rsa ml.R

jonah

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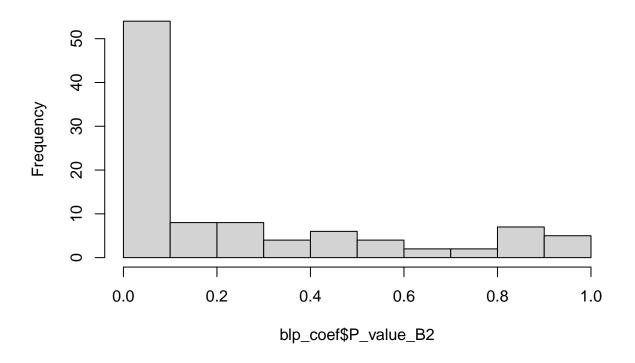
```
library(haven)
library(tidyverse)
## -- Attaching packages ------ tidyverse 1.3.0 --
## v ggplot2 3.3.2 v purrr
                               0.3.4
## v tibble 3.0.4 v dplyr 1.0.2
## v tidyr 1.1.2 v stringr 1.4.0
## v readr 1.4.0 v forcats 0.5.0
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                 masks stats::lag()
library(randomForest)
## randomForest 4.6-14
## Type rfNews() to see new features/changes/bug fixes.
## Attaching package: 'randomForest'
## The following object is masked from 'package:dplyr':
##
##
      combine
## The following object is masked from 'package:ggplot2':
##
      margin
library(ggthemes)
# dataset based on:
# Generating Skilled Self-Employment in Developing Countries: Experimental Evidence
# from Uganda (Blattman et al, 2014)
# outcome of interest is profit in last four weeks capped at 99th percentile;
# treatment is randomised assignment to the Program (intent to treat)
```

```
# method is based on:
# Generic Machine Learning Inference on Heterogenous Treatment Effects in Randomized
# Experiments (Chernozhukov et al, 2017)
# Note: This is merely preliminary EDA and will be refined once more of the data is
# understood
# Final code will be run with neural nets, LASSO, and boosted trees
#some data cleaning
x<-read_dta('https://www.dropbox.com/s/yxgigmtcrut9fii/yop_analysis.dta?dl=1') %>%
  filter(e2==1) %>% #filter results only from endline survey1
  select(assigned,S_K,S_H,S_P_m,
         admin_cost_us,groupsize_est_e,grantsize_pp_US_est3,group_existed,group_age,ingroup_hetero,ingr
         lowskill7da_zero,lowbus7da_zero,skilledtrade7da_zero,highskill7da_zero,acto7da_zero,aghours7da
         age,urban,ind_found_b,risk_aversion,inschool,
         D_1,D_2,D_3,D_4,D_5,D_6,D_7,D_8,D_9,D_10,D_11,D_12,D_13,
         profits4w_real_p99_e)
#compute proportion of all NA values
sum(x\%>\%is.na())/(ncol(x)*nrow(x))
## [1] 0.007195332
#compute proportion of missing outcome values
sum(x$profits4w_real_p99_e%>%is.na())/nrow(x)
## [1] 0.302204
#eliminate NA values (robustness checks with lee and manski bounds to be added)
df<-x[which(complete.cases(x)),]</pre>
# create empty dataframes to store values
n_split<-100
n group<-5
blp_coef<-data.frame(B1=1:n_split,SE_B1=1:n_split,</pre>
                     P_value_B1=1:n_split,
                     B2=1:n_split,SE_B2=1:n_split,
                     P_value_B2=1:n_split)
gate_coef<-matrix(ncol = n_group*3,nrow = n_split) %>% as.data.frame()
colnames(gate_coef)<-paste(c("G",'SE_G','P_value'), rep(1:n_group, each=3), sep = "")</pre>
gate_diff<-data.frame(diff=1:n_split,SE=1:n_split,P_value=1:n_split)</pre>
clan<-matrix(ncol = 6,nrow = n_split*(ncol(df)-15)) %>% as.data.frame()
colnames(clan)<-c('estimate','SE','lower_conf','upper_conf','P_value','var')</pre>
clan_os<-matrix(ncol = 7,nrow = n_split*(ncol(df)-15)) %>% as.data.frame()
colnames(clan_os)<-c('g1','g1_lower_conf','g1_upper_conf','gk','gk_lower_conf','gk_upper_conf','var')</pre>
#f<-which(sapply(df, class) == "factor")
```

```
\#mcol < -ncol(df[,-f])
mcol<-ncol(df)
gate_mean<-matrix(ncol = mcol*n_group,nrow = n_split)</pre>
colnames(gate_mean) <- rep(colnames(df[,]),n_group)</pre>
#split data
set.seed(55,sample.kind = 'Rounding')
## Warning in set.seed(55, sample.kind = "Rounding"): non-uniform 'Rounding'
## sampler used
for(i in 1:n_split){
  #randomly split data into main and auxiliary
  random<-runif(nrow(df))</pre>
  main_ind<-which(random>0.5)
  aux_ind<-which(random<0.5)</pre>
  aux df<-df[aux ind,]</pre>
  main_df<-df[main_ind,]</pre>
  # train data on auxiliary sample
  rftreat<-randomForest(profits4w_real_p99_e~., data = (aux_df%>%filter(assigned==1)),
                         ntree=500, nodesize=5)
  rfbase<-randomForest(profits4w_real_p99_e~., data = (aux_df%-%filter(assigned==0)),
                        ntree=500,nodesize=5)
  # predict baseline and treatment outcomes on main sample
  B<-predict(rfbase,main_df)</pre>
  treat<-predict(rftreat,main_df)</pre>
  # specifying regression variables
  S<-treat-B #CATE: what the algorithm predicts is an individual's treatment effect
  ES<-mean(S) # the average predicted treatment effect
  p<-mean(main_df$assigned) #take mean as propensity score</pre>
  x<-S-ES #excess CATE: how far one's predicted treatment effect is from the mean
  w<-main_df$assigned-p #weighted treatment var</pre>
  #derive Best Linear Predictor from main sample
  blp<-lm(profits4w_real_p99_e~B+w+I((w*x)),data=cbind(main_df,B,S,x,w))</pre>
  blp_coef[i,]<-c(blp$coefficients[3],summary(blp)$coefficients[3:4,c(2,4)][1,],
                  blp$coefficients[4], summary(blp)$coefficients[3:4,c(2,4)][2,])
  #Group Average Treatment Effect
  qt<-quantile(S,seq(0,1,length.out = n_group+1))
  diff<-cbind(main_df,B,S,w) %>%
    filter(S<=qt[2]|S>qt[n_group]) %>%
    mutate(G=ifelse((S>qt[n_group]),1,0))
  gate_diff[i,]<-summary(lm(profits4w_real_p99_e~B+I(w*G),data = diff))$coefficients[3,c(1,2,4)]</pre>
  for(k in 1:n_group){
    G<-ifelse(S>qt[k] & S<=qt[k+1],1,0)
    gate<-lm(profits4w_real_p99_e~B+I(w*G),data = cbind(main_df,B,S,x,w,G))</pre>
```

