

# BIOSTAT650\_Final\_Project

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## (1) Data cleaning

```
rm(list = ls())
gc()

##          used (Mb) gc trigger (Mb) max used (Mb)
## Ncells 469621 25.1   1011344 54.1   660860 35.3
## Vcells 878082  6.7    8388608 64.0   1800812 13.8

set.seed(123)
##### (1) Data cleaning #####
## select variables
library(NHANES)
df0 <- NHANES
df <- NHANES[NHANES$Age >= 18 & NHANES$Age < 60,]
# colSums(is.na(df)) / nrow(df)
df <- df[, which(colSums(is.na(df)) / nrow(df) < 0.3)]
# exclude duplication
df <- df[!duplicated(df),]
names(df)

## [1] "ID"          "SurveyYr"      "Gender"        "Age"
## [5] "AgeDecade"    "Race1"         "Education"     "MaritalStatus"
## [9] "HHIncome"     "HHIncomeMid"   "Poverty"       "HomeRooms"
## [13] "HomeOwn"      "Work"          "Weight"        "Height"
## [17] "BMI"          "BMI_WHO"       "Pulse"         "BPSysAve"
## [21] "BPDiaAve"     "BPSys1"        "BPDia1"        "BPSys2"
## [25] "BPDia2"       "BPSys3"        "BPDia3"        "DirectChol"
## [29] "TotChol"      "UrineVol1"     "UrineFlow1"    "Diabetes"
## [33] "HealthGen"    "DaysPhysHlthBad" "DaysMentHlthBad" "LittleInterest"
## [37] "Depressed"    "SleepHrsNight" "SleepTrouble"  "PhysActive"
## [41] "Alcohol12PlusYr" "AlcoholYear"   "Smoke100"      "Smoke100n"
## [45] "Marijuana"    "RegularMarij"  "HardDrugs"     "SexEver"
## [49] "SexAge"       "SexNumPartnLife" "SexNumPartYear" "SameSex"
## [53] "SexOrientation"

# df$BPSysAve
library(dplyr)

##
## Attaching package: 'dplyr'

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

```
##
##      filter, lag
## The following objects are masked from 'package:base':
##
##      intersect, setdiff, setequal, union
```

```
df2 <- df %>% select(
  SleepHrsNight,
  BMI,
  DirectChol,
  Age,
  Gender,
  Race1,
  TotChol,
  BPDiaAve,
  BPSysAve,
  AlcoholYear,
  Poverty,
  SexNumPartnLife,
  SexNumPartYear,
  DaysMentHlthBad,
  UrineFlow1,
  PhysActive,
  DaysPhysHlthBad,
  Smoke100,
  Depressed,
  HealthGen,
  SexAge
)

