

# SISCER Module 15 Practice 2: Matching/Weighting and Sensitivity Analysis

Ting Ye and Qingyuan Zhao

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## 1

```
library(tidyverse)

## -- Attaching packages ----- tidyverse 1.3.1 --
## v ggplot2 3.3.6      v purrr  0.3.4
## v tibble  3.1.7      v dplyr  1.0.9
## v tidyr   1.2.0      v stringr 1.4.0
## v readr   2.1.2      v forcats 0.5.1

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()

library(ggplot2)
library(tableone) # for creating table 1
source("optmatch.R")

## Loading required package: lattice

##
## Attaching package: 'caret'

## The following object is masked from 'package:purrr':
##
##   lift

df<-read.csv("nhanesi_class_dataset.csv",stringsAsFactors = TRUE)
df<-df[,-1]
summary(df)

##   physically.inactive years.lived.since.1971.up.to.1992      sex
##   Mode :logical      Min.   : 1.00                      Female:962
##   FALSE:1481         1st Qu.:14.00                      Male  :989
##   TRUE :470          Median :21.00
##                   Mean   :17.32
##                   3rd Qu.:21.00
##                   Max.   :21.00
##
##   smoking.status income.poverty.ratio age.at.interview      race
##   Current:604     Min.   :0.130         Min.   :45.00      Nonwhite: 274
##   Former :416     1st Qu.:1.215         1st Qu.:52.00      White   :1677
```

```

## Never :931 Median :2.280 Median :60.00
## Mean :2.706 Mean :59.84
## 3rd Qu.:3.510 3rd Qu.:67.00
## Max. :9.980 Max. :74.00
## NA's :544
## education working.last.three.months married
## 0-8 :739 Mode :logical Mode :logical
## 12 :439 FALSE:1063 FALSE:505
## 9-11 :415 TRUE :888 TRUE :1446
## College Grad:151
## Missing : 12
## Some College:195
##
## alcohol.consumption dietary.adequacy
## <1 time per month :377 Min. :0.000
## 1-4 times per month :410 1st Qu.:3.000
## 2+ times per week :205 Median :4.000
## Just about everyday/everyday:300 Mean :3.565
## Never :659 3rd Qu.:5.000
## Max. :5.000
## NA's :479

```

```

# (a) create missing indicators for income.poverty.ratio and dietary.adequacy and impute by observed
df<-df %>% mutate(income.poverty.ratio.missingind=1*(is.na(income.poverty.ratio)),
                 dietary.adequacy.missingind=1*(is.na(dietary.adequacy)))
df<-df %>% mutate(income.poverty.ratio=ifelse(is.na(income.poverty.ratio),
                                             mean(income.poverty.ratio,na.rm=TRUE),income.poverty.ratio),
                 dietary.adequacy=ifelse(is.na(dietary.adequacy),
                                             mean(dietary.adequacy,na.rm=TRUE),dietary.adequacy))
df<- df %>% mutate(physically.inactive=ifelse(physically.inactive,1,0))

```

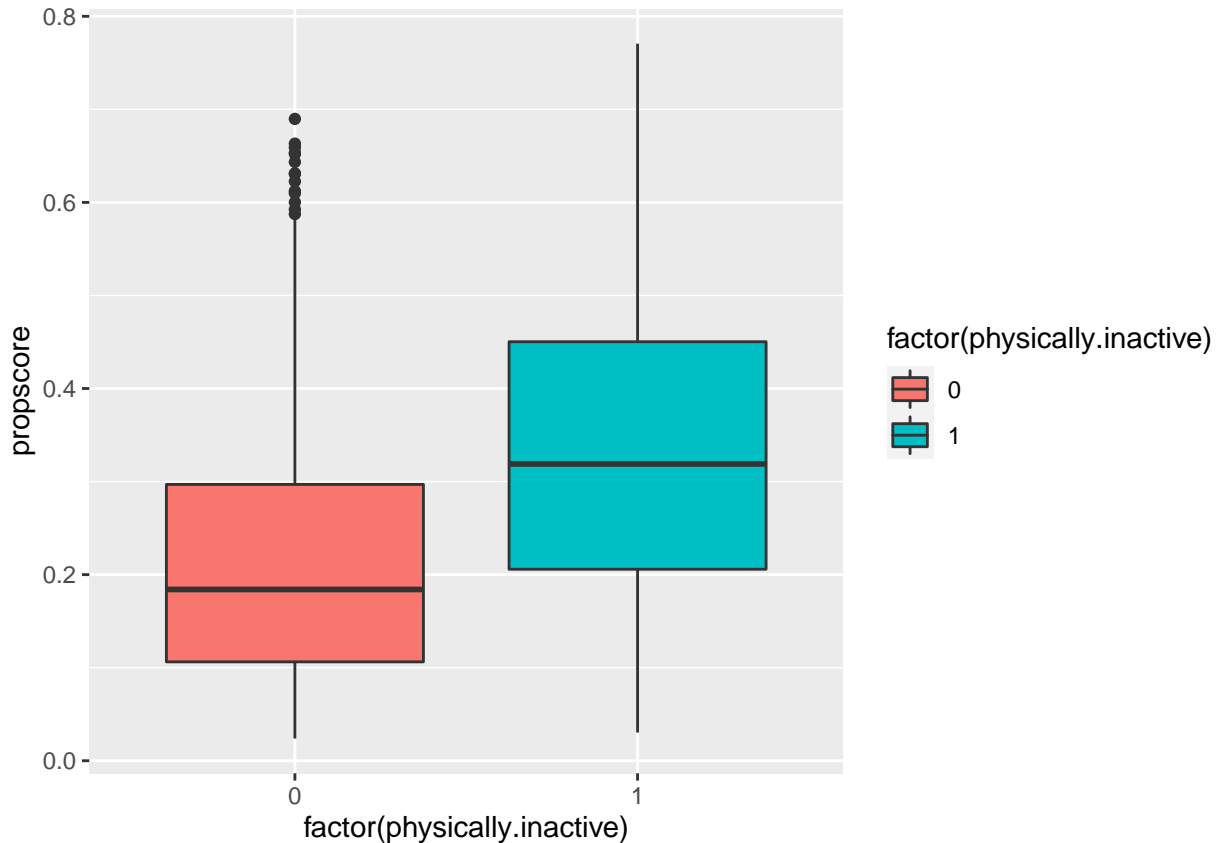
## 2

```

# PS model should include all confounders and missing indicators
ps.formula=physically.inactive~sex+smoking.status+income.poverty.ratio+
age.at.interview+race+education+working.last.three.months+married+
alcohol.consumption+dietary.adequacy+income.poverty.ratio.missingind+
dietary.adequacy.missingind
propscore.model=glm(ps.formula,family=binomial,data=df)
propscore<-propscore.model$fitted.values

ggplot(data=df, aes(x = factor(physically.inactive), y = propscore,
                    fill = factor(physically.inactive))) + geom_boxplot()

