median(pgr)"
## [4] "age <= 50"
                              "age <= 43"
## [7] "grade3"
## LMAX= 7
## Confounders per grf screening q6 q1 q7 q4 q2 q3 q5
## FSconfounders.name
                           vi.cs
## 6
                    q6 0.1931314
## 1
                    q1 0.1653027
## 7
                    q7 0.1500610
## 4
                    q4 0.1394943
## 2
                    q2 0.1347695
## 3
                    q3 0.1240801
                    q5 0.0931610
## Number of unique levels (L) and possible subgroups= 14 16383
## # of subgroups based on # variables > k.max and excluded (per million) 0.015914
## k.max= 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 164 270
## # of subgroups with sample size less than criteria 246
## # of subgroups meeting all criteria = 178
## # of subgroups fitted (Cox model estimable) = 178
## *Subgroup Searching Minutes=* 0.009366667
## Number of subgroups meeting HR threshold 14
## # of candidate subgroups (meeting HR criteria) = 14
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
##
      K n E d1 m1 m0 HR L(HR) U(HR) q6.0 q6.1 q1.0 q1.1 q7.0 q7.1 q4.0
## 1: 2 177 55 18 66.20
                        NA 1.53 0.87 2.69
                                                0
                                                    0
                                                          0
                                                              0
                                                                     0
## 2: 3 150 41 15 66.20
                        NA 1.55 0.82 2.94
                                                 1
                                                           0
## 3: 2 142 72 18 27.17 39.66 1.40 0.82 2.39
                                                 0
                                                      0
                                                           0
                                                                0
## 4: 3 139 41 13 66.20
                        NA 1.33 0.69 2.57
                                                                0
                                                 0
                                                      0
                                                           0
                                                                     1
                                                                          0
## 5: 3 84 43 12 27.17 44.88 1.59 0.81 3.10
                                                 0
                                                      0
                                                           0
                                                                0
                                                                     1
                                                                          0
## 6: 1 82 45 16 22.93 43.66 1.95 1.05 3.61
                                                 0
                                                                     0
                                                      1
                                                           0
                                                                0
## 7: 3 80 37 12 16.49 NA 2.51 1.25 5.01
                                                 0
                                                      0
                                                           0
                                                                0
                                                                     0
                                                                          0
                                                                               0
                                               0
                                                    0
                                                                0
                                                                     0
                                                                               0
## 8: 3 76 39 16 32.41 52.14 1.42 0.75 2.70
                                                           0
                                                                          0
                                                                0
                                                                     0
                                                                               0
## 9: 2 75 41 16 18.53 47.61 2.22 1.18 4.20 0 1
                                                           0
                                                                          0
## 10: 3 73 37 13 30.59 47.61 1.34 0.68 2.64
```

```
## q4.1 q2.0 q2.1 q3.0 q3.1 q5.0 q5.1
## 1: 1 0 0 0 0
## 2: 1
           0
                0
                   0
                         0
       1
           0
## 3:
                0
                    0
                        1
                             0
                                 0
          0
## 4:
      1
                0
                    0
                        0
                             1
                                 0
## 5: 1 0
              0 0
                        1
                           0
## 6: 0 0 0 0
                           0 0
## 7: 1 0 0 0 1 1 0
## 8: 1 1 0 0 0 1 0
## 9: 0 0 0 1 0 0
      0 0 0 0 0 1 0
## 10:
## Consistency 0.8525
## Consistency 0.7925
## Consistency 0.7625
## Consistency 0.59
## Consistency 0.8475
## Consistency 0.9625
## # of splits= 400
## Model, % Consistency Met= {er <= 0} 0.9625
## Consistency 0.975
## # of splits= 400
## Model, % Consistency Met= {age <= 50} {pgr <= median(pgr)} !{age <= 43} 0.975
## Consistency 0.63
## Consistency 0.9925
## # of splits= 400
## Model, % Consistency Met= {er <= 0} {pgr <= median(pgr)} 0.9925
## Number of subgroups meeting consistency criteria=
## Pcons NgmK
                   M.1
                                         M.2
                                                    М.З
## 1: 0.9625 82 1 6 1 {er <= 0}
## 2: 0.9750 80 8 7 3 {age <= 50} {pgr <= median(pgr)} !{age <= 43}
## 3: 0.9925 75 6 9 2 {er <= 0} {pgr <= median(pgr)}
```



```
## [1] "{er <= 0}"
## % consistency criteria met= 0.9625
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.2091833
## Subgroup found (FS)
## Minutes overall= 0.2381667

file_out <- c("output/gbsg_Nsg_results_b=1000_v0a.Rdata")

library(doParallel)
registerDoParallel(parallel::detectCores(logical = FALSE))

cox.formula.boot <- as.formula(paste("Surv(time_months,status)~hormon"))
max.minutes <- 3

# Suggest running 20, first ... to get timing estimate

NB <- 1000</pre>
```

```
df_boot_analysis <- fs.est$df.est</pre>
fitH <- get_Cox_sg(df_sg = subset(df_boot_analysis, treat.recommend == 0), cox.formula = cox.formula
H_obs <- fitH$est_obs # log(hr) scale</pre>
seH_obs <- fitH$se_obs
# Hc observed estimates
fitHc <- get_Cox_sg(df_sg = subset(df_boot_analysis, treat.recommend == 1), cox.formula = cox.formul
Hc_obs <- fitHc$est_obs</pre>
seHc_obs <- fitHc$se_obs</pre>
rm("fitH", "fitHc")
Ystar_mat <- bootYstar({</pre>
   ystar <- get_Ystar(boot)</pre>
}, boots = NB, seed = 8316951, counter = "boot", export = fun_arg_list_boot)
# Check dimension
if (dim(Ystar_mat)[1] != NB | dim(Ystar_mat)[2] != nrow(df_boot_analysis)) stop("Dimension of Ystar_
# Check 1st 10 bootstraps
ansB <- NULL
for (bb in 1:10) {
   boot <- bb
   ans <- fsboot_forparallel(boot)</pre>
   cat_line("***Bootstrap done, B***=", c(boot), col = "blue")
   print(ans)
   ansB <- rbind(ansB, c(bb, ans))</pre>
## tau, maxdepth= 66.69405 2
## leaf.node control.mean control.size control.se treated.mean treated.size
         2
## 1
                 5.188109 85.000000 4.845534 -5.188109
                                                                85.000000
                                        1.655657
                 -6.692222 601.000000
## 2
            3
                                                     6.692222 601.000000
                                                   16.214534
## 11
            4
                -16.214534
                             84.000000 4.989337
                                                                 84.000000
            5
                  7.998578 133.000000 3.475361 -7.998578 133.000000
## 21
## 4
            7 -8.991534 412.000000 1.924000 8.991534 412.000000
## treated.se diff Nsg depth
## 1 4.845534 10.37622 85
      1.655657 -13.38444 601
## 11 4.989337 -32.42907 84
## 21   3.475361   15.99716   133
     1.924000 -17.98307 412
                                 2
## leaf.node control.mean control.size control.se treated.mean treated.size
## 21
        5 7.998578 133.000000 3.475361 -7.998578 133.000000
                   diff Nsg depth
## treated.se
## Subgroup found
## [1] "age <= 48" "age <= 42" "er <= 0"
## [1] "age <= 42"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.01144389
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                    s0
## age -0.0067445197
```

```
## meno -0.0334900698
## size 0.0050479396
## grade3 0.3004935560
## nodes 0.0431947483
## pgr
         -0.0008205332
## er
        -0.0001016426
## Cox-LASSO selected: age meno size grade3 nodes pgr er
## Cox-LASSO not selected:
## Median cuts after Lasso: age size nodes pgr er
## Categorical after Lasso: meno grade3
## Factors per GRF: age <= 48 age <= 42 er <= 0
## Medians prior to removing if also in GRF: age size nodes pgr er
## ***cMed_flag***=age
## ***cMed_flag***=age
## ***cMed_flag***=er
## ***to_exclude***=TRUE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=TRUE
## ***conf.cont_medians***=size
## ***conf.cont_medians***=nodes
## ***conf.cont_medians***=pgr
## Factors after removing any duplicates also in GRF: size nodes pgr
## ***Factors per lasso after omitting GRF dups***=meno
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 48 age <= 42 er <= 0
## Factors included per GRF (not in lasso) age <= 48 age <= 42 er <= 0
## # of candidate subgroup factors= 8
## [1] "size <= median(size)"  "nodes <= median(nodes)"  "pgr <= median(pgr)"
                              "age <= 42"
## [4] "age <= 48"
                                                         "er <= 0"
## [7] "meno"
                                "grade3"
## LMAX= 7
## Confounders per grf screening q8 q6 q2 q3 q4 q1 q7
## FSconfounders.name vi.cs
## 8
                    q8 0.24823961
## 6
                    q6 0.21932881
## 2
                    q2 0.12182119
## 3
                    q3 0.11540089
## 4
                    q4 0.11348862
                    q1 0.11333843
## 1
                     q7 0.05908588
## 7
## Number of unique levels (L) and possible subgroups= 14 16383
## # of subgroups based on # variables > k.max and excluded (per million) 0.015914
## k.max= 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 148 287
## # of subgroups with sample size less than criteria 228
## # of subgroups meeting all criteria = 167
## # of subgroups fitted (Cox model estimable) = 167
## *Subgroup Searching Minutes=* 0.008433333
## Number of subgroups meeting HR threshold 4
```