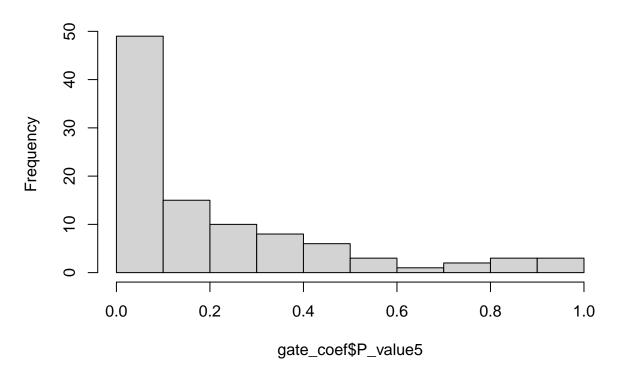
```
gate_coef[i,((3*k)-2):(3*k)]<-summary(gate)$coefficients[3,c(1,2,4)]
    # data preparation for later
    gate_mean[i,((k*mcol)-(mcol-1)):(k*mcol)]<-apply(main_df[which(G==1),],2,mean)</pre>
  #Classification Analysis (CLAN)
  diff2<-diff%>%select(-profits4w_real_p99_e,-assigned,-B,-w,-G,-ind_found_b,
                        -D_1, -D_2, -D_3, -D_4, -D_5, -D_6, -D_7, -D_8, -D_9, -D_{10}, -D_{11}, -D_{12}, -D_{13}
  n<-ncol(diff2)</pre>
  for (j in (1:n)){
    #two sample t test
    b<-t.test(diff2%>%filter(S>qt[n group])%>%.[,j],diff2%>%filter(S<qt[2])%>%.[,j],
              alternative="two.sided", var.equal=F)
    clan[((i*n)-n+j),1]<-b*estimate[1]-b$*estimate[2]
    clan[((i*n)-n+j),2]<-b$stderr
    clan[((i*n)-n+j),3:4]<-b$conf.int
    clan[((i*n)-n+j),5]<-b$p.value
    clan[((i*n)-n+j),6] < -colnames(diff2)[j]
    #one sample t test
    d<-t.test(diff2%>%filter(S<=qt[2])%>%.[,j])
    clan_os[((i*n)-n+j),1]<-d*estimate
    clan_os[((i*n)-n+j),2:3]<-d$conf.int
    d<-t.test(diff2%>%filter(S>qt[n group])%>%.[,j])
    clan_os[((i*n)-n+j),4]<-d*estimate
    clan_os[((i*n)-n+j),5:6]<-d$conf.int
    clan_os[((i*n)-n+j),7]<-colnames(diff2)[j]
  }
}
# check distribution of p value
hist(blp_coef$P_value_B2)
```