```



```
# excluding subjects lack of overlap
exclud1<-which(prop score>max(prop score[df$physically.inactive==0]))
exclud2<-which(prop score<min(prop score[df$physically.inactive==1]))
cbind(prop score,df)[c(exclud1,exclud2),] # check the subjects to be excluded
```

##	prop score	physically.inactive	years.lived.since.1971.up.to.1992	sex	
##	272	0.74720177	1	6 Female	
##	328	0.77056752	1	14 Female	
##	371	0.70173323	1	1 Male	
##	815	0.74575568	1	21 Female	
##	123	0.03025210	0	14 Female	
##	429	0.02386385	0	21 Male	
##	602	0.02913807	0	21 Male	
##	754	0.02532975	0	19 Male	
##	1282	0.02928994	0	21 Female	
##	smoking.status	income.poverty.ratio	age.at.interview	race	education
##	272	Current	2.70634	70 Nonwhite	0-8
##	328	Current	2.70634	63 White	College Grad
##	371	Current	0.73000	55 Nonwhite	0-8
##	815	Current	0.40000	68 Nonwhite	0-8
##	123	Never	9.05000	65 White	Some College
##	429	Former	8.89000	66 White	12
##	602	Former	8.86000	45 White	Some College
##	754	Never	9.98000	62 White	Some College
##	1282	Never	8.34000	57 White	9-11
##	working.last.three.months	married	alcohol.consumption		
##	272	FALSE	FALSE	Never	

```

## 328          FALSE  FALSE Just about everyday/everyday
## 371          FALSE  FALSE                               Never
## 815          FALSE  FALSE                               Never
## 123          TRUE   TRUE  Just about everyday/everyday
## 429          TRUE   TRUE           1-4 times per month
## 602          TRUE   TRUE           2+ times per week
## 754          TRUE   TRUE                               Never
## 1282         TRUE   TRUE           <1 time per month
##      dietary.adequacy income.poverty.ratio.missingind
## 272          4          1
## 328          4          1
## 371          1          0
## 815          1          0
## 123          4          0
## 429          5          0
## 602          5          0
## 754          3          0
## 1282         5          0
##      dietary.adequacy.missingind
## 272          0
## 328          0
## 371          0
## 815          0
## 123          0
## 429          0
## 602          0
## 754          0
## 1282         0

```

```
length(c(exclud1,exclud2)) # the number of units excluded
```

```
## [1] 9
```

```
df<-df[-c(exclud1,exclud2),]
```

### 3

```
##(d) Perform matching using optmatch
```

```
# Specify the model for propensity score (fitted by logistic regression)
```

```
ps.formula
```

```
## physically.inactive ~ sex + smoking.status + income.poverty.ratio +
```

```
## age.at.interview + race + education + working.last.three.months +
```

```
## married + alcohol.consumption + dietary.adequacy + income.poverty.ratio.missingind +
```

```
## dietary.adequacy.missingind
```

```
# Specify all the variables you want to use in the Mahalanobis distance
```

```
# on the right handside of ~
```

```
mahal.formula=physically.inactive~sex+smoking.status+age.at.interview
```

```
# Perform paired matching
```

```
# Print standardized difference (both before and after matching) and the Love plot
```

```
match_res1<-optmatch_caliper(df,nocontrols.per.match = 1, calipersd=0.5,
```

