## Group Project

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# libraries: ---
library(Hmisc)
library(data.table)
library(splines)
                           _____
path = "/Users/Sabrina/Documents/2019UMICH/STATS506/group project/data/"
sleep = as.data.table(sasxport.get(paste0(path, "SLQ_D.XPT")))
physical = as.data.table(sasxport.get(paste0(path, "PAQ_D.XPT")))
physical_indv = as.data.table(sasxport.get(paste0(path, "PAQIAF_D.XPT")))
demo = as.data.table(sasxport.get(paste0(path, "DEMO D.XPT")))
dietary1 = as.data.table(sasxport.get(paste0(path, "DR1TOT_D.XPT")))
dietary2 = as.data.table(sasxport.get(paste0(path, "DR2TOT_D.XPT")))
# keep the variables we need
sleep = sleep[, .(seqn, sld010h)]
physical = physical[, .(seqn, pad080, paq520)]
physical_indv = physical_indv[, .(seqn, padtimes, paddurat)]
demo = demo[, .(seqn, riagendr, ridageyr, ridreth1, indfminc, ridexmon)]
dietary1 = dietary1[, .(seqn, dr1tkcal, dr1tsugr, dr1tcaff, day = 1)]
dietary2 = dietary2[, .(seqn, dr2tkcal, dr2tsugr, dr2tcaff, day = 2)]
# rename colnames
names(dietary1) = c("seqn", "drtkcal", "drtsugr", "drtcaff", "day")
names(dietary2) = c("seqn", "drtkcal", "drtsugr", "drtcaff", "day")
# merge day 1 data
data1 = merge(sleep, physical, all = TRUE)
data1 = merge(data1, physical_indv, all = TRUE)
data1 = merge(data1, demo, all = TRUE)
data1 = merge(data1, dietary1, all = TRUE)
# merge day 2 data
data2 = merge(sleep, physical, all = TRUE)
data2 = merge(data2, physical_indv, all = TRUE)
data2 = merge(data2, demo, all = TRUE)
data2 = merge(data2, dietary2, all = TRUE)
# row bind day 1 and day 2 data
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data = rbind(data1, data2)
# omit the rows that have missing values
data_naomit = data[complete.cases(data[])]
# take mean for repeated seqn
avg_value = data_naomit[, lapply(.SD, mean), by = .(seqn),
                   .SDcols = c("sld010h", "pad080", "paq520", "padtimes",
                       "paddurat", "ridageyr", "ridreth1", "indfminc",
                       "ridexmon", "drtkcal", "drtsugr", "drtcaff")]
# fit linear model
model = lm(sld010h ~ pad080 + padtimes + ridageyr + as.factor(ridreth1) +
            indfminc + as.factor(ridexmon) + drtkcal + drtsugr + drtcaff,
          data = avg_value)
summary(model)
##
## Call:
## lm(formula = sld010h ~ pad080 + padtimes + ridageyr + as.factor(ridreth1) +
      indfminc + as.factor(ridexmon) + drtkcal + drtsugr + drtcaff,
##
      data = avg_value)
##
## Residuals:
## How much sleep do you get (hours)?
      Min 1Q Median 3Q
## -5.8500 -0.8438 -0.0088 0.8833 5.2411
## Coefficients:
##
                        Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                      7.786e+00 1.476e-01 52.743 < 2e-16 ***
                      -1.922e-03 6.707e-04 -2.866 0.004228 **
## pad080
                       4.931e-03 2.764e-03 1.784 0.074679 .
## padtimes
## ridageyr
                      -6.926e-03 2.085e-03 -3.322 0.000919 ***
## as.factor(ridreth1)2 -3.866e-01 2.019e-01 -1.915 0.055757 .
## as.factor(ridreth1)3 2.258e-02 1.061e-01 0.213 0.831471
## as.factor(ridreth1)4 -5.699e-01 1.063e-01 -5.360 9.83e-08 ***
## as.factor(ridreth1)5 -1.190e-01 1.827e-01 -0.651 0.514866
                       1.454e-03 3.435e-03 0.423 0.672248
## as.factor(ridexmon)2 -9.247e-02 7.793e-02 -1.187 0.235635
## drtkcal
                      -7.351e-05 5.338e-05 -1.377 0.168705
                      -1.995e-04 6.589e-04 -0.303 0.762048
## drtsugr
                       -7.186e-04 2.666e-04 -2.696 0.007107 **
## drtcaff
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.319 on 1318 degrees of freedom
## Multiple R-squared: 0.05457, Adjusted R-squared: 0.04596
## F-statistic: 6.339 on 12 and 1318 DF, p-value: 5.119e-11
# do spline for "padtimes" variable
spline_model = lm(sld010h ~ pad080 + bs(padtimes) + ridageyr +
                   as.factor(ridreth1) + indfminc + as.factor(ridexmon) +
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drtkcal + drtsugr + drtcaff,
                 data = avg_value)
summary(spline model)
##
## lm(formula = sld010h ~ pad080 + bs(padtimes) + ridageyr + as.factor(ridreth1) +
      indfminc + as.factor(ridexmon) + drtkcal + drtsugr + drtcaff,
##
      data = avg_value)
##
## Residuals:
## How much sleep do you get (hours)?
               1Q Median
## -5.9177 -0.8271 -0.0118 0.8720 5.2470
##
## Coefficients:
##
                         Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                        7.680e+00 1.564e-01 49.094 < 2e-16 ***
## pad080
                       -1.998e-03 6.709e-04 -2.978 0.002953 **
## bs(padtimes)1
                       1.201e+00 4.673e-01 2.571 0.010247 *
## bs(padtimes)2
                       -1.816e+00 1.171e+00 -1.552 0.120965
## bs(padtimes)3
                       1.500e+00 8.613e-01
                                             1.742 0.081812 .
## ridageyr
                       -6.925e-03 2.092e-03 -3.310 0.000958 ***
## as.factor(ridreth1)2 -3.864e-01 2.018e-01 -1.915 0.055694 .
## as.factor(ridreth1)3 2.850e-02 1.060e-01 0.269 0.788049
## as.factor(ridreth1)4 -5.680e-01 1.066e-01 -5.331 1.15e-07 ***
## as.factor(ridreth1)5 -1.082e-01 1.826e-01 -0.592 0.553636
## indfminc
                        1.500e-03 3.432e-03 0.437 0.662095
## as.factor(ridexmon)2 -9.660e-02 7.800e-02 -1.238 0.215775
## drtkcal
                       -6.912e-05 5.337e-05 -1.295 0.195474
## drtsugr
                       -2.791e-04 6.594e-04 -0.423 0.672157
## drtcaff
                       -6.888e-04 2.667e-04 -2.582 0.009923 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.317 on 1316 degrees of freedom
## Multiple R-squared: 0.05798,
                                  Adjusted R-squared: 0.04795
## F-statistic: 5.785 on 14 and 1316 DF, p-value: 4.423e-11
# plot splines results
plot(avg_value$sld010h, avg_value$padtimes, col="grey",xlab="Sleep (hr.)",ylab="Padtimes")
fit1 = smooth.spline(avg_value$sld010h, avg_value$padtimes, df=12)
lines(fit1,col="red",lwd=2)
```