Histogram of blp_coef\$P_value_B2



hist(gate_coef\$P_value5)

Histogram of gate_coef\$P_value5

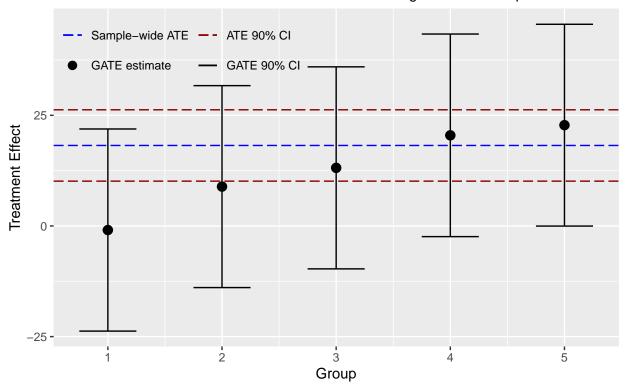


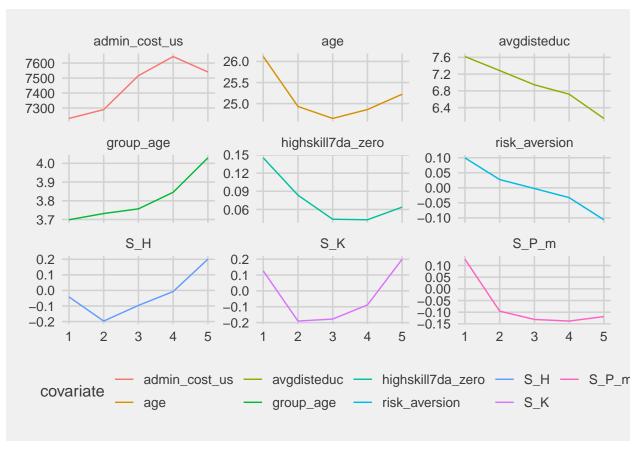
```
# obtain median values
# data for each column does not correspond to the same split
apply(blp_coef,2,median) #median for Best Linear Predictor
##
                  SE_B1 P_value_B1
                                                   SE_B2 P_value_B2
## 11.71240001 6.17144994 0.06023404 0.30983913 0.18788793 0.08355674
apply(gate_coef,2,median) #median for Grouped Average Treatment Effect (GATE)
                 SE_G1
                        P_value1
                                       G2
                                               SE_G2
                                                      P value2
## -0.9219174 13.8653376 0.4793581 8.8925038 13.8523339 0.4551271 13.1287592
       SE G3
              P value3
                             G4
                                     SE G4
                                            P value4
##
    P value5
  0.1041146
apply(gate_diff,2,median) #median for difference in GATE between G1 and Gk
##
       diff
                      P_value
## 22.376135 14.907757 0.122659
# plot GATE with confidence bands
ci<-data.frame(group = 1:5,estimate = NA,SE = NA,P_value=NA)</pre>
```

```
for (k in 1:5){
ci[k,2:4] < -apply(gate_coef,2,median)[((k*3)-2):(k*3)]
}
labels <- c(point = "GATE estimate", error = "GATE 90% CI",</pre>
            blue = "Sample-wide ATE", darkred = "ATE 90% CI")
breaks <- c("blue", "darkred", "point", "error")</pre>
ggplot(ci, aes(x = group, y = estimate)) +
  geom_point(aes(color = "point"), size = 3) +
  geom_errorbar(width = .5, aes(
   ymin = estimate - (1.647 * SE),
   ymax = estimate + (1.647 * SE),
   color = "error"
  )) +
  scale_color_manual(values = c(
   point = "black",
   error = "black",
   blue = "blue",
   darkred = "darkred"
  ), labels = labels, breaks = breaks) +
  labs(
   title = "GATE with confidence bands",
   subtitle = "Point estimates and confidence bands are derived using median of all splits",
   x = "Group",
   y = "Treatment Effect",
   color = NULL, linetype = NULL, shape = NULL
  ) +
  geom_hline(
   data = data.frame(yintercept = 18.19+c(-1,1)*(1.646*4.898)),
   aes(yintercept = yintercept, color = "darkred"), linetype = "longdash"
  ) + # ATE and CI from original paper
  geom_hline(
   data = data.frame(yintercept = 18.19),
   aes(yintercept = yintercept, color = "blue"), linetype = "longdash"
  guides(color = guide_legend(override.aes = list(
   shape = c(NA, NA, 16, NA),
   linetype = c("longdash", "longdash", "blank", "solid")
  ), nrow = 2, byrow = TRUE)) +
  theme(legend.position = c(0, 1),
       legend.justification = c(0, 1),
        legend.background = element_rect(fill = NA),
        legend.key = element_rect(fill = NA)) +
# examining heterogeneity
for(k in 1:n_group) {
 nam <- paste("gate_mean", k, sep = "")</pre>
  assign(nam, gate_mean[,((k*mcol)-(mcol-1)):(k*mcol)])
}
```