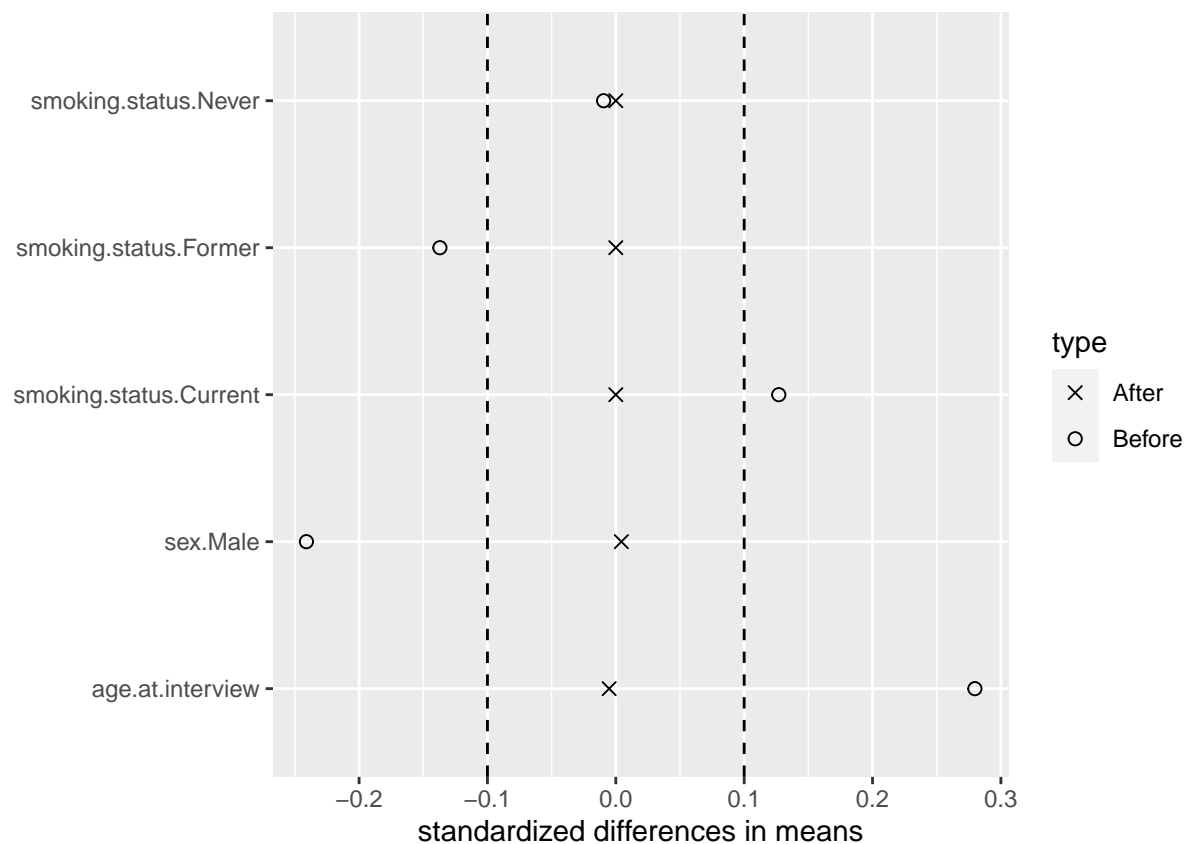
```
ps.formula=ps.formula,mahal.formula=mahal.formula)
```

```
##
```

```
## Attaching package: 'MASS'

## The following object is masked from 'package:dplyr':
##
##   select
##
##           stand.diff.before stand.diff.after
## sex.Female           0.24           0.00
## sex.Male             -0.24           0.00
## smoking.status.Current  0.13           0.00
## smoking.status.Former  -0.14           0.00
## smoking.status.Never   -0.01           0.00
## age.at.interview       0.28           -0.01
##
##           treatmean.after controlmean.before controlmean.after
## sex.Female           0.58           0.46           0.59
## sex.Male             0.42           0.54           0.41
## smoking.status.Current  0.35           0.29           0.35
## smoking.status.Former  0.17           0.23           0.17
## smoking.status.Never   0.47           0.48           0.47
## age.at.interview       61.69           59.25           61.73
```

```
match_res1$p
```



```
# Matched samples
df_treated<-match_res1$datatemp[match_res1$treated.subject.index,] # the treated data
df_control<-match_res1$datatemp[match_res1$matched.control.subject.index,] # the matched controls
df_matched<-rbind(df_treated,df_control)
```

```
# create a table 1
```

```

tb1<-CreateTableOne(vars=names(df)[-c(1:2)],strata="physically.inactive",data=df_matched)
tab1Mat<-print(tb1,showAllLevels = TRUE,smd=FALSE,quote=FALSE, noSpaces = TRUE, printToggle = FALSE)
tab1Mat # note this SMD is different than the standardized mean difference discussed in the lecture,

```

```

##                               Stratified by physically.inactive
##                               level
## n                               ""
## sex (%)                         "Female"
##                               "Male"
## smoking.status (%)             "Current"
##                               "Former"
##                               "Never"
## income.poverty.ratio (mean (SD)) ""
## age.at.interview (mean (SD))   ""
## race (%)                       "Nonwhite"
##                               "White"
## education (%)                  "0-8"
##                               "12"
##                               "9-11"
##                               "College Grad"
##                               "Missing"
##                               "Some College"
## working.last.three.months (%)  "FALSE"
##                               "TRUE"
## married (%)                   "FALSE"
##                               "TRUE"
## alcohol.consumption (%)       "<1 time per month"
##                               "1-4 times per month"
##                               "2+ times per week"
##                               "Just about everyday/everyday"
##                               "Never"
## dietary.adequacy (mean (SD))   ""
## income.poverty.ratio.missingind (mean (SD)) ""
## dietary.adequacy.missingind (mean (SD)) ""
##                               Stratified by physically.inactive
##                               0           1
## n                               "466"           "466"
## sex (%)                         "273 (58.6)"      "272 (58.4)"
##                               "193 (41.4)"      "194 (41.6)"
## smoking.status (%)             "165 (35.4)"    "165 (35.4)"
##                               "80 (17.2)"       "80 (17.2)"
##                               "221 (47.4)"      "221 (47.4)"
## income.poverty.ratio (mean (SD)) "2.36 (1.29)"    "2.20 (1.51)"
## age.at.interview (mean (SD))   "61.73 (8.75)"  "61.69 (8.79)"
## race (%)                       "73 (15.7)"     "96 (20.6)"
##                               "393 (84.3)"     "370 (79.4)"
## education (%)                  "205 (44.0)"    "229 (49.1)"
##                               "85 (18.2)"     "78 (16.7)"
##                               "81 (17.4)"     "86 (18.5)"
##                               "46 (9.9)"       "39 (8.4)"
##                               "2 (0.4)"        "3 (0.6)"
##                               "47 (10.1)"      "31 (6.7)"
## working.last.three.months (%)  "346 (74.2)"    "350 (75.1)"
##                               "120 (25.8)"     "116 (24.9)"