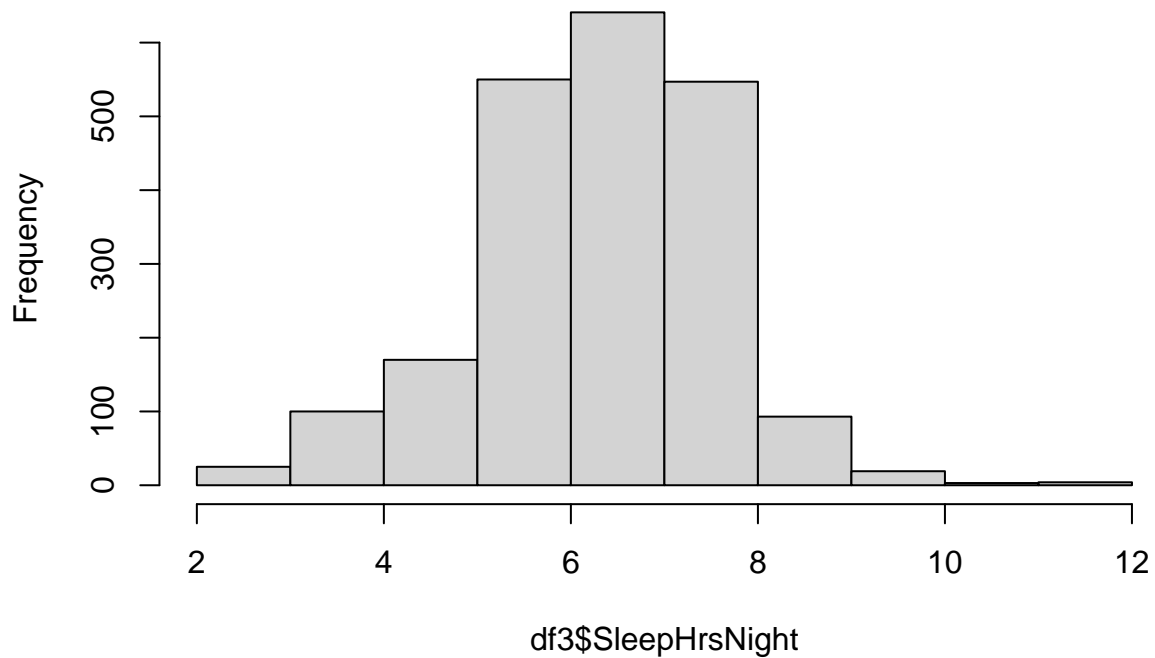
df3 <- na.omit(df2)
#df3$SleepHrsNight <- df3$SleepHrsNight * 60
#df3 <- df3[, -which(names(df3) %in% "SleepHrsNight")]
# cor(df3$BPSysAve, df3$BPDiaAve)
psych::describe(df3)
```

##	vars	n	mean	sd	median	trimmed	mad	min	max
## SleepHrsNight	1	2152	6.78	1.31	7.00	6.85	1.48	2.00	12.00
## BMI	2	2152	28.77	6.75	27.60	28.09	5.78	15.02	69.00
## DirectChol	3	2152	1.35	0.41	1.29	1.31	0.39	0.39	3.83
## Age	4	2152	39.18	11.33	39.00	39.15	14.83	20.00	59.00
## Gender*	5	2152	1.53	0.50	2.00	1.54	0.00	1.00	2.00
## Race1*	6	2152	3.43	1.15	4.00	3.57	0.00	1.00	5.00
## TotChol	7	2152	5.07	1.05	4.99	5.01	1.04	1.53	13.65
## BPDiaAve	8	2152	71.19	11.84	71.00	71.28	10.38	0.00	116.00
## BPSysAve	9	2152	117.43	14.28	116.00	116.50	13.34	78.00	209.00
## AlcoholYear	10	2152	70.59	94.22	24.00	50.94	35.58	0.00	364.00
## Poverty	11	2152	2.84	1.69	2.78	2.89	2.49	0.00	5.00
## SexNumPartnLife	12	2152	16.73	66.13	7.00	8.91	5.93	0.00	2000.00
## SexNumPartYear	13	2152	1.38	2.59	1.00	1.04	0.00	0.00	69.00
## DaysMentHlthBad	14	2152	4.47	8.02	0.00	2.40	0.00	0.00	30.00
## UrineFlow1	15	2152	1.07	0.97	0.81	0.91	0.60	0.00	10.14
## PhysActive*	16	2152	1.58	0.49	2.00	1.60	0.00	1.00	2.00
## DaysPhysHlthBad	17	2152	3.16	7.19	0.00	1.12	0.00	0.00	30.00

```
## Smoke100*      18 2152   1.46  0.50   1.00   1.45  0.00  1.00   2.00
## Depressed*     19 2152   1.30  0.58   1.00   1.16  0.00  1.00   3.00
## HealthGen*     20 2152   2.64  0.94   3.00   2.65  1.48  1.00   5.00
## SexAge         21 2152  17.10  3.39  17.00  16.80  2.97  9.00  44.00
##               range skew kurtosis se
## SleepHrsNight 10.00 -0.30   0.69 0.03
## BMI           53.98  1.28   2.96 0.15
## DirectChol     3.44  1.09   2.27 0.01
## Age           39.00  0.02  -1.15 0.24
## Gender*        1.00 -0.12  -1.99 0.01
## Race1*         4.00 -1.13   0.08 0.02
## TotChol        12.12  0.92   3.47 0.02
## BPDiaAve       116.00 -0.39   3.13 0.26
## BPSysAve       131.00  1.00   2.94 0.31
## AlcoholYear    364.00  1.66   1.98 2.03
## Poverty         5.00 -0.01  -1.47 0.04
## SexNumPartnLife 2000.00 18.82  456.62 1.43
## SexNumPartYear  69.00 14.07  293.16 0.06
## DaysMentHlthBad 30.00  2.16   3.76 0.17
## UrineFlow1     10.14  2.89  14.06 0.02
## PhysActive*     1.00 -0.32  -1.90 0.01
## DaysPhysHlthBad 30.00  2.80   7.06 0.15
## Smoke100*       1.00  0.15  -1.98 0.01
## Depressed*       2.00  1.83   2.21 0.01
## HealthGen*       4.00  0.11  -0.33 0.02
## SexAge          35.00  1.51   5.56 0.07
```

```
# psych::pairs.panels(df3)
hist(df3$SleepHrsNight)
```

## Histogram of df3\$SleepHrsNight



```
# colSums(is.na(df2)) / nrow(df2)
fit0 <-
  lm(SleepHrsNight ~ .,
      data = df3)
#data type
df3$Gender <- ifelse(df3$Gender == "male", 0, 1)
df3 <- df3 %>%
  mutate(
    Race1 = case_when(
      Race1 == 'Black' ~ 1,
      Race1 == 'Hispanic' ~ 2,
      Race1 == 'Mexican' ~ 3,
      Race1 == 'White' ~ 4,
      Race1 == 'Other' ~ 5,
      TRUE ~ NA_integer_ # Default value if none of the conditions are met
    )
  )
```

## (2) Baseline characteristics