GATE with confidence bands

Point estimates and confidence bands are derived using median of all splits





```
#classification analysis (CLAN)
col<-clan$var %>% unique
clan_os_med<-matrix(ncol = 7,nrow = length(col)) %>% as.data.frame()
colnames(clan_os_med)<-c('g1','g1_lower_conf','g1_upper_conf','gk','gk_lower_conf','gk_upper_conf','var</pre>
clan_med<-matrix(ncol = 6,nrow = length(col)) %>% as.data.frame()
colnames(clan_med)<-c('estimate','SE','lower_conf','upper_conf','P_value','var')</pre>
#obtain medians of means and confidence interval for G1 and GK (1 sample t test)
for (i in col){
  clan_os_med[which(col==i),]<-clan_os %>% filter(var==i) %>% select(-var) %>%
    apply(2,median,na.rm=T) %>%
    as.list() %>% as.data.frame() %>% mutate(var=i)
}
# obtain medians for difference in means for G1 and GK (2 sample t test)
for (i in col){
  clan_med[which(col==i),]<-clan %>% filter(var==i) %>% select(-var) %>% apply(2,median) %>%
  as.list() %>% as.data.frame() %>% mutate(var=i)
}
clan_med[,c(6,1,2,3,4,5)] %>% filter(var!='S',P_value<=0.05) %>%
  arrange(-desc(P_value))
```

var estimate SE lower_conf upper_conf P_value

```
## 1
               S_P_m -0.2630031 0.10246666 -0.457633983 -0.0620133243 0.003258999
## 2
               urban 0.1129148 0.04285560 0.032098859 0.1963831716 0.005223420
## 3
       risk aversion -0.2123593 0.10699286 -0.418020250 0.0005123978 0.006294560
## 4 ingroup_dynamic 0.2967003 0.10454470 0.080664569 0.5091663230 0.007139508
                 S_H 0.2178325 0.10896574 0.001175123 0.4324383948 0.011417867
## 5
## 6
         avgdisteduc -1.4349099 0.65722910 -2.723171482 -0.2028861262 0.013389239
## 7
        acto7da zero 1.4120640 0.58653225 0.258238219 2.6566699792 0.013924192
## 8
          zero_hours 0.1239901 0.05033799 0.023329934 0.2243181488 0.014748294
## 9
       ingroup_hetero 0.2446042 0.10843910 0.032249766 0.4633295549 0.015551000
                 age -0.7354318 0.55885226 -1.815550785 0.3621369451 0.036670426
## 10
## 11 aghours7da_zero -1.7647849 1.14683252 -3.927053655 0.5721026982 0.047526597
     chores7da_zero -3.6623076 1.82046628 -7.140949831 0.1275594687 0.048649724
     lowbus7da_zero -0.6532051 0.78865306 -2.156656801 0.8516187287 0.048990597
```