```

```

## married (%) "152 (32.6)" "174 (37.3)"
## "314 (67.4)" "292 (62.7)"
## alcohol.consumption (%) "95 (20.4)" "93 (20.0)"
## "84 (18.0)" "81 (17.4)"
## "45 (9.7)" "42 (9.0)"
## "65 (13.9)" "62 (13.3)"
## "177 (38.0)" "188 (40.3)"
## dietary.adequacy (mean (SD)) "3.40 (1.26)" "3.28 (1.33)"
## income.poverty.ratio.missingind (mean (SD)) "0.35 (0.48)" "0.27 (0.45)"
## dietary.adequacy.missingind (mean (SD)) "0.32 (0.47)" "0.23 (0.42)"
## Stratified by physically.inactive
## p test
## n "" ""
## sex (%) "1.000" ""
## "" ""
## smoking.status (%) "1.000" ""
## "" ""
## "" ""
## income.poverty.ratio (mean (SD)) "0.089" ""
## age.at.interview (mean (SD)) "0.937" ""
## race (%) "0.061" ""
## "" ""
## education (%) "0.323" ""
## "" ""
## "" ""
## "" ""
## "" ""
## "" ""
## working.last.three.months (%) "0.821" ""
## "" ""
## married (%) "0.149" ""
## "" ""
## alcohol.consumption (%) "0.965" ""
## "" ""
## "" ""
## "" ""
## dietary.adequacy (mean (SD)) "0.168" ""
## income.poverty.ratio.missingind (mean (SD)) "0.013" ""
## dietary.adequacy.missingind (mean (SD)) "0.002" ""

```

```

# the latter uses the pre-matching data to estimate the pooled standard error
write.csv(tab1Mat, file = "Table1_nhanesi.csv") # saving this to your computer, so it is easy to paste
head(df)

```

```

## physically.inactive years.lived.since.1971.up.to.1992 sex smoking.status
## 1 1 8 Female Never
## 2 0 16 Male Current
## 3 0 21 Male Current
## 4 0 13 Male Never
## 5 0 21 Female Current
## 6 0 17 Male Current
## income.poverty.ratio age.at.interview race education
## 1 2.70634 74 Nonwhite 0-8

```

```
## 2          1.52000          74    White          9-11
## 3          3.11000          53 Nonwhite          12
## 4          9.18000          65    White College Grad
## 5          2.62000          47 Nonwhite          9-11
## 6          2.62000          53 Nonwhite          9-11
##   working.last.three.months married          alcohol.consumption
## 1                FALSE    FALSE                Never
## 2                FALSE    TRUE Just about everyday/everyday
## 3                TRUE     TRUE          1-4 times per month
## 4                TRUE    FALSE          2+ times per week
## 5                TRUE     TRUE                Never
## 6                TRUE     TRUE Just about everyday/everyday
##   dietary.adequacy income.poverty.ratio.missingind dietary.adequacy.missingind
## 1          2          1          0
## 2          5          0          0
## 3          3          0          0
## 4          2          0          0
## 5          4          0          0
## 6          2          0          0
```

```
catVars<-c("cm1ethrace","cm1lage_c","cm1edu",
           "cm1relf","cm1hhinc_c","cm1bsex",
           "cf1ethrace","cf1lage_c","cf1edu")
tb1<-CreateTableOne(vars=names(df),data=df,factorVars = catVars)
```

```
## Warning in ModuleReturnVarsExist(factorVars, data): The data frame does not
## have: cm1ethrace cm1lage_c cm1edu cm1relf cm1hhinc_c cm1bsex cf1ethrace cf1lage_c
## cf1edu Dropped
```

```
tbl1Mat<-print(tb1,showAllLevels = TRUE,quote=FALSE, noSpaces = TRUE, printToggle = FALSE)
```

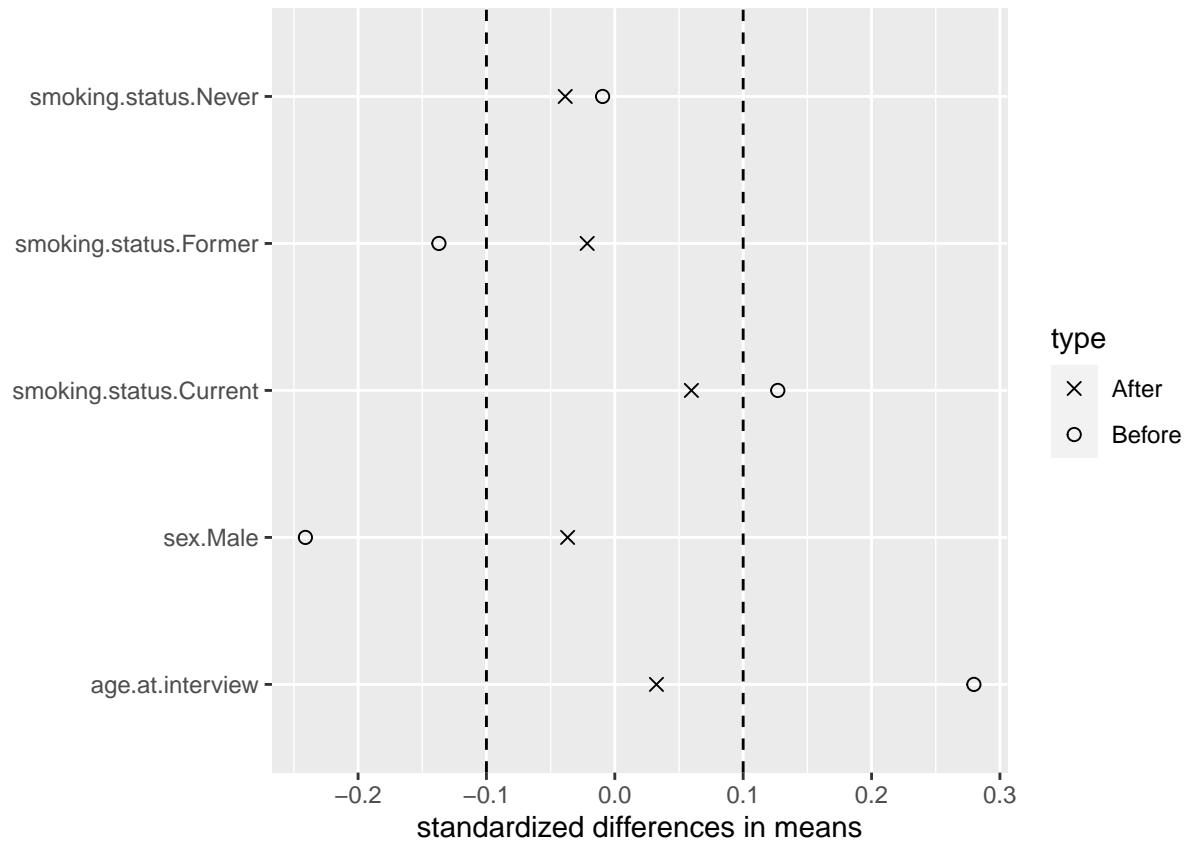
## 4

```
# Consider matching 2 controls to each treated unit. Still found adequate balance,
# all absolute standardized diff < 0.1
match_res2<-optmatch_caliper(df,nocontrols.per.match = 2, calipersd=0.5,
                             ps.formula=ps.formula,mahal.formula=mahal.formula)
```

```
##          stand.diff.before stand.diff.after
## sex.Female          0.24          0.04
## sex.Male           -0.24         -0.04
## smoking.status.Current          0.13          0.06
## smoking.status.Former         -0.14         -0.02
## smoking.status.Never         -0.01         -0.04
## age.at.interview          0.28          0.03
##          treatmean.after controlmean.before controlmean.after
## sex.Female          0.58          0.46          0.57
## sex.Male           0.42          0.54          0.43
## smoking.status.Current          0.35          0.29          0.33
## smoking.status.Former          0.17          0.23          0.18
## smoking.status.Never          0.47          0.48          0.49
## age.at.interview          61.69          59.25          61.40
```