```
## # of candidate subgroups (meeting HR criteria) = 4
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
      K n E d1
                m1 m0 HR L(HR) U(HR) q8.0 q8.1 q6.0 q6.1 q2.0 q2.1 q3.0
## 1:
      2 90 48 13 23.72 38.24 1.44 0.76 2.73
                                          0
                                               1
                                                    0
## 2: 1 85 55 19 18.53 40.25 1.84 1.05 3.24
                                          0
                                                0
                                                         1
                                                             0
                                                    0
                                                                      0
## 3: 2 75 47 19 18.53 40.25 2.26 1.24 4.09
                                               0
                                                             0
                                          0
                                                    0
                                                        1
                                                                0
                                                                    0
## 4: 3 61 37 13 18.00 31.05 1.83 0.93 3.62
                                          0
                                               1
                                                    0
                                                        0
                                                            0
                                                                0
## 5: NA NA NA NA
                 NA
                      NA
                           NA
                                NA
                                       NA NA
                                               NA
                                                   NA
                                                        NA
                                                            NA
                                                                 NA
                                                                     NA
## 6: NA NA NA NA
                 NA
                      NA
                           NΑ
                                 NΑ
                                      NA NA NA
                                                   NA
                                                        NΑ
                                                            NΑ
                                                                 NΑ
                                                                     NΑ
## 7: NA NA NA NA
                 NA
                      NA
                                      NA
                                           NA NA
                           NA
                                 NA
                                                    NA
                                                        NA
                                                            NA
                                                                 NA
                                                                     NA
## 8: NA NA NA NA
                  NA
                        NA
                            NA
                                 NA
                                       NA
                                           NA
                                               NA
                                                    NA
                                                        NA
                                                            NA
                                                                 NA
                                                                     NA
                                                       NA NA
## 9: NA NA NA NA
                  NA
                       NA
                            NA
                                 NA
                                      NA
                                           NA NA
                                                    NA
                                                                 NA
                                                                     NA
                           NA
                                                  NA
## 10: NA NA NA
                 NA
                      NA
                                 NA
                                      NA NA NA
                                                       NA NA
                                                                 NA
                                                                     NA
##
      q3.1 q4.0 q4.1 q1.0 q1.1 q7.0 q7.1
##
      0 0
               0 1 0
## 2:
       0
           0
                 0
                   0
                          0
                              0
## 3:
            0
                    0
                         0
                                0
       1
                 0
                              0
## 4:
       0
            1
                         0
                                  0
                0
                    1
                             0
## 5:
      NA
           NA
               NA
                   NA
                         NA
                             NA
                                 NA
## 6:
      NA
           NA
               NA NA
                        NA
                            NA
                                 NA
      NA
## 7:
           NA NA NA
                        NA NA
                                NΑ
## 8:
      NA NA
               NA NA
                        NA NA
                                NA
## 9:
      NA NA
               NA NA
                        NA NA
                                 NΑ
## 10:
      NA NA
               NΑ
                   NA NA NA
                                NΑ
## Consistency 0.6575
## Consistency 0.9375
## # of splits= 400
## Model, % Consistency Met= {er <= 0} 0.9375
## Consistency 0.9925
## # of splits= 400
## Model, % Consistency Met= {er <= 0} {pgr <= median(pgr)} 0.9925
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.0657
## Subgroup found (FS)
## Minutes overall= 0.08753333
## ***Bootstrap done, B***=1
## H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
## 1: 0.5765919 0.6342285 -0.423406 -0.3271736 0.0084
## tau, maxdepth= 66.69405 2
## leaf.node control.mean control.size control.se treated.mean treated.size
## 1
        2
             11.355910 87.000000 4.306807 -11.355910 87.000000
                                               6.735298
## 2
           3
               -6.735298 599.000000 1.605803
                                                         599.000000
              12.168174
                                                         84.000000
## 3
          4
                        84.000000 4.409105
                                              -12.168174
              -9.961720 294.000000 2.411649
## 4
          5
                                               9.961720 294.000000
## 5
           6 10.058065 70.000000 4.176711
                                              -10.058065 70.000000
              -7.747583 238.000000 2.370839 7.747583 238.000000
          7
  treated.se
                 diff Nsg depth
    4.306807 22.71182 87
## 1
## 2
     1.605803 -13.47060 599
     4.409105 24.33635 84
## 4
     2.411649 -19.92344 294
                             2
## 5
     4.176711 20.11613 70
## 6 2.370839 -15.49517 238
## leaf.node control.mean control.size control.se treated.mean treated.size
       4 12.168174 84.000000 4.409105 -12.168174
## 3
                                                           84.000000
```

```
## treated.se diff Nsg depth
## 3 4.409105 24.33635 84
## Subgroup found
## [1] "pgr <= 43" "er <= 0" "pgr <= 74"
## [1] "er <= 0"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.009164127
## 7 \times 1 \text{ sparse Matrix of class "dgCMatrix"}
##
                     s0
## age
## meno
          0.0655875411
## size 0.0105561880
## grade3 0.3796314845
## nodes 0.0421993621
## pgr
         -0.0017537694
## er
        -0.0001772295
## Cox-LASSO selected: meno size grade3 nodes pgr er
## Cox-LASSO not selected: age
## Median cuts after Lasso: size nodes pgr er
## Categorical after Lasso: meno grade3
## Factors per GRF: pgr <= 43 er <= 0 pgr <= 74
## Medians prior to removing if also in GRF: size nodes pgr er
## ***cMed_flag***=pgr
## ***cMed_flag***=er
## ***cMed_flag***=pgr
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=TRUE
## ***to_exclude***=TRUE
## ***conf.cont_medians***=size
## ***conf.cont_medians***=nodes
## Factors after removing any duplicates also in GRF: size nodes
## ***Factors per lasso after omitting GRF dups***=meno
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## Initial GRF cuts included pgr <= 43 er <= 0 pgr <= 74
## Factors included per GRF (not in lasso) pgr <= 43 er <= 0 pgr <= 74
## # of candidate subgroup factors= 7
## [1] "size <= median(size)" "nodes <= median(nodes)" "pgr <= 43"
                                "pgr <= 74"
## [4] "er <= 0"
## [7] "grade3"
## LMAX= 7
## Confounders per grf screening q4 q7 q2 q1 q3 q6 q5
## FSconfounders.name vi.cs
## 4
                    q4 0.40460324
## 7
                    q7 0.15888880
## 2
                    q2 0.09889540
## 1
                     q1 0.09737508
## 3
                     q3 0.08252573
## 6
                     q6 0.07979085
## 5
                     q5 0.07792090
## Number of unique levels (L) and possible subgroups= 14 16383
```

```
## # of subgroups based on # variables > k.max and excluded (per million) 0.015914
## k.max= 3
## Events criteria for control, exp= 10 10
## # of subgroups with events less than criteria: control, experimental 164 248
## # of subgroups with sample size less than criteria 216
## # of subgroups meeting all criteria = 184
## # of subgroups fitted (Cox model estimable) = 184
## *Subgroup Searching Minutes=* 0.01205
\#\# Number of subgroups meeting HR threshold 16
## # of candidate subgroups (meeting HR criteria) = 16
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
      K n E d1
                 m1 m0 HR L(HR) U(HR) q4.0 q4.1 q7.0 q7.1 q2.0 q2.1 q1.0
                                                         1
## 1: 2 154 85 24 18.00 39.20 1.47 0.91 2.36
                                                    0
                                            0 0
## 2: 2 144 82 23 17.81 39.20 1.60 0.99 2.60
                                                  0
                                           0
                                                      0
                                                         1
## 3: 2 107 76 23 12.88 20.50 1.38 0.83 2.27 0
## 4: 3 101 59 15 18.00 39.20 1.31 0.72 2.35
                                           0
                                                  0
                                                    0
                                                         1
                                                                        1
## 5: 3 95 59 15 17.18 36.40 1.37 0.76 2.47
                                           0
                                                    0
                                                               0
                                                  0
                                                          1
                                                                    0
                                                                        1
## 6: 3 91 66 20 12.88 18.76 1.42 0.84 2.41
                                           0
                                                    0
                                                 0
                                                          1
                                                               1
                                                                   0
                                                                        0
                                           0
## 7: 1 87 57 24 12.88 47.61 3.39 1.97 5.83
                                                 1
                                                      0
                                                           0
                                                               0
                                                                   0
                                                0
## 8: 3 85 64 20 12.88 17.58 1.57 0.92 2.67
                                           0
                                                      0
                                                          1
                                                              1
                                                                   0
                                                                        0
## 9: 3 84 52 18 18.00 29.27 1.26 0.70 2.25 0 0 1 0
                                                                  0
                                                                        0
## 10: 3 82 49 10 17.51 27.66 1.36 0.65 2.82 0 0 0
                                                               0
      q1.1 q3.0 q3.1 q6.0 q6.1 q5.0 q5.1
## 1:
      0
           0
                 0 0
                          0
                             0 1
## 2:
        0
             0
                 1
                      0
                          0
                               0
## 3:
        0
             0
                 0
                      0
                          0
                               0
                                   0
## 4:
        0
             0
                 0
                      0
                          0
                               0
                                   1
## 5: 0
            0
                 1
                      0
                          0
                               0
           0
## 6: 0
                 0 0
                          \cap
                               0 1
## 7: 0 0
               0 0
                          \cap
                               0 0
## 8:
      0 0
                1 0
## 9:
                    0 1
       0
           0
                 0
                             0 1
       0
                    1
## 10:
            0
                          0
                 0
                               0
## Consistency 0.8625
## Consistency 0.93
## # of splits= 400
## Model, % Consistency Met= {grade3} {pgr <= 43} 0.93
## Consistency 0.7675
## Consistency 0.5825
## Consistency 0.6625
## Consistency 0.755
## Consistency 1
## # of splits= 400
## Model, % Consistency Met= {er <= 0} 1
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.1547833
## Subgroup found (FS)
## Minutes overall= 0.1734667
## ***Bootstrap done, B***=2
## H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
## 1: 0.7232828 0.1723476 -0.4346078 -0.3464593 0.012
## tau, maxdepth= 66.69405 2
## leaf.node control.mean control.size control.se treated.mean treated.size
## 1
      2 13.599979 78.000000 4.540939 -13.599979 78.000000
            3 -8.051902 608.000000 1.601027 8.051902
## 2
                                                            608.000000
```