```
Hmisc::describe(df3)
```

```
## df3
##
```

```

## 21 Variables      2152 Observations
## -----
## SleepHrsNight
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      11      0.94      6.781      1.415      4      5
##      .25      .50      .75      .90      .95
##      6      7      8      8      9
##
## lowest : 2 3 4 5 6, highest: 8 9 10 11 12
##
## Value      2      3      4      5      6      7      8      9      10      11      12
## Frequency      3      22      100      170      550      641      547      93      19      3      4
## Proportion 0.001 0.010 0.046 0.079 0.256 0.298 0.254 0.043 0.009 0.001 0.002
## -----
## BMI
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      1072      1      28.77      7.223      20.18      21.50
##      .25      .50      .75      .90      .95
##    24.00      27.60      32.00      37.36      41.22
##
## lowest : 15.02 15.80 15.98 16.51 16.70, highest: 62.80 63.30 63.91 67.83 69.00
## -----
## DirectChol
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      98      0.999      1.346      0.4446      0.80      0.91
##      .25      .50      .75      .90      .95
##      1.06      1.29      1.58      1.89      2.09
##
## lowest : 0.39 0.41 0.52 0.54 0.57, highest: 3.13 3.41 3.44 3.59 3.83
## -----
## Age
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      40      0.999      39.18      13.08      21      23
##      .25      .50      .75      .90      .95
##      30      39      49      55      57
##
## lowest : 20 21 22 23 24, highest: 55 56 57 58 59
## -----
## Gender
##      n missing distinct      Info      Sum      Mean      Gmd
##    2152      0      2      0.747      1011      0.4698      0.4984
##
## -----
## Race1
##      n missing distinct      Info      Mean      Gmd
##    2152      0      5      0.758      3.428      1.115
##
## lowest : 1 2 3 4 5, highest: 1 2 3 4 5
##
## Value      1      2      3      4      5
## Frequency      289      145      230      1333      155
## Proportion 0.134 0.067 0.107 0.619 0.072
## -----
## TotChol

```

```

##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0      208         1      5.069      1.151      3.57      3.85
##          .25      .50      .75      .90      .95
##      4.32      4.99      5.69      6.36      6.83
##
## lowest :  1.53  2.69  2.74  2.79  2.82, highest:  9.31  9.34  9.90 12.28 13.65
## -----
## BPDiaAve
##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0       84      0.999      71.19      12.83       53       57
##          .25      .50      .75      .90      .95
##          64       71       78       85       89
##
## lowest :    0  20  21  22  25, highest: 108 109 110 114 116
## -----
## BPSysAve
##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0       98      0.999      117.4      15.44       97      101
##          .25      .50      .75      .90      .95
##          108      116      125      134      142
##
## lowest :   78  83  84  85  86, highest: 182 184 191 202 209
## -----
## AlcoholYear
##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0       56      0.993      70.59      91.9         0         0
##          .25      .50      .75      .90      .95
##          4       24      104      208      260
##
## lowest :    0   1   2   3   4, highest: 260 300 312 360 364
## -----
## Poverty
##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0      393      0.988      2.841      1.931      0.340      0.660
##          .25      .50      .75      .90      .95
##      1.277      2.780      4.817      5.000      5.000
##
## lowest : 0.00 0.02 0.03 0.04 0.05, highest: 4.95 4.96 4.97 4.99 5.00
## -----
## SexNumPartnLife
##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0       81      0.995      16.73      22.47         1         1
##          .25      .50      .75      .90      .95
##          3         7       15       30       50
##
## lowest :    0   1   2   3   4, highest:  600  800  999 1000 2000
## -----
## SexNumPartYear
##          n missing distinct      Info      Mean      Gmd      .05      .10
##      2152         0       21      0.645      1.381      1.18         0         0
##          .25      .50      .75      .90      .95
##          1         1         1         2         3
##
## lowest :    0   1   2   3   4, highest:  19  20  30  50  69

```

```

## -----
## DaysMentHlthBad
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      28    0.844    4.475    6.894      0      0
##      .25      .50      .75      .90      .95
##      0      0      5      15      30
##
## lowest :  0  1  2  3  4, highest: 25 26 27 29 30
## -----
## UrineFlow1
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0    1337      1    1.074    0.9061    0.1960    0.2775
##      .25      .50      .75      .90      .95
##    0.4580    0.8100    1.3618    2.1929    2.7780
##
## lowest :  0.000  0.006  0.011  0.014  0.016, highest:  7.325  7.826  8.730  9.410 10.143
## -----
## PhysActive
##      n missing distinct
##    2152      0      2
##
## Value      No  Yes
## Frequency   906 1246
## Proportion 0.421 0.579
## -----
## DaysPhysHlthBad
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      24    0.708    3.165    5.318    0.00    0.00
##      .25      .50      .75      .90      .95
##      0.00    0.00    2.00    10.00    24.45
##
## lowest :  0  1  2  3  4, highest: 24 25 26 28 30
## -----
## Smoke100
##      n missing distinct
##    2152      0      2
##
## Value      No  Yes
## Frequency  1155  997
## Proportion 0.537 0.463
## -----
## Depressed
##      n missing distinct
##    2152      0      3
##
## Value      None Several      Most
## Frequency   1657    355    140
## Proportion  0.770  0.165  0.065
## -----
## HealthGen
##      n missing distinct
##    2152      0      5
##
## lowest : Excellent Vgood      Good      Fair      Poor

```