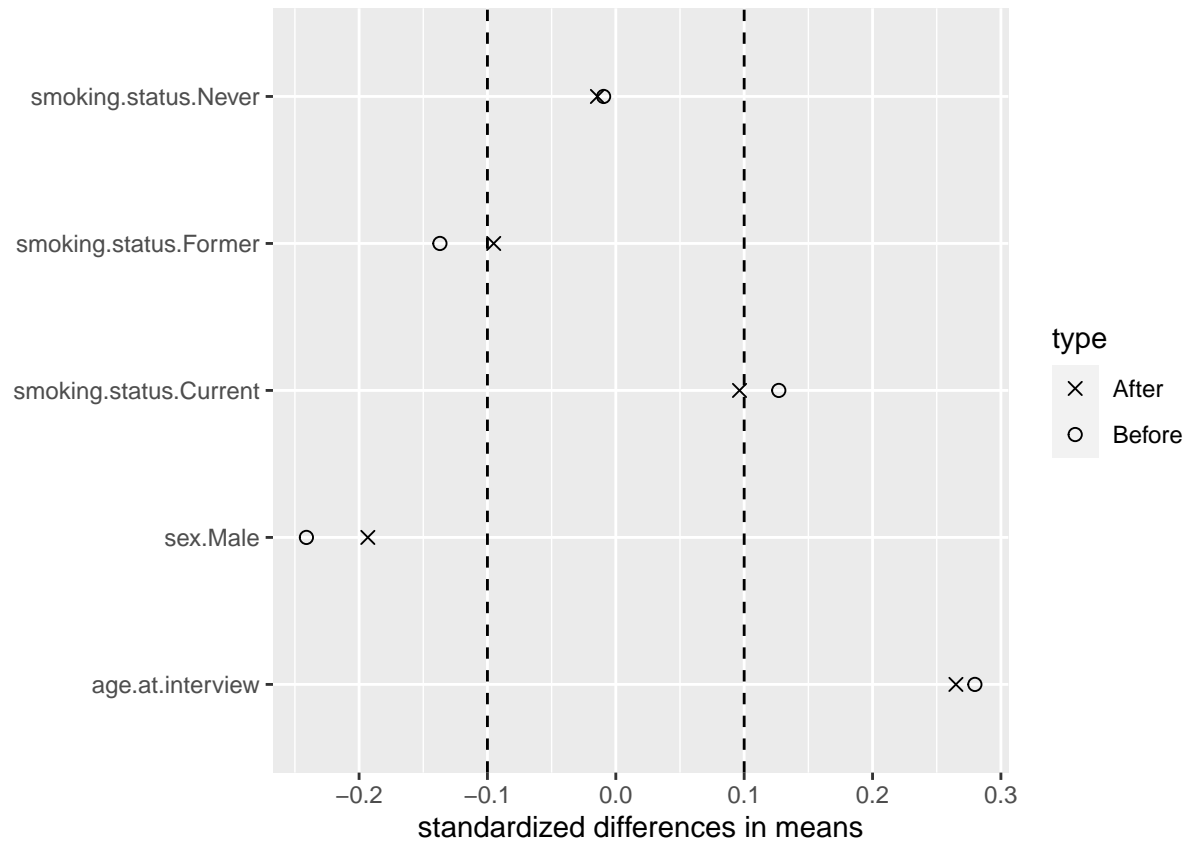
```
match_res2$p
```





```
#consider matching 3 controls to each treated unit. Not adequate balance
match_res3<-optmatch_caliper(df,nocontrols.per.match = 3, calipersd=0.5,
                             ps.formula=ps.formula,mahal.formula=mahal.formula)
```

```
##          stand.diff.before stand.diff.after
## sex.Female          0.24          0.19
## sex.Male           -0.24         -0.19
## smoking.status.Current          0.13          0.10
## smoking.status.Former         -0.14         -0.10
## smoking.status.Never          -0.01         -0.01
## age.at.interview          0.28          0.26
##          treatmean.after controlmean.before controlmean.after
## sex.Female          0.58          0.46          0.49
## sex.Male           0.42          0.54          0.51
## smoking.status.Current          0.35          0.29          0.31
## smoking.status.Former          0.17          0.23          0.21
## smoking.status.Never          0.47          0.48          0.48
## age.at.interview        61.69          59.25          59.38
match_res3$p
```



## 5

I chose 1-v-2 match. Because it uses more controls and all standardized diff is below 0.1.