```
## 11 4 -14.420029 122.000000 3.745840 14.420029
                                                                 122.000000
## 21
            5
                 9.000873 155.000000 3.142448 -9.000873
                                                                 155.000000
            7 -10.957677 370.000000 1.950020 10.957677
## 4
                                                                 370.000000
     treated.se
                  diff Nsg depth
     4.540939 27.19996 78
      1.601027 -16.10380 608
## 11 3.745840 -28.84006 122
## 21 3.142448 18.00175 155
## 4 1.950020 -21.91535 370
## leaf.node control.mean control.size control.se treated.mean treated.size
## 1 2 13.599979 78.000000 4.540939 -13.599979 78.000000
## treated.se
                  diff Nsg depth
## 1   4.540939 27.19996 78
## Subgroup found
## [1] "er <= 0"
## [1] "er <= 0"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.01765235
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                    s0
## age
         -0.0003698075
## meno .
## size 0.0050380318
## grade3 0.1759831893
## nodes 0.0524226554
         -0.0012033411
## pgr
## er
## Cox-LASSO selected: age size grade3 nodes pgr
## Cox-LASSO not selected: meno er
## Median cuts after Lasso: age size nodes pgr
## Categorical after Lasso: grade3
## Factors per GRF: er <= 0
## Medians prior to removing if also in GRF: age size nodes pgr
## Factors after removing any duplicates also in GRF: age size nodes pgr
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=age <= median(age)
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included er <= 0
## Factors included per GRF (not in lasso) er <= 0
## # of candidate subgroup factors= 6
## [1] "age <= median(age)"
                              "size <= median(size)"
                                                      "nodes <= median(nodes)"
## [4] "pgr <= median(pgr)"
                              "er <= 0"
                                                      "grade3"
## LMAX= 6
## Confounders per grf screening q3 q4 q6 q5 q1 q2
## FSconfounders.name
                        vi.cs
## 3
                    q3 0.2048646
## 4
                    q4 0.1945507
## 6
                   q6 0.1866930
## 5
                   q5 0.1816801
## 1
                   q1 0.1191562
## 2
                   q2 0.1130554
```

```
## Number of unique levels (L) and possible subgroups= 12 4095
## # of subgroups based on # variables > k.max and excluded (per million) 0.003797
## k.max= 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 112 156
## # of subgroups with sample size less than criteria 138
## # of subgroups meeting all criteria = 136
## # of subgroups fitted (Cox model estimable) = 136
## *Subgroup Searching Minutes=* 0.0057
## Number of subgroups meeting HR threshold 12
## # of candidate subgroups (meeting HR criteria) = 12
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
##
      K n E d1 m1 m0 HR L(HR) U(HR) q3.0 q3.1 q4.0 q4.1 q6.0 q6.1 q5.0
## 1: 2 177 91 22 28.22 39.16 1.39 0.86 2.25
                                              0
                                                   0
                                                        0
                                                            1
## 2: 2 165 95 33 27.17 36.40 1.36 0.89 2.08
                                                    0
                                                            1
                                                         0
## 3: 3 97 72 29 18.00 24.38 1.62 1.01 2.60
                                                    0
                                                         0
                                                             1
                                               1
## 4: 3 97 44 12 34.27 44.88 1.40 0.72 2.72
                                                                  0
                                                                            0
                                               0
                                                    0
                                                         0
                                                             1
                                                                       0
## 5: 3 92 63 32 23.72 28.42 1.44 0.87 2.37
                                                              0
                                                                  0
                                                                            0
                                               1
                                                    0
                                                        0
                                                                       0
## 6: 3 87 62 17 16.43 20.50 1.37 0.78 2.41
                                               1
                                                    0
                                                        0
                                                              1
                                                                  0
                                                                       0
## 7: 3 82 59 19 16.36 20.50 1.35 0.78 2.34
                                               1
                                                    0
                                                        0
                                                             1
                                                                  0
                                                                       1
                                                                            0
                                                             1
## 8: 3 80 47 10 27.70 36.40 1.41 0.70 2.84
                                             0 0
                                                       0
                                                                  \cap
                                                                           0
                                                                       \cap
## 9: 1 78 38 15 18.53 NA 3.04 1.56 5.92 0 0 0 0
                                                                          0
                                                                       0
## 10: 3 78 45 13 18.00 38.24 1.85 0.97 3.54 0 0 1
      q5.1 q1.0 q1.1 q2.0 q2.1
## 1:
       0
             0
                  1
                       0
## 2:
         0
              0
                  0
                       1
## 3:
         0
              0
                  0
                       1
## 4:
        0
             0
                  1
                       0
## 5: 0
                  0
            1
                       1
## 6: 0
            0
                  1
                       \cap
## 7:
      0
           0
                  0
## 8:
       0
            0
                       1
                  1
## 9:
                          0
              0
                       0
        1
                  0
## 10:
         0
              0
                  0
## Consistency 0.8
## Consistency 0.82
## Consistency 0.975
## # of splits= 400
## Model, % Consistency Met= !{nodes <= median(nodes)} {pgr <= median(pgr)} !{size <= median(size)}
## Consistency 0.6225
## Consistency 0.835
## Consistency 0.65
## Consistency 0.6875
## Consistency 0.62
## Consistency 1
## # of splits= 400
## Model, % Consistency Met= {er <= 0} 1
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.1997333
## Subgroup found (FS)
## Minutes overall= 0.2149
## ***Bootstrap done, B***=3
     H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
       0.7619819 0.3176834
                             -0.3713988 -0.2553875 0.005666667
## tau, maxdepth= 75.10472 2
```

```
## leaf.node control.mean control.size control.se treated.mean treated.size
## 1 2 -9.044924 660.000000 1.834078 9.044924 660.000000
           4 -22.426090 102.000000 4.532073
                                                  22.426090 102.000000
## 2
                12.170876 113.000000 3.802054
## 3
           5
                                                   -12.170876 113.000000
## 4
            6
                 6.350012
                            75.000000 5.676159
                                                    -6.350012
                                                                75.000000
## 5
           7 -12.853668 396.000000 2.351096
                                                  12.853668 396.000000
## treated.se diff Nsg depth
## 1 1.834078 -18.08985 660
## 2 4.532073 -44.85218 102
## 3 3.802054 24.34175 113
## 4
     5.676159 12.70002 75
## 5 2.351096 -25.70734 396
## leaf.node control.mean control.size control.se treated.mean treated.size
## 3
           5 12.170876 113.000000 3.802054 -12.170876 113.000000
## treated.se
                 diff Nsg depth
## 3 3.802054 24.34175 113
## Subgroup found
## [1] "age <= 47" "age <= 43" "er <= 0"
## [1] "age <= 43"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.02203198
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                   s0
## age
## meno
          0.009736886
## size
## grade3 0.276082230
## nodes 0.051834823
## pgr -0.001316396
## er
## Cox-LASSO selected: meno grade3 nodes pgr
## Cox-LASSO not selected: age size er
## Median cuts after Lasso: nodes pgr
## Categorical after Lasso: meno grade3
## Factors per GRF: age <= 47 age <= 43 er <= 0
## Medians prior to removing if also in GRF: nodes pgr
## Factors after removing any duplicates also in GRF: nodes pgr
## ***Factors per lasso after omitting GRF dups***=meno
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 47 age <= 43 er <= 0 \,
## Factors included per GRF (not in lasso) age <= 47 age <= 43 er <= 0
## # of candidate subgroup factors= 7
## [1] "nodes <= median(nodes)" "pgr <= median(pgr)"</pre>
                                                      "age <= 47"
## [4] "age <= 43"
                             "er <= 0"
                                                      "meno"
## [7] "grade3"
## LMAX= 7
## Confounders per grf screening q7 q5 q1 q2 q6 q3 q4
## FSconfounders.name vi.cs
## 7
                   q7 0.32940684
## 5
                    q5 0.24197554
## 1
                    q1 0.15598301
```