```
## highest: Excellent Vgood      Good      Fair      Poor
##
## Value      Excellent      Vgood      Good      Fair      Poor
## Frequency      240      697      854      313      48
## Proportion      0.112      0.324      0.397      0.145      0.022
## -----
## SexAge
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    2152      0      28      0.985      17.1      3.463      13.00      14.00
##      .25      .50      .75      .90      .95
##    15.00      17.00      18.00      21.00      23.45
##
## lowest :  9 10 11 12 13, highest: 32 34 35 37 44
## -----
```

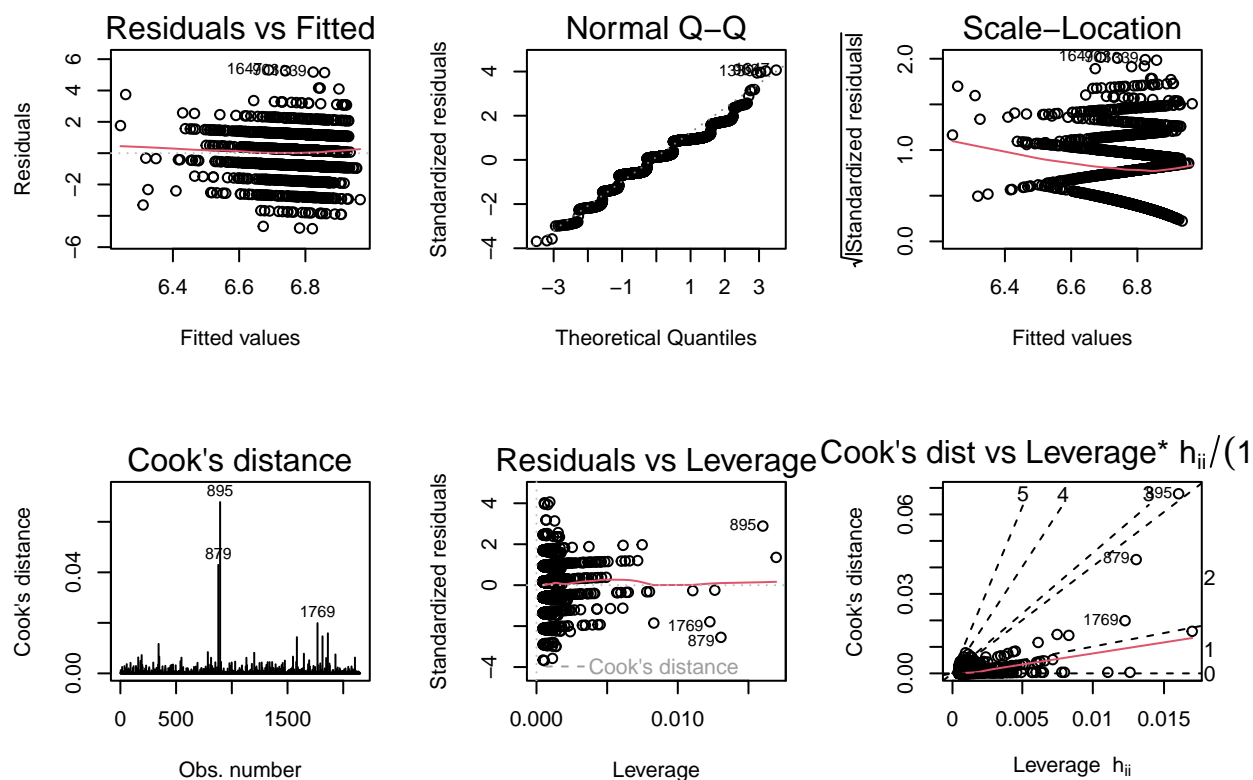
### (3) linear regression model

```
##simple linear regression##
model1 = lm(df3$SleepHrsNight ~ df3$BMI, data = df3)
summary(model1)

##
## Call:
## lm(formula = df3$SleepHrsNight ~ df3$BMI, data = df3)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.8209 -0.8022  0.1710  1.1494  5.3105
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  7.166900   0.123331  58.111 < 2e-16 ***
## df3$BMI      -0.013409   0.004174  -3.213  0.00133 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.307 on 2150 degrees of freedom
## Multiple R-squared:  0.004778, Adjusted R-squared:  0.004315
## F-statistic: 10.32 on 1 and 2150 DF, p-value: 0.001334

par(mfrow = c(2, 3)) #read more from ?plot.lm
plot(model1, which = 1)
plot(model1, which = 2)
plot(model1, which = 3)
plot(model1, which = 4)
plot(model1, which = 5)
plot(model1, which = 6)
```