## 6

```
# For paired matching
df_treated<-match_res1$datatemp[match_res1$treated.subject.index,] # the treated data
df_control<-match_res1$datatemp[match_res1$matched.control.subject.index,] # the matched controls
df_matched<-rbind(df_treated,df_control)
# Wilcoxon signed rank test, a randomization test.
# This is only applicable to matched pairs
# Exact = FALSE implements the large sample approximation (the familiar chi-square test)
# which is fine for most samples of practical size.
library(exactRankTests)

## Package 'exactRankTests' is no longer under development.
## Please consider using package 'coin' instead.
wilcox.exact(df_treated$years.lived.since.1971.up.to.1992,
             df_control$years.lived.since.1971.up.to.1992,
             paired=TRUE,conf.int=TRUE,exact = TRUE)

##
## Exact Wilcoxon signed rank test
##
```

```
## data: df_treated$years.lived.since.1971.up.to.1992 and df_control$years.lived.since.1971.up.to.1992
## V = 16134, p-value = 3.206e-11
## alternative hypothesis: true mu is not equal to 0
## 95 percent confidence interval:
## -4.5 -2.5
## sample estimates:
## (pseudo)median
## -3.25
```

```
# this is the confidence interval about location parameter mu, not difference in means
```

```
# Regression method, adjusting for all match set indicators
```

```
matched.reg.model=lm(years.lived.since.1971.up.to.1992~physically.inactive+matchvec+
sex+smoking.status+income.poverty.ratio+age.at.interview+race+
education+working.last.three.months+married+alcohol.consumption+
dietary.adequacy+income.poverty.ratio.missingind+
dietary.adequacy.missingind,data=df_matched)
```

```
coef(matched.reg.model)[2] # Point estimate of treatment effect
```

```
## physically.inactive
## -1.916665
```

```
confint(matched.reg.model)[2,] # Confidence interval
```

```
## 2.5 % 97.5 %
## -2.646187 -1.187144
```

```
# For 1-v-2 matching
```

```
df_treated2<-match_res2$datatemp[match_res2$treated.subject.index,] # the treated data
df_control2<-match_res2$datatemp[match_res2$matched.control.subject.index,] # the matched controls
df_matched2<-rbind(df_treated2,df_control2)
```

```
# Regression method, adjusting for all match set indicators
```

```
matched.reg.model2=lm(years.lived.since.1971.up.to.1992~physically.inactive+matchvec+
sex+smoking.status+income.poverty.ratio+age.at.interview+race+
education+working.last.three.months+married+alcohol.consumption+
dietary.adequacy+income.poverty.ratio.missingind+
dietary.adequacy.missingind,data=df_matched2)
```

```
coef(matched.reg.model2)[2] # Point estimate of treatment effect
```

```
## physically.inactive
## -1.415638
```

```
confint(matched.reg.model2)[2,] # Confidence interval
```

```
## 2.5 % 97.5 %
## -2.2782274 -0.5530493
```

## 7

Potential unmeasured confounders: stress, ...

## 8

```
#### Sensitivity Analysis for Signed Rank Statistic
#### Assumes that a one-sided test of whether the treatment has
#### a positive effect is being done.
#### diff is the difference between the treated and control units
#### gamma is the sensitivity parameter

library(DOS)
diff=df_treated$years.lived.since.1971.up.to.1992-
  df_control$years.lived.since.1971.up.to.1992
senWilcox(diff, gamma = 1, conf.int = FALSE, alpha = 0.05, alternative = "less")
```

```
## $pval
## [1] 9.539214e-11
```

```
# gamma=1 is the primary analysis assuming no unmeasured confounders
senWilcox(diff, gamma = 1.7, conf.int = FALSE, alpha = 0.05, alternative = "less")
```

```
## $pval
## [1] 0.02630124
```

```
# unmeasured bias up to gamma=1.7 cannot explain away the treatment effect
senWilcox(diff, gamma = 1.8, conf.int = FALSE, alpha = 0.05, alternative = "less")
```

```
## $pval
## [1] 0.07017079
```

```
# gamma=1.8 would explain away the treatment effect because the p-value is now larger than alpha=0.05
```

## 9: Alternative estimators: outcome regression, IPW, AIPW

```
# unadjusted. Would be biased due to confounding
reg.model=lm(years.lived.since.1971.up.to.1992~physically.inactive,data=df)
coef(reg.model)[2] # Point estimate of treatment effect
```

```
## physically.inactive
## -2.889424
```

```
confint(reg.model)[2,] # Confidence interval
```

```
## 2.5 % 97.5 %
## -3.434285 -2.344563
```

```
library(CausalGAM) # this package implements the stablized version
```

```
## Loading required package: gam
```

```
## Loading required package: splines
```

```
## Loading required package: foreach
```

```
##
```

```
## Attaching package: 'foreach'
```

```
## The following objects are masked from 'package:purrr':
```