```
## 2
                   q2 0.09636494
## 6
                   q6 0.07942378
## 3
                   q3 0.05017508
## 4
                   q4 0.04667081
## Number of unique levels (L) and possible subgroups= 14 16383
## # of subgroups based on # variables > k.max and excluded (per million) 0.015914
## k.max= 3
## Events criteria for control, exp= 10 10
## # of subgroups with events less than criteria: control, experimental 169 288
\#\# ## of subgroups with sample size less than criteria 253
## # of subgroups meeting all criteria = 148
## # of subgroups fitted (Cox model estimable) = 148
## *Subgroup Searching Minutes=* 0.01145
## Number of subgroups meeting HR threshold 24
## # of candidate subgroups (meeting HR criteria) = 24
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
      K n E d1 m1 m0 HR L(HR) U(HR) q7.0 q7.1 q5.0 q5.1 q1.0 q1.1 q2.0
## 1: 2 140 61 25 64.89 52.14 1.41 0.84 2.35
                                               0
                                                    1
                                                         0
                                                              0
## 2: 2 113 25 12 64.95 NA 3.07 1.40 6.76
                                                              0
                                                0
                                                     0
                                                         0
                                                                   0
                                                                        0
## 3: 2 108 56 16 27.17 38.24 1.37
                                  0.76 2.46
                                                0
                                                     0
                                                         0
                                                              0
                                                                   0
## 4: 3 107 53 19 18.14 40.25 1.53 0.87 2.69
                                                0
                                                     1
                                                         0
                                                              0
                                                                   0
                                                                            0
                                                       0
## 5: 1 104 58 16 18.53 38.57 2.09 1.17 3.74
                                                \cap
                                                    0
                                                              1
                                                                   \cap
                                                                            0
                                              0
                                                            0
                                                                   0
## 6: 3 97 55 15 17.81 35.91 1.48 0.80 2.71
                                                   0
                                                       0
## 7: 2 97 54 10 16.71 38.24 1.32 0.66 2.63 0 1
## 8: 2 94 53 16 18.14 40.25 2.79 1.52 5.10 0
                                                         0 1
                                                                0
                                                                       0
                                                                            0
## 9: 3 94 39 10 27.17 52.14 1.65 0.78 3.51 0 0
                                                         0 0 0
                                                                       0
                                                                            0
## 10: 3 93 22 11 64.95 NA 4.04 1.74 9.40
                                               0 0
                                                             0
                                                         0
                                                                   0
                                                                       0
                                                                            0
      q2.1 q6.0 q6.1 q3.0 q3.1 q4.0 q4.1
       0
## 1:
             0
                  0 0
                            0
                               1
## 2:
         0
             0
                  0
                       0
                            1
                                 1
## 3:
       1
             0
                  0
                       0
                            1
                                     0
## 4:
      1
             0
                  0
                     0
                     0
## 5:
       0
            0
                  0
                            \cap
                                 \cap
       1
## 6:
                       0
             1
                  0
                            1
                                 0
        0
## 7:
              1
                  0
                       0
                            0
                                 0
## 8:
         1
              0
                  0
                       0
                            0
                                 0
## 9:
         1
              1
                  0
                       0
                            0
                                 1
                                     0
## 10:
         0
             1
                  0
                                 1
## Consistency 0.7925
## Consistency 0.9825
## # of splits= 400
## Model, % Consistency Met= {age <= 47} !{age <= 43} 0.9825
## Consistency 0.685
## Consistency 0.8125
## Consistency 0.9675
## # of splits= 400
## Model, % Consistency Met= {er <= 0} 0.9675
## Consistency 0.7675
## Consistency 0.485
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.1554167
## Subgroup found (FS)
## Minutes overall= 0.1735333
## ***Bootstrap done, B***=4
     H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
```

```
## 1: 0.3394555 0.2703407 -0.3735976 -0.1709955 0.01145 4.041984
## tau, maxdepth= 66.69405 2
   leaf.node control.mean control.size control.se treated.mean treated.size
          2
                -8.201299 506.000000 1.923748
                                                  8.201299 506.000000
## 2
            3
                 3.137727
                            180.000000 2.579216
                                                    -3.137727
                                                               180.000000
## 3
            4 -11.231654 131.000000 3.819340
                                                  11.231654 131.000000
## 4
            5
                9.829112 128.000000 3.597026
                                                  -9.829112 128.000000
## 5
            6 -14.563108 280.000000 2.449286
                                                  14.563108 280.000000
           7
                4.801527 147.000000
                                        2.869781
                                                    -4.801527 147.000000
                    diff Nsg depth
## treated.se
## 1 1.923748 -16.402597 506
      2.579216
                6.275454 180
      3.819340 -22.463308 131
## 4   3.597026   19.658224   128
## 5 2.449286 -29.126217 280
## 6 2.869781 9.603053 147
   leaf.node control.mean control.size control.se treated.mean treated.size
          5
                9.829112 128.000000 3.597026 -9.829112 128.000000
## treated.se
                  diff Nsg depth
## 4 3.597026 19.65822 128
## Subgroup found
## [1] "age <= 48" "age <= 44" "er <= 101"
## [1] "age <= 44"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.01499184
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                    s0
## age
         -0.0014966338
## meno
          0.0026819002
## size
## grade3 .
## nodes 0.0685438158
## pgr
         -0.0015853423
## er
         -0.0007426255
## Cox-LASSO selected: age size nodes pgr er
## Cox-LASSO not selected: meno grade3
## Median cuts after Lasso: age size nodes pgr er
## Categorical after Lasso:
## Factors per GRF: age <= 48 age <= 44 er <= 101
## Medians prior to removing if also in GRF: age size nodes pgr er
## ***cMed_flag***=age
## ***cMed_flag***=age
## ***cMed_flag***=er
## ***to_exclude***=TRUE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=TRUE
## ***conf.cont_medians***=size
## ***conf.cont_medians***=nodes
## ***conf.cont_medians***=pgr
## Factors after removing any duplicates also in GRF: size nodes pgr
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
```

```
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 48 age <= 44 er <= 101
## Factors included per GRF (not in lasso) age <= 48 age <= 44 er <= 101
## # of candidate subgroup factors= 6
## [4] "age <= 48"
                            "age <= 44"
                                                   "er <= 101"
## LMAX= 5
## Confounders per grf screening q6 q1 q4 q3 q2
## FSconfounders.name
                         vi.cs
## 6
                  q6 0.3127067
## 1
                  q1 0.2439341
## 4
                  q4 0.1771667
## 3
                   q3 0.1242483
## 2
                   q2 0.1174634
## Number of unique levels (L) and possible subgroups= 10 1023
## # of subgroups based on # variables > k.max and excluded (per million) 0.000848
## k.max= 3
## Events criteria for control, exp= 10 10
## # of subgroups with events less than criteria: control, experimental 63 87
## # of subgroups with sample size less than criteria 68
## # of subgroups meeting all criteria = 84
## # of subgroups fitted (Cox model estimable) = 84
## *Subgroup Searching Minutes=* 0.003716667
## Number of subgroups meeting HR threshold 11
## # of candidate subgroups (meeting HR criteria) = 11
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
##
      K n E d1 m1 m0 HR L(HR) U(HR) q6.0 q6.1 q1.0 q1.1 q4.0 q4.1
## 1: 2 219 102 23 47.93 68.76 1.33 0.83 2.12
                                             0
                                                  1
                                                       0
                                                           0
## 2: 2 147 50 24 NA 66.99 1.36 0.78 2.36
                                               1
                                                   0
                                                        \cap
                                                            0
## 3: 2 128 66 19 17.51 42.58 1.96 1.15 3.35
## 4: 3 120 62 19 17.51 42.58 1.96 1.14 3.38
                                               0
                                                        0
                                                            0
                                                   1
                                                                      1
## 5: 3 117 59 14 27.17 68.76 1.62 0.88 2.96
                                               0
                                                   1
                                                        1
                                                            0
                                                                      1
## 6: 3 96 61 16 17.74 28.45 1.29 0.72 2.28
                                               0
                                                   1
                                                        0
                                                            0
## 7: 2 80 26 15 55.16
                        NA 2.12 0.97 4.63
                                               1
                                                   0
                                                        1
                                                            0
                                             1 0
                                                      0
                                                               0
## 8: 2 76 25 14 NA 59.60 1.26 0.57 2.78
                                                            0
                                                                      0
## 9: 3 66 36 10 11.73 39.66 3.30 1.57 6.93
                                              0 0 1
                                                            0 0 1
## 10: 3 64 44 14 9.76 18.30 2.08 1.09 3.97
                                             0 0 0
      q3.0 q3.1 q2.0 q2.1
## 1:
       0
            0
                 0
## 2:
        0
             0
                  0
## 3:
        0
             1
                  0
## 4:
        0
             1
                  0
## 5: 0
             0
                  \cap
## 6:
      0
           0
                 1
## 7:
       0 0
## 8:
       0
           0
                 1
## 9:
       0
                      0
             1
                 0
## 10:
        0
             1
                  1
## Consistency 0.72
## Consistency 0.71
## Consistency 0.955
## # of splits= 400
## Model, % Consistency Met= {age <= 48} {pgr <= median(pgr)} 0.955
## Consistency 0.9775
```

```
## # of splits= 400
## Model, % Consistency Met= {er <= 101} {age <= 48} {pgr <= median(pgr)} 0.9775
## Consistency 0.8625
## Consistency 0.555
## Consistency 0.91
## # of splits= 400
## Model, % Consistency Met= !{er <= 101} !{size <= median(size)} 0.91
## Consistency 0.4475
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.1811
## Subgroup found (FS)
## Minutes overall= 0.19155
## ***Bootstrap done, B***=5
     H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
## 1: 0.4531755 0.7810623 -0.3841067 -0.4618613 0.003666667
## tau, maxdepth= 75.10472 2
## leaf.node control.mean control.size control.se treated.mean treated.size
               -9.004349 543.000000 1.970610 9.004349 543.000000
## 1
          2
                3.519685 143.000000 4.016152
                                                  -3.519685 143.000000
## 2
           3
           4 -11.220478 254.000000 2.998119
                                                11.220478 254.000000
## 3
## 4
           5
                5.642840 100.000000 4.278444
                                                  -5.642840 100.000000
## 5
           6 -11.509218 265.000000 2.746475 11.509218 265.000000
          7 14.173336 67.000000 5.744199 -14.173336 67.000000
## 6
## treated.se diff Nsg depth
## 1 1.970610 -18.00870 543
                             1
               7.03937 143
## 2 4.016152
## 3 2.998119 -22.44096 254
      4.278444 11.28568 100
2
                               2
## 6 5.744199 28.34667 67
## leaf.node control.mean control.size control.se treated.mean treated.size
       7 14.173336 67.000000 5.744199 -14.173336
## treated.se
                diff Nsg depth
## 6 5.744199 28.34667 67
## Subgroup found
## [1] "age <= 53" "er <= 44" "size <= 36"
## [1] "size <= 36"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.01673217
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
## age
       0.041853121
## meno
## size 0.008292060
## grade3 0.323439843
## nodes 0.040103619
        -0.002528803
## pgr
## Cox-LASSO selected: meno size grade3 nodes pgr
## Cox-LASSO not selected: age er
## Median cuts after Lasso: size nodes pgr
## Categorical after Lasso: meno grade3
## Factors per GRF: age <= 53 er <= 44 size <= 36
```