```
par(mfrow = c(1, 1)) # reset

dummy_b = 1 * (df3$Race1 == "Black")
dummy_h = 1 * (df3$Race1 == "Hispanic")
dummy_m = 1 * (df3$Race1 == "Mexican")
dummy_w = 1 * (df3$Race1 == "White")
dummy_o = 1 * (df3$Race1 == "Other")

age_quant = quantile(df3$Age)
df3$AgeC = 0
df3$AgeC[df3$Age > age_quant[2] & df3$Age <= age_quant[3]] = 1
df3$AgeC[df3$Age > age_quant[3] & df3$Age <= age_quant[4]] = 2
df3$AgeC[df3$Age > age_quant[4]] = 3

### multiple linear regression###
# model_1 add demographic
m_1 = lm(BMI ~ SleepHrsNight + Age + Gender + factor(Race1), df3)
summary(m_1)
```

```
##
## Call:
## lm(formula = BMI ~ SleepHrsNight + Age + Gender + factor(Race1),
##     data = df3)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
```

```

## -14.347 -4.497 -1.201 3.190 40.277
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  30.78080   0.97780  31.480 < 2e-16 ***
## SleepHrsNight -0.29383   0.11031  -2.664 0.007785 **
## Age          0.05055   0.01282   3.944 8.26e-05 ***
## Gender       0.25869   0.28895   0.895 0.370740
## factor(Race1)2 -2.28054   0.67704  -3.368 0.000769 ***
## factor(Race1)3 -1.02309   0.59140  -1.730 0.083782 .
## factor(Race1)4 -2.51942   0.43385  -5.807 7.30e-09 ***
## factor(Race1)5 -4.14341   0.66274  -6.252 4.88e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.643 on 2144 degrees of freedom
## Multiple R-squared:  0.03564, Adjusted R-squared:  0.03249
## F-statistic: 11.32 on 7 and 2144 DF, p-value: 3.698e-14
## model_2 add known risk factors
m_2 = lm(
  BMI ~ SleepHrsNight + Age + Gender + Race1 + TotChol + BPDiaAve + BPSysAve + AlcoholYear + Smoke100 +
  DaysPhysHlthBad + PhysActive,
  df3
)
summary(m_2)

##
## Call:
## lm(formula = BMI ~ SleepHrsNight + Age + Gender + Race1 + TotChol +
##      BPDiaAve + BPSysAve + AlcoholYear + Smoke100 + DaysPhysHlthBad +
##      PhysActive, data = df3)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -14.752  -4.236  -0.849   3.055  37.857
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  21.023150   1.610401  13.055 < 2e-16 ***
## SleepHrsNight -0.212193   0.107400  -1.976 0.048314 *
## Age          0.012839   0.013495   0.951 0.341528
## Gender       0.514621   0.291331   1.766 0.077463 .
## Race1       -0.622971   0.122615  -5.081 4.09e-07 ***
## TotChol      0.076572   0.139325   0.550 0.582658
## BPDiaAve     0.054500   0.014049   3.879 0.000108 ***
## BPSysAve     0.066004   0.012027   5.488 4.55e-08 ***
## AlcoholYear  -0.009762   0.001533  -6.368 2.34e-10 ***
## Smoke100Yes  -0.507830   0.287921  -1.764 0.077911 .
## DaysPhysHlthBad 0.066309   0.019785   3.352 0.000818 ***
## PhysActiveYes -1.260928   0.292769  -4.307 1.73e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.413 on 2140 degrees of freedom

```

```
## Multiple R-squared:  0.1029, Adjusted R-squared:  0.09826
## F-statistic: 22.31 on 11 and 2140 DF,  p-value: < 2.2e-16
```

```
#LINE
```

```
#influential observations
```

```
#multicollinearity
```

```
## model_3 add additional risk factors
```

```
m_3 = lm(
  BMI ~ SleepHrsNight + Age + Gender + Race1 + Poverty + TotChol + BPDiaAve + BPSysAve + AlcoholYear +
  DaysPhysHlthBad + HealthGen + PhysActive,
  df3
)
summary(m_3)
```

```
##
```

```
## Call:
```

```
## lm(formula = BMI ~ SleepHrsNight + Age + Gender + Race1 + Poverty +
##     TotChol + BPDiaAve + BPSysAve + AlcoholYear + Smoke100 +
##     UrineFlow1 + DaysMentHlthBad + DaysPhysHlthBad + HealthGen +
##     PhysActive, data = df3)
```

```
##
```

```
## Residuals:
```

```
##      Min       1Q   Median       3Q      Max
## -16.838  -4.054  -0.646   3.203  35.902
```