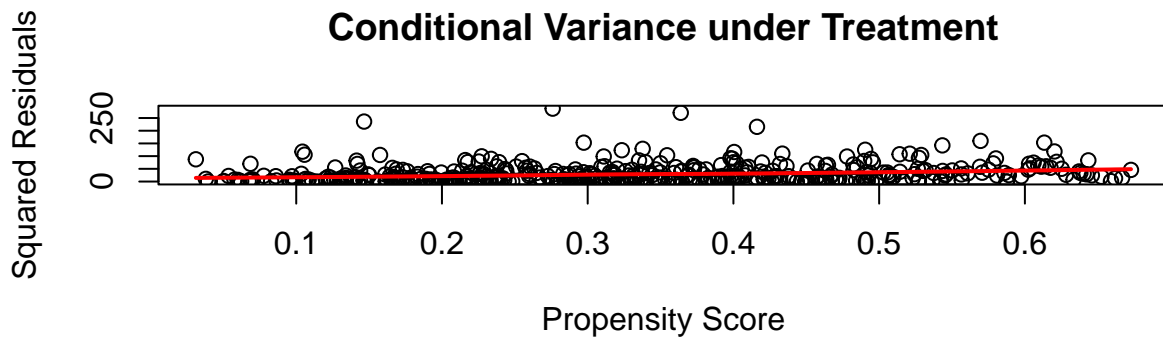
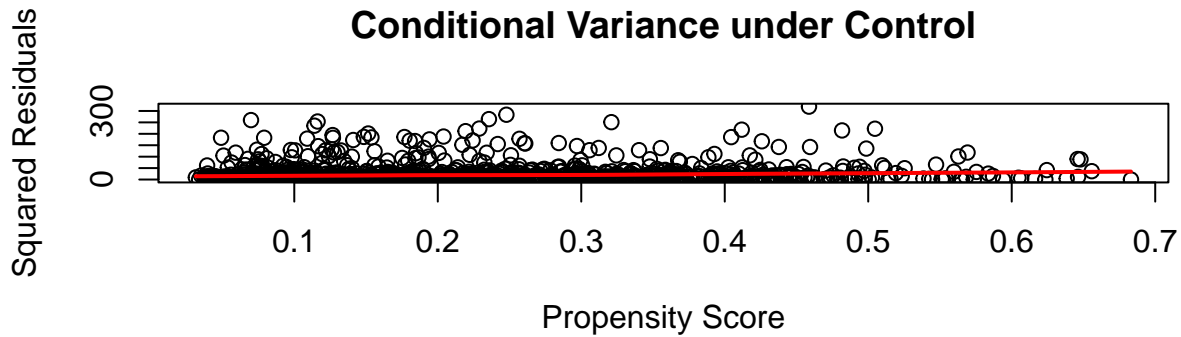
```
##
```

```
## accumulate, when
```

```
## Loaded gam 1.20.2
```

```
## ##
## ## CausalGAM Package
## ## Copyright (C) 2009 Adam Glynn and Kevin Quinn
out1.formula<-years.lived.since.1971.up.to.1992~sex+smoking.status+income.poverty.ratio+
age.at.interview+race+education+working.last.three.months+married+
alcohol.consumption+dietary.adequacy+income.poverty.ratio.missingind+
dietary.adequacy.missingind

ATE.out<-estimate.ATE(pscore.formula=ps.formula,pscore.family = binomial,
outcome.formula.t=out1.formula,outcome.formula.c=out1.formula,
outcome.family=gaussian,
treatment.var="physically.inactive",data=df,nboot=0)
```



```
print(ATE.out)

## #####
## AIPW Estimator:
## -----
## Estimated ATE:  -1.8731
##                SE           z-statistic      Pr(>|z|)
## Emp. Sandwich   0.3144           -5.9571      < 0.0001
## Estim. Asymptotic 0.2828           -6.6231      < 0.0001
## #####
## #####
## IPW Estimator:
## -----
## Estimated ATE:  -1.6107
##                SE           z-statistic      Pr(>|z|)
```

```

## Estim. Asymptotic  0.2829          -5.6946          < 0.0001
## #####
##
## #####
## Regression Estimator:
## -----
## Estimated ATE:  -1.8375
##                SE          z-statistic          Pr(>|z|)
## Estim. Asymptotic  0.2828          -6.4975          < 0.0001
## #####
##
## #####
## General Information
## -----
## Control Value:  physically.inactive = 0
## Treated Value:  physically.inactive = 1
## Number of Discarded Units:  0
## Number of Truncated Propensity Scores:  0
## Number of Treated Units Before Discards/Truncations:  466
## Number of Treated Units not Discarded/Truncated:  466
## Number of Control Units Before Discards/Truncations:  1476
## Number of Control Units not Discarded/Truncated:  1476
## #####
# validate the IPW estimator from the CausalGAM package
ps.fitted<-glm(ps.formula,family=binomial,data=df)$fitted.values
my.ipw<-mean(with(df,physically.inactive*years.lived.since.1971.up.to.1992/ps.fitted))-
mean(with(df,(1-physically.inactive)*years.lived.since.1971.up.to.1992/(1-ps.fitted))) # non-stablized
print(paste("my ipw =", round(my.ipw,4)))

## [1] "my ipw = -1.1186"

my.sipw<-sum(with(df,physically.inactive*years.lived.since.1971.up.to.1992/ps.fitted))/
sum(with(df,physically.inactive/ps.fitted))-
sum(with(df,(1-physically.inactive)*years.lived.since.1971.up.to.1992/(1-ps.fitted)))/
sum(with(df,(1-physically.inactive)/(1-ps.fitted))) # stablized IPW
print(paste("my sipw =", round(my.sipw,4)))

## [1] "my sipw = -1.6107"

#entropy balancing
library(ebal)

## ##
## ## ebal Package: Implements Entropy Balancing.
## ## See http://www.stanford.edu/~jhain/ for additional information.

A<-df$physically.inactive
X<-model.matrix(glm(ps.formula,family=binomial,data=df))
X<-X[,-1]
Y<-df$years.lived.since.1971.up.to.1992

eb.out<-ebalance(Treatment = A, X=X)

## Converged within tolerance