```
## Medians prior to removing if also in GRF: size nodes pgr
## ***cMed_flag***=size
## ***to_exclude***=TRUE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***conf.cont_medians***=nodes
## ***conf.cont_medians***=pgr
## Factors after removing any duplicates also in GRF: nodes pgr
## ***Factors per lasso after omitting GRF dups***=meno
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 53 er <= 44 size <= 36
## Factors included per GRF (not in lasso) age <= 53 er <= 44 size <= 36
## # of candidate subgroup factors= 7
## [1] "nodes <= median(nodes)" "pgr <= median(pgr)"</pre>
                                                       "age <= 53"
                              "size <= 36"
## [4] "er <= 44"
                                                        "meno"
## [7] "grade3"
## LMAX= 7
## Confounders per grf screening q5 q7 q1 q4 q3 q2 q6
## FSconfounders.name vi.cs
                   q5 0.33550687
## 5
## 7
                    q7 0.14473257
## 1
                    q1 0.14380982
## 4
                    q4 0.11692239
## 3
                    q3 0.10767578
## 2
                    q2 0.09568279
                    q6 0.05566979
## Number of unique levels (L) and possible subgroups= 14 16383
## # of subgroups based on # variables > k.max and excluded (per million) 0.015914
## k.max= 3
## Events criteria for control, exp= 10 10
## # of subgroups with events less than criteria: control, experimental 167 243
## # of subgroups with sample size less than criteria 219
## # of subgroups meeting all criteria = 192
## # of subgroups fitted (Cox model estimable) = 192
## *Subgroup Searching Minutes=* 0.009133333
## Number of subgroups meeting HR threshold 13
## # of candidate subgroups (meeting HR criteria) = 13
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
                         m0 HR L(HR) U(HR) q5.0 q5.1 q7.0 q7.1 q1.0 q1.1 q4.0
##
      K n E d1 m1
## 1: 1 143 65 28 31.41
                           NA 1.34 0.82 2.19
                                                 1
                                                      0
                                                           0
                                                                0
                                                                     0
## 2: 2 128 74 17 20.76 31.41 1.27 0.74 2.18
                                                 0
                                                       0
                                                           0
                                                                1
                                                                     0
## 3: 3 115 71 17 16.36 31.41 1.75 1.01 3.03
                                                 0
                                                      0
                                                           0
                                                                1
                                                                     0
                                                                          0
                                                                               0
## 4: 2 98 55 23 21.75 28.45 1.35 0.78 2.32
                                                 1
                                                      \cap
                                                           0
                                                                0
                                                                     1
                                                                               0
                                                                          \cap
## 5: 2 95 38 22 44.78
                         NA 1.53 0.80 2.92
                                                 1
                                                      0
                                                           1
                                                                0
                                                                     \cap
                                                                               0
## 6: 2 86 43 18 29.37
                         NA 1.36 0.74 2.50
                                                 1
## 7: 2 84 41 20 31.41 47.61 1.31 0.71 2.44
                                                                0
                                                                     0
                                                                               0
                                                 1
                                                      0
                                                           0
## 8: 3 78 36 21 59.37 59.33 1.36 0.70 2.64
                                                                0
                                                 0
                                                      0
                                                           0
                                                                               0
                                                                     1
                                                                          0
## 9: 3 71 48 29 23.72 26.18 1.25 0.70 2.25
                                                 0
                                                      0
                                                            0
                                                                0
                                                                     1
                                                                          0
                                                                               0
## 10: 3 70 46 15 12.88 20.50 1.47 0.79
                                          2.73
                                                  0
                                                      0
      q4.1 q3.0 q3.1 q2.0 q2.1 q6.0 q6.1
## 1:
       0 0
                   0 0 0
                                0
                                      0
             0
                             0
                                       0
## 2:
       1
                   0
                        0
                                  0
## 3:
              0
                   0
```

```
0
           0
                     0
                       0
                            0
  5:
       0
             0
                  0
## 6:
        1
             0
                  0
                       0
                            0
                                0
   7:
        0
             0
                  0
                      0
                            0
                                0
## 8:
        0
             0
                  0
                      1
                                0
                                     1
## 9:
                  0
                      0
                         0
                              0
                                   0
       1
             1
## 10:
             0
                     0
                              0
        1
                  0
## Consistency 0.715
## Consistency 0.53
## Consistency 0.95
## # of splits= 400
## Model, % Consistency Met= {grade3} {er <= 44} {pgr <= median(pgr)} 0.95
## Consistency 0.715
## Consistency 0.77
## Consistency 0.6525
## Consistency 0.615
## Consistency 0.6225
## Consistency 0.5575
## Consistency 0.79
## Consistency 0.835
## Consistency 0.9725
## # of splits= 400
## Model, % Consistency Met= !{size <= 36} !{age <= 53} 0.9725
## Consistency 0.6925
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.28805
## Subgroup found (FS)
## Minutes overall= 0.3041167
## ***Bootstrap done, B***=6
     H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
## 1: 0.8799324 0.9842622 -0.5793025 -0.6175556
                                                         0.0091
## tau, maxdepth= 68.76386 2
   leaf.node control.mean control.size control.se treated.mean treated.size
        2
               -6.794352 594.000000 1.708990 6.794352 594.000000
## 1
                            92.000000 4.483674
## 2
           3
                8.388327
                                                  -8.388327
                                                              92.000000
              -10.782268
                          126.000000 3.865454
## 3
           4
                                                  10.782268 126.000000
## 4
           5 10.598959
                          168.000000 3.103245 -10.598959 168.000000
## 5
           6 -13.474525 319.000000 2.199612 13.474525 319.000000
          7
                8.386231
                           73.000000 5.125491
                                                -8.386231 73.000000
## treated.se diff Nsg depth
## 1 1.708990 -13.58870 594
     4.483674 16.77665 92
      3.865454 -21.56454 126
## 4
      3.103245 21.19792 168
## 5 2.199612 -26.94905 319
## 6 5.125491 16.77246 73
## leaf.node control.mean control.size control.se treated.mean treated.size
          5
               10.598959 168.000000 3.103245 -10.598959 168.000000
## treated.se
                 diff Nsg depth
## 4 3.103245 21.19792 168
## Subgroup found
## [1] "age <= 50"
                  "age <= 43" "size <= 36"
## [1] "age <= 43"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
```

```
## Categorical characteristics: meno grade3
## CV lambda = 0.02006424
## 7 x 1 sparse Matrix of class "dgCMatrix"
                   s0
## age
## meno 0.155530395
## size
        0.004009881
## grade3 .
## nodes 0.050334712
## pgr
       -0.001934956
## er
## Cox-LASSO selected: meno size nodes pgr
## Cox-LASSO not selected: age grade3 er
## Median cuts after Lasso: size nodes pgr
## Categorical after Lasso: meno
## Factors per GRF: age <= 50 age <= 43 size <= 36
## Medians prior to removing if also in GRF: size nodes pgr
## ***cMed_flag***=size
## ***to_exclude***=TRUE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***conf.cont_medians***=nodes
## ***conf.cont_medians***=pgr
## Factors after removing any duplicates also in GRF: nodes pgr
## ***Factors per lasso after omitting GRF dups***=meno
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 50 age <= 43 size <= 36
## Factors included per GRF (not in lasso) age <= 50 age <= 43 size <= 36
## # of candidate subgroup factors= 6
## [1] "nodes <= median(nodes)" "pgr <= median(pgr)"</pre>
                                                        "age <= 50"
## [4] "age <= 43"
                              "size <= 36"
                                                        "meno"
## LMAX= 6
## Confounders per grf screening q3 q5 q1 q2 q4 q6
## FSconfounders.name vi.cs
                    q3 0.34373668
## 3
## 5
                    q5 0.23846945
## 1
                    q1 0.17924039
## 2
                    q2 0.14397947
                    q4 0.05345684
                    q6 0.04111717
## Number of unique levels (L) and possible subgroups= 12 4095
## # of subgroups based on # variables > k.max and excluded (per million) 0.003797
## k.max = 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 119 169
\#\# # of subgroups with sample size less than criteria 150
## # of subgroups meeting all criteria = 107
## # of subgroups fitted (Cox model estimable) = 107
## *Subgroup Searching Minutes=* 0.005833333
## Number of subgroups meeting HR threshold 27
## # of candidate subgroups (meeting HR criteria) = 27
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
## K n E d1 m1 m0 HR L(HR) U(HR) q3.0 q3.1 q5.0 q5.1 q1.0 q1.1 q2.0
## 1: 2 168 57 27 24.80 NA 2.88 1.71 4.85 0 1 0 0 0 0
```