```
##
```

```
## Coefficients:
```

```
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   18.47102    1.621565  11.391 < 2e-16 ***
## SleepHrsNight  -0.121393    0.106352  -1.141  0.25382
## Age            0.010806    0.013725   0.787  0.43118
## Gender         0.532917    0.286537   1.860  0.06304 .
## Race1         -0.500763    0.122151  -4.100 4.29e-05 ***
## Poverty        0.073370    0.090958   0.807  0.41997
## TotChol        0.030653    0.136000   0.225  0.82170
## BPDiaAve       0.058458    0.013721   4.260 2.13e-05 ***
## BPSysAve       0.053724    0.011806   4.550 5.65e-06 ***
## AlcoholYear   -0.008337    0.001515  -5.503 4.18e-08 ***
## Smoke100Yes   -0.807332    0.287264  -2.810  0.00499 **
## UrineFlow1    -0.113369    0.142545  -0.795  0.42652
## DaysMentHlthBad -0.030360    0.017984  -1.688  0.09153 .
## DaysPhysHlthBad 0.014779    0.020974   0.705  0.48112
```

```
## HealthGenVgood    1.922013    0.470923    4.081 4.64e-05 ***
## HealthGenGood     3.569501    0.468730    7.615 3.93e-14 ***
## HealthGenFair     5.283476    0.575334    9.183 < 2e-16 ***
## HealthGenPoor     7.546146    1.078147    6.999 3.43e-12 ***
## PhysActiveYes    -0.818408    0.294015   -2.784 0.00542 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.251 on 2133 degrees of freedom
## Multiple R-squared:  0.1504, Adjusted R-squared:  0.1432
## F-statistic: 20.97 on 18 and 2133 DF, p-value: < 2.2e-16
```

```
# model_4 add additional risk factors
```

```
m_full = lm(
  BMI ~ SleepHrsNight + Age + Gender + Race1 + Poverty + TotChol + BPDiaAve + BPSysAve + AlcoholYear +
    DaysPhysHlthBad + HealthGen + PhysActive + SleepHrsNight * Age + SleepHrsNight *
    Gender + SleepHrsNight * factor(Race1),
  df3
)
summary(m_full)
```

```
##
## Call:
## lm(formula = BMI ~ SleepHrsNight + Age + Gender + Race1 + Poverty +
##     TotChol + BPDiaAve + BPSysAve + AlcoholYear + Smoke100 +
##     UrineFlow1 + DaysMentHlthBad + DaysPhysHlthBad + HealthGen +
##     PhysActive + SleepHrsNight * Age + SleepHrsNight * Gender +
##     SleepHrsNight * factor(Race1), data = df3)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -16.958  -4.088  -0.576   3.191  36.357
##
## Coefficients: (1 not defined because of singularities)
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    22.175411    3.663849   6.052 1.68e-09 ***
## SleepHrsNight   -0.672481    0.440017  -1.528  0.12658
## Age            -0.080205    0.063471  -1.264  0.20649
## Gender          3.956938    1.441705   2.745  0.00611 **
## Race1           0.266774    0.812363   0.328  0.74265
## Poverty         0.054070    0.091689   0.590  0.55544
## TotChol         0.012933    0.135840   0.095  0.92416
## BPDiaAve        0.057750    0.013676   4.223 2.52e-05 ***
## BPSysAve        0.052227    0.011793   4.429 9.96e-06 ***
## AlcoholYear    -0.009047    0.001517  -5.966 2.84e-09 ***
## Smoke100Yes    -0.847770    0.287236  -2.951  0.00320 **
## UrineFlow1     -0.088739    0.142102  -0.624  0.53238
## DaysMentHlthBad -0.032621    0.017991  -1.813  0.06993 .
## DaysPhysHlthBad  0.014998    0.020905   0.717  0.47319
## HealthGenVgood   1.882401    0.469175   4.012 6.23e-05 ***
## HealthGenGood    3.613081    0.467141   7.734 1.59e-14 ***
## HealthGenFair    5.346537    0.574641   9.304 < 2e-16 ***
## HealthGenPoor    7.518320    1.075750   6.989 3.69e-12 ***
## PhysActiveYes   -0.891431    0.294530  -3.027  0.00250 **
## factor(Race1)2   -1.143338    2.946825  -0.388  0.69806
```

```
## factor(Race1)3          -3.941114    2.751175   -1.433    0.15214
## factor(Race1)4          -5.899855    2.349469   -2.511    0.01211 *
## factor(Race1)5           NA           NA        NA        NA
## SleepHrsNight:Age       0.012971    0.009134    1.420    0.15574
## SleepHrsNight:Gender   -0.508514    0.207897   -2.446    0.01453 *
## SleepHrsNight:factor(Race1)2 -0.160437    0.452870   -0.354    0.72317
## SleepHrsNight:factor(Race1)3  0.334059    0.403929    0.827    0.40832
## SleepHrsNight:factor(Race1)4  0.544607    0.287394    1.895    0.05823 .
## SleepHrsNight:factor(Race1)5 -0.629379    0.475731   -1.323    0.18599
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.217 on 2124 degrees of freedom
## Multiple R-squared:  0.1631, Adjusted R-squared:  0.1525
## F-statistic: 15.33 on 27 and 2124 DF,  p-value: < 2.2e-16
```

## (4) Diagnosis: 10-fold CV