```

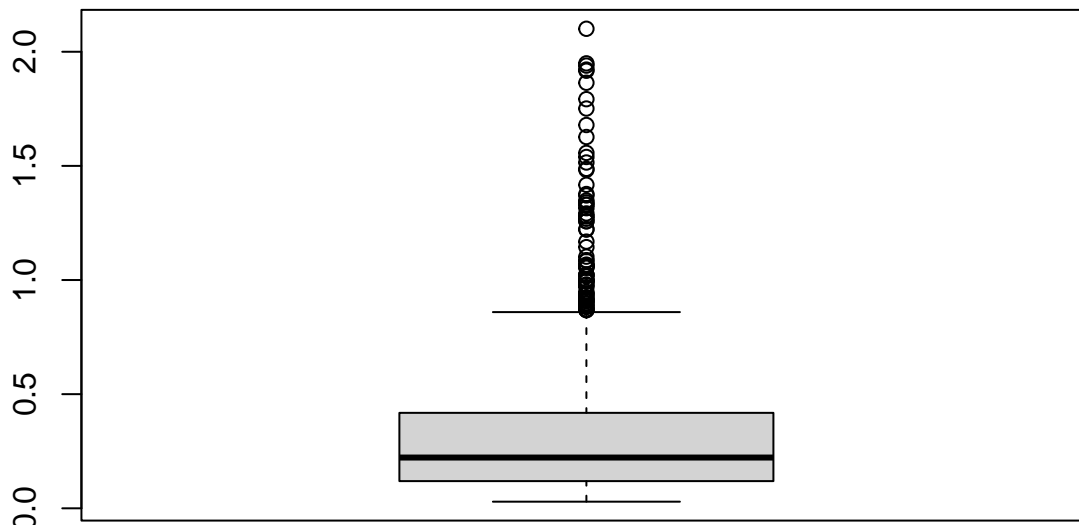
```
res<-data.frame(treated=apply(X[A==1,],2,mean),weighted.control=apply(X[A==0,],2,weighted.mean,w=eb.out$w))
res
```

	treated	weighted.control
## sexMale	0.416309013	0.416309736
## smoking.statusFormer	0.171673820	0.171674149
## smoking.statusNever	0.474248927	0.474248555
## income.poverty.ratio	2.202049669	2.202054549
## age.at.interview	61.686695279	61.686678489
## raceWhite	0.793991416	0.793991578
## education12	0.167381974	0.167382219
## education9-11	0.184549356	0.184549418
## educationCollege Grad	0.083690987	0.083691073
## educationMissing	0.006437768	0.006437763
## educationSome College	0.066523605	0.066523842
## working.last.three.monthsTRUE	0.248927039	0.248928948
## marriedTRUE	0.626609442	0.626609654
## alcohol.consumption1-4 times per month	0.173819742	0.173819909
## alcohol.consumption2+ times per week	0.090128755	0.090128826
## alcohol.consumptionJust about everyday/everyday	0.133047210	0.133047498
## alcohol.consumptionNever	0.403433476	0.403433044
## dietary.adequacy	3.277570442	3.277571376
## income.poverty.ratio.missingind	0.272532189	0.272531904
## dietary.adequacy.missingind	0.225321888	0.225321630

```
summary(eb.out$w)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 0.0291 0.1192 0.2223 0.3157 0.4181 2.1006
```

```
boxplot(eb.out$w)
```



10

```
# sensitivity analysis using bootsens
library(bootsens)
# ipw
```

```

A<-df$physically.inactive
X<-model.matrix(glm(ps.formula,family=binomial,data=df))
X<-X[,-1]
Y<-df$years.lived.since.1971.up.to.1992

## IPW, assuming no unmeasure confounder (i.e. gamma = 0 or Gamma = e^0 = 1)
extrema.os(A, X, Y) # point estimate

## [1] -1.610729 -1.610729
bootsens.os(A, X, Y, parallel = FALSE) # bootstrap confidence interval

##      2.5%      97.5%
## -2.257772 -1.023293
## IPW, Sensitivity analysis (Gamma = exp(gamma))
extrema.os(A, X, Y, gamma = log(1.2)) # point estimate

## [1] -2.5119367 -0.7512871
bootsens.os(A, X, Y, gamma = log(1.2), parallel = FALSE) # bootstrap CI

##      2.5%      97.5%
## -3.1532805 -0.2151229
# the IPW estimator is robust to Gamma=1.2

## AIPW, assuming no unmeasure confounder (i.e. gamma = 0 or Gamma = e^0 = 1)
extrema.os(A, X, Y, reg.adjust = TRUE) # point estimate

## [1] -1.872209 -1.872209
options(warn=-1) # there is a category with small number,
# which leads to some bootstrap samples without that category and generates warnings.
# this is to temporarily turn off the warning
bootsens.os(A, X, Y, reg.adjust = TRUE, parallel = FALSE) # bootstrap confidence interval

##      2.5%      97.5%
## -2.556446 -1.274457
## AIPW, Sensitivity analysis (Gamma = exp(gamma))
extrema.os(A, X, Y, reg.adjust = TRUE,gamma = log(1.2)) # point estimate

## [1] -2.555729 -1.206745
bootsens.os(A, X, Y, reg.adjust = TRUE, gamma = log(1.2),parallel = FALSE) # bootstrap CI

##      2.5%      97.5%
## -3.1896053 -0.6580529
bootsens.os(A, X, Y, reg.adjust = TRUE, gamma = log(1.4),parallel = FALSE) # bootstrap CI

##      2.5%      97.5%
## -3.8506294 -0.0941925
# the AIPW estimator is robust to Gamma=1.4, because it has higher power compared to the IPW
options(warn=0)

```