```
## 2: 2 159 48 15 64.95 NA 1.81 0.98 3.35 O
                                                       0
                                                                         0
## 3: 2 155 74 21 16.49 66.83 2.20 1.32 3.69
                                              0
                                                   1
## 4: 2 140 64 14 17.74 66.83 1.74 0.95
                                                            0
                                                                \cap
                                       3.20
                                              0
                                                   0
                                                       0
                                                                         0
## 5: 3 137 63 14 17.74 66.83 1.67
                                 0.90
                                       3.10
                                              0
                                                   1
                                                       0
                                                            0
                                                                0
                                                                     0
                                                                         0
   6: 3 135 39 15 24.80
                         NA 2.70
                                 1.41 5.19
                                              0
                                                  1
                                                       0
                                                                0
                                                                         0
## 7: 1 131 62 29 31.41
                         NA 1.54 0.93 2.54
                                             0
                                                  0
                                                            0
                                                                0
                                                      1
                                                                     0
                                                                         0
                                            0
                                                 1
## 8: 3 130 44 19 37.65
                         NA 2.42 1.33 4.39
                                                      0
                                                         1
                                                                0
                                                                         0
                                                                0
                                                                         0
## 9: 3 127 40 11
                 NA
                         NA 1.57 0.78 3.15 0 0 0
                                                         1
## 10: 3 122 55 14 17.74 66.83 2.11 1.14 3.91 0 1 0
      q2.1 q4.0 q4.1 q6.0 q6.1
## 1:
       0
                 0
                      0
           1
## 2:
         0
             1
                  0
                      1
## 3:
        1
             0
                  0
                      0
      1
## 4:
             \cap
                  0
                      1
## 5: 1
             0
                 0
                    1
## 6:
      0
           1
                 0 1
## 7:
      0 0
                 0 0
## 8:
       0
                0
                      0
             1
## 9:
                          0
       0
             1
                 0
                      1
## 10:
        1
             0
                  0
                      0
                           0
## Consistency 1
## # of splits= 400
## Model, % Consistency Met= {age <= 50} !{age <= 43} 1
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.02288333
## Subgroup found (FS)
## Minutes overall= 0.0351
## ***Bootstrap done, B***=7
     H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
##
## 1: 0.3242974 0.6546308 -0.3003587 -0.4287666 0.0058 5.6219
## tau, maxdepth= 75.10472 2
   leaf.node control.mean control.size control.se treated.mean treated.size
## 1
          2
                8.897110
                          80.000000 5.460171 -8.897110 80.000000
## 2
           3
                -7.075530 606.000000 1.823131
                                                  7.075530 606.000000
## 3
           4
              -19.209861
                           90.000000 4.805198
                                                19.209861
                                                             90.000000
                          167.000000 3.438575
## 4
           5
                8.717088
                                                 -8.717088 167.000000
## 5
           6
               14.051673
                           66.000000 5.800078
                                                -14.051673
                                                            66.000000
           7 -11.653661 363.000000 2.272171
## 6
                                               11.653661
                                                            363.000000
                diff Nsg depth
## treated.se
## 1 5.460171 17.79422 80
     1.823131 -14.15106 606
     4.805198 -38.41972 90
## 3
      3.438575 17.43418 167
      5.800078 28.10335 66
## 6 2.272171 -23.30732 363
                              2
## leaf.node control.mean control.size control.se treated.mean treated.size
          6 14.051673 66.000000 5.800078 -14.051673
## treated.se
                 diff Nsg depth
## 5 5.800078 28.10335 66
## Subgroup found
## [1] "age <= 50" "age <= 43" "er <= 3"
## [1] "er <= 3"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
```

```
## CV lambda = 0.03936338
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                    s0
## age
## meno
## size
          0.0102478421
## grade3 .
## nodes 0.0518139829
## pgr
       -0.0006167849
## er
## Cox-LASSO selected: size nodes pgr
## Cox-LASSO not selected: age meno grade3 er
## Median cuts after Lasso: size nodes pgr
## Categorical after Lasso:
## Factors per GRF: age <= 50 age <= 43 er <= 3
## Medians prior to removing if also in GRF: size nodes pgr
## Factors after removing any duplicates also in GRF: size nodes pgr
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## Initial GRF cuts included age <= 50 age <= 43 er <= 3 \,
## Factors included per GRF (not in lasso) age <= 50 age <= 43 er <= 3
## # of candidate subgroup factors= 6
## [1] "size <= median(size)" "nodes <= median(nodes)" "pgr <= median(pgr)"
## [4] "age <= 50"
                               "age <= 43"
                                                       "er <= 3"
## LMAX= 6
## Confounders per grf screening q6 q4 q1 q2 q3 q5
## FSconfounders.name vi.cs
## 6
                    q6 0.3517081
## 4
                    q4 0.1696282
## 1
                    q1 0.1421386
## 2
                    q2 0.1199226
## 3
                    q3 0.1147136
## 5
                    q5 0.1018889
## Number of unique levels (L) and possible subgroups= 12 4095
## # of subgroups based on # variables > k.max and excluded (per million) 0.003797
## k.max = 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 105 164
## # of subgroups with sample size less than criteria 150
## # of subgroups meeting all criteria = 113
## # of subgroups fitted (Cox model estimable) = 113
## *Subgroup Searching Minutes=* 0.0056
## Number of subgroups meeting HR threshold 19
## # of candidate subgroups (meeting HR criteria) = 19
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
      K n E d1 m1 m0 HR L(HR) U(HR) q6.0 q6.1 q4.0 q4.1 q1.0 q1.1
## 1: 2 272 132 65 44.12 50.10 1.41 1.00 1.98
                                                0
                                                      0
                                                            0
                                                                    1
                                                                0
## 2: 2 167 54 23 64.95
                         NA 1.95 1.14 3.35
                                                  0
                                                       0
                                                            0
                                                                1
## 3: 3 160 95 52 29.34 28.48 1.45 0.97 2.18
                                                  0
                                                       0
                                                            0
                                                                0
                                                                      1
## 4: 3 141 83 45 28.22 31.05 1.26 0.82 1.95
                                                  0
                                                       0
                                                            0
                                                                 0
## 5: 3 133 33 14 NA NA 1.94 0.97 3.88
                                                 1
                                                       0
                                                            0
                                                                 1
                                                                      0
                                                       0
                                                            0
                                                                           0
## 6: 3 131 49 20 49.05
                         NA 1.30 0.74 2.30
                                                0
                                                                0
                                                                     1
                                                0 0
## 7: 3 122 74 42 28.09 28.42 1.37 0.87 2.18
                                                          1
                                                                0
                                                                   1
                                                                           0
## 8: 1 121 66 31 22.93 38.24 1.29 0.80 2.10
```

```
## 9: 2 112 64 31 18.53 38.24 1.68 1.03 2.75
                                               0
                                                   1
## 10: 3 107 65 38 28.22 31.05 1.28 0.78 2.11
                                                 0
                                                     0
                                                          1
      q2.0 q2.1 q3.0 q3.1 q5.0 q5.1
## 1:
        0
             0
                  0
                       0
## 2:
         0
             0
                  0
                       0
## 3:
             0
                       0
         1
                  0
                            1
                                 0
       0
            0
## 4:
                     1
                  0
                            1
                                 0
## 5:
      0
            0
                 0 0
                           1
## 6:
      0
            0
                 1 0
                           1
## 7:
        1
             0
                  0
                       0
                           0
                                 0
## 8:
        0
             0
                       0
                           0
                  0
                                 0
## 9:
         0
             0
                  0
                       1
                            0
                                 0
## 10:
         0
             0
                  0
                       1
## Consistency 0.9575
## # of splits= 400
## Model, % Consistency Met= !{size <= median(size)} !{age <= 43} 0.9575
## Consistency 0.985
## # of splits= 400
## Model, % Consistency Met= {age <= 50} !{age <= 43} 0.985
## Consistency 0.9175
## # of splits= 400
## Model, % Consistency Met= !{size <= median(size)} !{nodes <= median(nodes)} !{age <= 43} 0.9175
## Consistency 0.6525
## Consistency 0.885
## Consistency 0.6025
## Consistency 0.7775
## Consistency 0.69
## Consistency 0.945
## # of splits= 400
## Model, % Consistency Met= {er <= 3} {pgr <= median(pgr)} 0.945
## Consistency 0.55
## Consistency 0.95
## # of splits= 400
## Model, % Consistency Met= {er <= 3} !{age <= 43} 0.95
## Consistency 0.9975
## # of splits= 400
## Model, % Consistency Met= {er <= 3} {pgr <= median(pgr)} !{age <= 43} 0.9975
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.2859833
## Subgroup found (FS)
## Minutes overall= 0.2982167
## ***Bootstrap done, B***=8
     H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
## 1: 0.6732497 0.6862775 -0.1973118 -0.1846732
                                                          0.0056
## tau, maxdepth= 75.10472 2
   leaf.node control.mean control.size control.se treated.mean treated.size
           2
                           82.000000 5.262939
                                                 -6.820985 82.000000
                 6.820985
## 2
           3
                -5.489502
                          604.000000 1.811470
                                                    5.489502 604.000000
                            82.000000 5.262939
## 3
           4
                 6.820985
                                                   -6.820985
                                                               82.000000
                                                             315.000000
## 4
           5
                -9.515197
                           315.000000 2.558821
                                                    9.515197
                           129.000000 3.803021
## 5
            6
                 9.749882
                                                   -9.749882
                                                              129.000000
## 6
            7
                -9.850669
                            160.000000 3.261377
                                                   9.850669 160.000000
                   diff Nsg depth
## treated.se
## 1 5.262939 13.64197 82 1
## 2 1.811470 -10.97900 604
```