```
library(caret)
```

```
## Loading required package: ggplot2
```

```
## Loading required package: lattice
```

```
splitIndex <-
  createDataPartition(df3$SleepHrsNight, p = 0.7, list = FALSE)
trainData <- df3[splitIndex, ]
testData <- df3[-splitIndex, ]
predictions <- predict(m_3, newdata = testData)
mse <- mean((testData$SleepHrsNight - predictions) ^ 2)
control <-
  trainControl(method = "cv", number = 10) # 10-fold cross-validation
cv_model <-
  train(
    SleepHrsNight ~ .,
    data = df3,
    method = "lm",
    trControl = control
  )
cv_model
```

```
## Linear Regression
```

```
##
```

```
## 2152 samples
```

```
## 21 predictor
```

```
##
```

```
## No pre-processing
```

```
## Resampling: Cross-Validated (10 fold)
```

```
## Summary of sample sizes: 1937, 1938, 1936, 1937, 1937, 1937, ...
```

```
## Resampling results:
```

```
##
```

```
## RMSE      Rsquared    MAE
```

```
## 1.280209  0.05043061  0.9931499
```

```
##
## Tuning parameter 'intercept' was held constant at a value of TRUE
(cv_results <- cv_model$results)

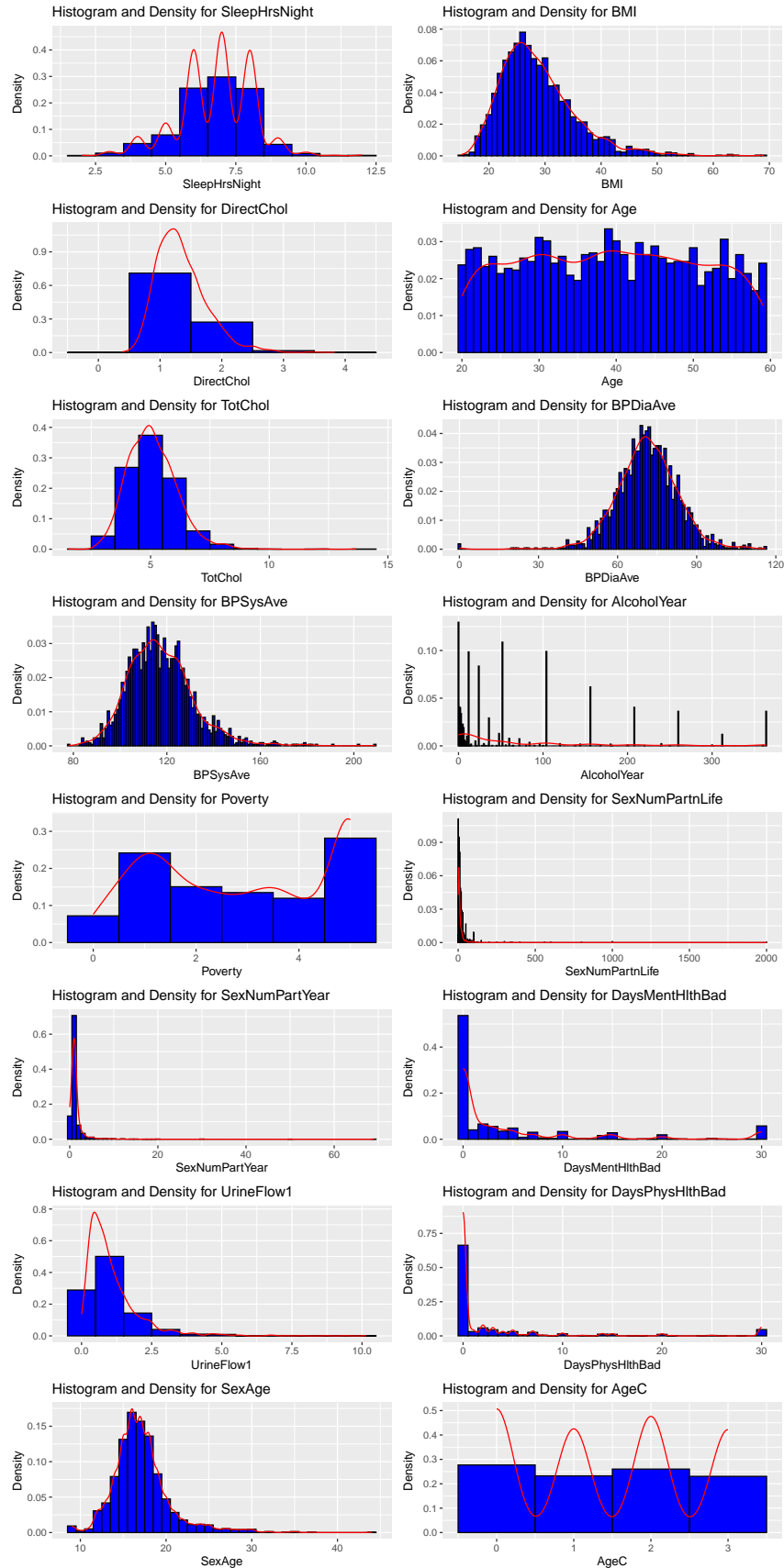
##      intercept      RMSE    Rsquared      MAE      RMSESD RsquaredSD      MAESD
## 1          TRUE 1.280209 0.05043061 0.9931499 0.04543809 0.02732622 0.02794626
```

## (4) Diagnosis: Normality Assumption