```
## 3 5.262939 13.64197 82 2
## 4 2.558821 -19.03039 315
## 5  3.803021  19.49976  129
     3.261377 -19.70134 160
## leaf.node control.mean control.size control.se treated.mean treated.size
## 5 6 9.749882 129.000000 3.803021 -9.749882 129.000000
## treated.se
                 diff Nsg depth
## 5 3.803021 19.49976 129
## Subgroup found
## [1] "er <= 49" "er <= 0" "pgr <= 80"
## [1] "pgr <= 80"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.009035554
## 7 x 1 sparse Matrix of class "dgCMatrix"
                   s0
       -0.0009486819
## age
## meno
## size
         0.0090564154
## grade3 0.2221186352
## nodes 0.0373909670
## pgr -0.0023773376
## er
         -0.0002635831
## Cox-LASSO selected: age size grade3 nodes pgr er
## Cox-LASSO not selected: meno
## Median cuts after Lasso: age size nodes pgr er
## Categorical after Lasso: grade3
## Factors per GRF: er <= 49 er <= 0 pgr <= 80
## Medians prior to removing if also in GRF: age size nodes pgr er
## ***cMed_flag***=er
## ***cMed_flag***=er
## ***cMed_flag***=pgr
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=TRUE
## ***to_exclude***=TRUE
## ***conf.cont_medians***=age
## ***conf.cont_medians***=size
## ***conf.cont_medians***=nodes
## Factors after removing any duplicates also in GRF: age size nodes
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=age <= median(age)
## ***Factors per lasso after omitting GRF dups***=size <= median(size)
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## Initial GRF cuts included er <= 49 er <= 0 pgr <= 80
## Factors included per GRF (not in lasso) er <= 49 er <= 0 pgr <= 80
## # of candidate subgroup factors= 7
## [4] "er <= 49"
                             "er <= 0"
                                                     "pgr <= 80"
## [7] "grade3"
## LMAX= 7
## Confounders per grf screening q5 q6 q4 q2 q3 q1 q7
## FSconfounders.name
                        vi.cs
```

```
## 5
                   q5 0.19259817
## 6
                   q6 0.18963213
## 4
                   q4 0.17126809
## 2
                    q2 0.13796966
## 3
                   q3 0.13348107
## 1
                   q1 0.09689287
## 7
                   q7 0.07815800
## Number of unique levels (L) and possible subgroups= 14 16383
## # of subgroups based on # variables > k.max and excluded (per million) 0.015914
## k.max= 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 155 243
## # of subgroups with sample size less than criteria 197
## # of subgroups meeting all criteria = 196
## # of subgroups fitted (Cox model estimable) = 196
## *Subgroup Searching Minutes=* 0.01026667
## Number of subgroups meeting HR threshold 32
## # of candidate subgroups (meeting HR criteria) = 32
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
                   m1 m0 HR L(HR) U(HR) q5.0 q5.1 q6.0 q6.1 q4.0 q4.1
      K n
             E d1
## 1: 2 303 163 84 44.78 45.60 1.31 0.96 1.78
                                               0
                                                     0
                                                          0
## 2: 3 265 144 77 47.93 45.60 1.25 0.90 1.74
                                                      \cap
                                                          0
                                                1
                                                               1
## 3: 2 221 113 36 47.93 66.83 1.44 0.97 2.15
                                                          0
                                                 0
                                                      0
## 4: 2 218 77 30 63.01
                         NA 1.39 0.88 2.19
## 5: 3 185 65 28 63.01
                           NA 1.45 0.89 2.38
                                                      0
                                                          0
                                                                    0
                                                                         0
                                                1
## 6: 3 175 94 27 38.87 39.20 1.29 0.82 2.02
                                                0
                                                      0
                                                          0
                                                                    0
                                                               1
                                                                         1
## 7: 2 171 54 23 77.93
                           NA 1.31 0.76 2.25
                                                 0
                                                      0
                                                          0
                                                               0
                                                                    1
                                                                         0
   8: 3 171 54 23 77.93
                                   0.76 2.25
                                                      0
                                                          0
                           NA 1.31
                                                 1
                                                               0
                                                                    1
## 9: 2 170 61 23 66.20
                           NA 1.37 0.82 2.31
                                                 0
                                                     0
                                                         0
                                                                    0
                                                                         0
                                                               0
                                                      0
                                                          0
                                                                         \cap
## 10: 3 156 58 26 63.01
                           NA 1.46 0.87 2.45
                                                               1
      q2.0 q2.1 q3.0 q3.1 q1.0 q1.1 q7.0 q7.1
  1:
        0 0
                  0
                     0
                           0
                                     1
## 2:
       0
            0
                  0
                       0
                            \cap
                                 0
                                     1
## 3:
        0
                       0
                            0
                                     0
                                          0
            0
                  0
                                 1
## 4:
        0
             0
                  0
                       1
                            0
                                 0
                                     0
                                          0
## 5:
        0
             0
                  0
                       1
                            0
                                 0
                                     0
## 6:
       0
            0
                  0
                       0
                            0
                                 1
                                     0
                                          0
      0
## 7:
            0
                  0 1
                           0
                               0
                                     0
## 8:
      0 0
                  0
                     1 0
                               0
                                     0
## 9:
        0
            1
                  0
                       0 0
                               1
                                     0
                                          0
## 10:
        0
             0
                 0
                       1
                          0
                                 0
                                     1
                                          0
## Consistency 0.8975
## Consistency 0.79
## Consistency 0.9175
## # of splits= 400
## Model, % Consistency Met= {pgr <= 80} {age <= median(age)} 0.9175
## Consistency 0.8425
## Consistency 0.8375
## Consistency 0.73
## Consistency 0.6675
## Consistency 0.6675
## Consistency 0.7525
## Consistency 0.8625
## Consistency 0.74
## Consistency 0.6975
```

```
## Consistency 0.87
## Consistency 1
## # of splits= 400
## Model, % Consistency Met= {pgr <= 80} {age <= median(age)} !{grade3} 1
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.3254667
## Subgroup found (FS)
## Minutes overall= 0.3426333
## ***Bootstrap done, B***=9
## H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
## 1: 0.6083361 0.5365407 -0.6760903 -0.9570188 0.01023333 3.284921
## tau, maxdepth= 77.93018 2
## leaf.node control.mean control.size control.se treated.mean treated.size
         2 -7.835404 592.000000 1.991329 7.835404 592.000000
## 1
## 2
           3
                5.466376 94.000000 5.595669 -5.466376 94.000000
## 11
           4 -25.394637 88.000000 5.497696 25.394637
                                                              88.000000
           6 10.129600 156.000000 3.827421 -10.129600 156.000000
           7 -8.983529 431.000000 2.280911
                                                   8.983529 431.000000
## 4
     treated.se
                  diff Nsg depth
## 1
      1.991329 -15.67081 592
## 2
      5.595669 10.93275 94
                                1
## 3 3.827421 20.25920 156
## 4 2.280911 -17.96706 431
## leaf.node control.mean control.size control.se treated.mean treated.size
## 3 6 10.129600 156.000000 3.827421 -10.129600 156.000000
               diff Nsg depth
## treated.se
## 3 3.827421 20.2592 156
## Subgroup found
## [1] "age <= 42" "size <= 40" "age <= 48"
## [1] "age <= 48"
## # of continuous/categorical characteristics 5 2
## Continuous characteristics: age size nodes pgr er
## Categorical characteristics: meno grade3
## CV lambda = 0.003651612
## 7 x 1 sparse Matrix of class "dgCMatrix"
##
                   s0
## age
       -0.0284297067
## meno 0.3320838001
## size 0.0058941277
## grade3 0.0830456319
## nodes 0.0663269703
         -0.0026267022
## pgr
        0.0006369094
## er
## Cox-LASSO selected: age meno size grade3 nodes pgr er
## Cox-LASSO not selected:
## Median cuts after Lasso: age size nodes pgr er
## Categorical after Lasso: meno grade3
## Factors per GRF: age <= 42 size <= 40 age <= 48
## Medians prior to removing if also in GRF: age size nodes pgr er
## ***cMed_flag***=age
## ***cMed_flag***=size
## ***cMed_flag***=age
## ***to_exclude***=TRUE
## ***to_exclude***=TRUE
```

```
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***to_exclude***=FALSE
## ***conf.cont_medians***=nodes
## ***conf.cont_medians***=pgr
## ***conf.cont_medians***=er
## Factors after removing any duplicates also in GRF: nodes pgr er
## ***Factors per lasso after omitting GRF dups***=meno
## ***Factors per lasso after omitting GRF dups***=grade3
## ***Factors per lasso after omitting GRF dups***=nodes <= median(nodes)
## ***Factors per lasso after omitting GRF dups***=pgr <= median(pgr)
## ***Factors per lasso after omitting GRF dups***=er <= median(er)
## Initial GRF cuts included age <= 42 size <= 40 age <= 48
## Factors included per GRF (not in lasso) age <= 42 size <= 40 age <= 48
## # of candidate subgroup factors= 8
## [1] "nodes <= median(nodes)" "pgr <= median(pgr)"</pre>
                                                       "er <= median(er)"</pre>
## [4] "age <= 42"
                              "size <= 40"
                                                       "age <= 48"
## [7] "meno"
                               "grade3"
## LMAX= 8
## Confounders per grf screening q8 q1 q2 q5 q3 q6 q7 q4
## FSconfounders.name
                         vi.cs
## 8
                   q8 0.22417497
## 1
                   q1 0.17373353
## 2
                   q2 0.14888770
## 5
                    q5 0.13268615
## 3
                    q3 0.11571098
## 6
                    q6 0.08551550
## 7
                    q7 0.08246532
                    q4 0.03682585
## Number of unique levels (L) and possible subgroups= 16 65535
## # of subgroups based on # variables > k.max and excluded (per million) 0.064839
## k.max= 3
## Events criteria for control,exp= 10 10
## # of subgroups with events less than criteria: control, experimental 250 418
## # of subgroups with sample size less than criteria 362
## # of subgroups meeting all criteria = 225
## # of subgroups fitted (Cox model estimable) = 225
## *Subgroup Searching Minutes=* 0.01591667
## Number of subgroups meeting HR threshold 30
## # of candidate subgroups (meeting HR criteria) = 30
## Subgroups (1st 10) meeting overall screening thresholds (HR, m1) sorted by HRs= Inf
                        m0 HR L(HR) U(HR) q8.0 q8.1 q1.0 q1.1 q2.0 q2.1 q5.0
      K n E d1 m1
## 1: 2 156 49 14 37.65
                          NA 2.14 1.14 4.00
                                                 0
                                                      0
                                                           0
                                                                0
## 2: 2 148 75 18 17.81 42.58 1.37 0.80 2.35
                                                 0
                                                      0
                                                           0
                                                                0
                                                                     0
                                                                          1
                                                                0
                                                                     0
## 3: 3 143 47 12 64.95
                        NA 2.03 1.04 3.94
                                                 0
                                                      \cap
                                                           0
                                                                          0
                                                                               0
## 4: 1 143 63 18 27.78 66.83 1.55 0.89 2.67
                                                 0
                                                      1
                                                           0
                                                                0
                                                                     0
## 5: 2 135 68 17 17.51 44.88 1.90 1.08 3.33
                                                 0
## 6: 3 127 41 11 64.95
                         NA 1.77 0.88 3.56
                                                                0
                                                 1
                                                      0
                                                         0
## 7: 3 125 67 16 17.51 38.24 1.64 0.92 2.91
                                                 0
                                                      0
                                                           0
                                                                0
                                                                     0
                                                                               0
                                                                          1
## 8: 2 116 53 14 20.76 40.25 1.55 0.84 2.86
                                                 0
                                                      1
                                                           0
                                                                0
                                                                     0
                                                                               0
## 9: 2 111 44 16 34.79 NA 1.75 0.95 3.24
                                                 0
                                                      1
                                                           0
                                                                0
                                                                     0
## 10: 3 99 60 17 17.51 36.40 1.59 0.89 2.82
                                                 0
                                                      0
                                                           0
                                                                0
                                                                     0
## q5.1 q3.0 q3.1 q6.0 q6.1 q7.0 q7.1 q4.0 q4.1
## 1: 0 0 0 1 0 0 1
              0
                   0
                        0
                             0
```

```
0
                  0
                       0
                       0
                            0
## 4:
       0
             0
                  0
## 5:
       0
              0
                  0
                       0
                            1
                                 0
                                     0
                                          0
        0
## 6:
             0
                  0
                       0
                            1
                                 0
                                     0
                                          1
                                               0
##
   7:
        0
             0
                  0
                       0
                            1
                                1
                                     0
## 8:
       0
            0
                     0
                          0
                               0
                                   0
                                        0
                                              0
                  0
                                        1
## 9:
       0
            0
                  0
                     0 0
                               0 0
                                              0
                                   0
                                         0
## 10:
        0
            0
                  1
                       0
                          1
                                 0
                                               0
## Consistency 0.97
## # of splits= 400
## Model, % Consistency Met= {age <= 48} !{age <= 42} 0.97
## Consistency 0.73
## Consistency 0.91
## # of splits= 400
## Model, % Consistency Met= {age <= 48} !{meno} !{age <= 42} 0.91
## Consistency 0.8875
## Consistency 0.9475
## # of splits= 400
## Model, % Consistency Met= {pgr <= median(pgr)} {age <= 48} 0.9475
## Consistency 0.8525
## Consistency 0.875
## Consistency 0.835
## Consistency 0.9225
## # of splits= 400
## Model, % Consistency Met= {grade3} !{age <= 42} 0.9225
## Consistency 0.845
## Consistency 0.85
## Consistency 0.9425
## # of splits= 400
## Model, % Consistency Met= {grade3} {pgr <= median(pgr)} {er <= median(er)} 0.9425
## Consistency 0.9425
## # of splits= 400
## Model, % Consistency Met= {pgr <= median(pgr)} !{meno} !{age <= 42} 0.9425
## Consistency 0.8675
## Consistency 0.565
## Consistency 0.915
## # of splits= 400
## Model, % Consistency Met= !{grade3} {pgr <= median(pgr)} {age <= 48} 0.915
## Consistency 0.9025
## # of splits= 400
## Model, % Consistency Met= {er <= median(er)} {age <= 48} !{age <= 42} 0.9025
## Consistency 0.6225
## Consistency 0.675
## Consistency 1
## # of splits= 400
## Model, % Consistency Met= {pgr <= median(pgr)} {age <= 48} !{age <= 42} 1
## SG focus= Nsg_only
## Subgroup Consistency Minutes= 0.4495167
## Subgroup found (FS)
## Minutes overall= 0.4723167
## ***Bootstrap done, B***=10
## H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search max_sg_est
        ## 1:
print(ansB)
```

```
H_biasadj_1 H_biasadj_2 Hc_biasadj_1 Hc_biasadj_2 tmins_search
## [1,] 1 0.5765919 0.6342285 -0.423406
                                              -0.3271736
                                                            0.0084
## [2,] 2 0.7232828 0.1723476 -0.4346078 -0.3464593
                                                            0.012
## [3,] 3 0.7619819 0.3176834
                                  -0.3713988 -0.2553875
                                                            0.005666667
## [4,] 4 0.3394555 0.2703407
                                  -0.3735976 -0.1709955
                                                           0.01145
## [5,] 5 0.4531755 0.7810623 -0.3841067 -0.4618613 0.003666667
## [6,] 6 0.8799324 0.9842622 -0.5793025 -0.6175556 0.0091
## [7,] 7 0.3242974 0.6546308 -0.3003587 -0.4287666 0.0058
## [8,] 8 0.6732497 0.6862775 -0.1973118 -0.1846732
                                                            0.0056
## [9,] 9 0.6083361 0.5365407 -0.6760903 -0.9570188
                                                            0.01023333
## [10,] 10 0.530357
                                 -0.4321159 -0.4546499
                      0.7034314
                                                            0.01591667
##
        max_sg_est
## [1,] 2.257594
## [2,] 3.385423
## [3,] 3.877166
## [4,] 4.041984
## [5,] 3.296798
## [6,] 2.222482
## [7,] 5.6219
## [8,] 2.407159
## [9,] 3.284921
## [10,] 5.123865
tB.start <- proc.time()[3]</pre>
# Bootstraps
resB <- bootPar({</pre>
   ans <- fsboot_forparallel(boot)</pre>
}, boots = NB, seed = 8316951, counter = "boot", export = fun_arg_list_boot)
tB.now <- proc.time()[3]
tB.min <- (tB.now - tB.start)/60
doParallel::stopImplicitCluster()
cat("Minutes for Boots", c(NB, tB.min), "\n")
## Minutes for Boots 1000 4.869583
cat("Projection per 1000", c(tB.min * (1000/NB)), "\n")
## Projection per 1000 4.869583
cat("Propn bootstrap subgroups found =", c(sum(!is.na(resB$H_biasadj_1))/NB), "\n")
## Propn bootstrap subgroups found = 0.864
# How many timmed out
cat("Number timmed out=", c(sum(is.na(resB$H_biasadj_1) & resB$tmins_search > max.minutes)),
    "\n")
## Number timmed out= 0
H_estimates <- get_dfRes(Hobs = H_obs, seHobs = seH_obs, H1_adj = resB$H_biasadj_2,
   ystar = Ystar_mat, cov_method = "standard", cov_trim = 0, est.scale = "hr")
Hc_estimates <- get_dfRes(Hobs = Hc_obs, seHobs = seHc_obs, H1_adj = resB$Hc_biasadj_2,</pre>
   ystar = Ystar_mat, cov_method = "standard", cov_trim = 0, est.scale = "hr")
print(H_estimates)
           HO
                   sdHO HO_lower HO_upper
                                               H1
                                                       sdH1 H1_lower H1_upper
## 1: 1.951393 0.6130758 1.054192 3.612182 1.495498 0.5088262 0.7676696 2.913383
```

```
print(Hc_estimates)
            НО
                    sdHO HO_lower HO_upper
                                                            sdH1 H1_lower
                                                 H1
## 1: 0.6149954 0.08408495 0.4704264 0.8039925 0.6488922 0.1033531 0.4748936
## H1_upper
## 1: 0.886643
bootit <- list(H_estimates = H_estimates, Hc_estimates = Hc_estimates)</pre>
tall.min <- (tB.now - t.start.all)/60
cat("Overall minutes for analysis", c(tall.min), "\n")
## Overall minutes for analysis 7.423333
if (!is.null(file_out)) save(df.analysis, fs.est, bootit, tall.min, resB, cox.formula.boot,
file = file_out)
## H un-adjusted estimates----: 1.95 (95% CI=1.05,3.61)
## H bias-corrected estimates--: 1.5 (95% CI=0.77,2.91)
## H^c un-adjusted estimates---: 0.61 (95% CI=0.47,0.8)
## H^c bias-corrected estimates: 0.65 (95% CI=0.47,0.89)
```