```
library(ggplot2)
library(patchwork)
# Initializes an empty patchwork object
plot_list <- list()

# Draw a histogram for each numeric variable (except Race1 and Gender) and add it to the list
for (var in names(df3)) {
  if (is.numeric(df3[[var]]) && !(var %in% c("Race1", "Gender"))) {
    p <- ggplot(df3, aes(x = .data[[var]])) +
      geom_histogram(
        aes(y = after_stat(density)),
        binwidth = 1,
        fill = "blue",
        color = "black"
      ) +
      geom_density(col = "red") +
      ggtitle(paste("Histogram and Density for", var)) +
      xlab(var) +
      ylab("Density")
    plot_list[[length(plot_list) + 1]] <- p
  }
}

# Use patchwork to put all the charts together
combined_plot <- wrap_plots(plot_list, ncol = 2)
print(combined_plot)
```



```

df3 <- data.frame(df3)
library(dplyr)
# Shapiro-Wilk normality test is performed for each numerical variable in df3
results <- sapply(df3, function(x) {
  if (is.numeric(x)) {
    shapiro_test <- shapiro.test(x)
    return(c(shapiro_test$statistic, shapiro_test$p.value))
  } else {
    return(c(NA, NA))
  }
})
# Convert the result to a data box and name the column
results_df <- as.data.frame(t(results))
names(results_df) <- c("W", "p.value")
# Add a variable name as a new column
results_df$Variable <- rownames(results_df)
# Rearrange the order of columns
results_df <- results_df[, c("Variable", "W", "p.value")]
# Calculate the corrected P-value (for example, using Bonferroni correction)
results_df$p.adjusted <-
  p.adjust(results_df$p.value, method = "bonferroni")
print(results_df)

```

##	Variable	W	p.value	p.adjusted
## SleepHrsNight	SleepHrsNight	0.9347691	1.022342e-29	1.840215e-28
## BMI	BMI	0.9263898	2.950926e-31	5.311666e-30
## DirectChol	DirectChol	0.9439221	7.552977e-28	1.359536e-26
## Age	Age	0.9579654	1.832383e-24	3.298290e-23
## Gender	Gender	0.6352876	1.636740e-55	2.946133e-54
## Race1	Race1	0.7327797	3.104346e-50	5.587823e-49
## TotChol	TotChol	0.9642744	1.175111e-22	2.115200e-21
## BPDiaAve	BPDiaAve	0.9718079	3.709893e-20	6.677808e-19
## BPSysAve	BPSysAve	0.9554033	3.865527e-25	6.957949e-24
## AlcoholYear	AlcoholYear	0.7454040	1.944127e-49	3.499428e-48
## Poverty	Poverty	0.8942742	4.092136e-36	7.365845e-35
## SexNumPartnLife	SexNumPartnLife	0.1496531	2.951432e-71	5.312577e-70
## SexNumPartYear	SexNumPartYear	0.2562318	1.244353e-68	2.239836e-67
## DaysMentHlthBad	DaysMentHlthBad	0.6112779	1.254550e-56	2.258190e-55
## UrineFlow1	UrineFlow1	0.7555438	8.969094e-49	1.614437e-47
## PhysActive	PhysActive	NA	NA	NA
## DaysPhysHlthBad	DaysPhysHlthBad	0.4968273	2.926552e-61	5.267794e-60
## Smoke100	Smoke100	NA	NA	NA
## Depressed	Depressed	NA	NA	NA
## HealthGen	HealthGen	NA	NA	NA
## SexAge	SexAge	0.8954434	5.842918e-36	1.051725e-34
## AgeC	AgeC	0.8533480	8.034125e-41	1.446143e-39

## Standardized residuals, Studentized residuals

```

# Regular residuals
residual_1 <- m_full$residuals

```



```

# Standardized residuals
residual_2 <- rstandard(m_full)

# Studentized residuals
residual_3 <- rstudent(m_full)

# Externally studentized residuals
# Note: Externally studentized residuals are the same as studentized residuals in most cases
residual_4 <- rstudent(m_full)

# Creating a data frame to summarize these residuals
residual_summary <- data.frame(
  Residuals = c("Regular", "Standardized", "Studentized", "Externally Studentized"),
  Mean = c(mean(residual_1), mean(residual_2), mean(residual_3), mean(residual_4)),
  SD = c(sd(residual_1), sd(residual_2), sd(residual_3), sd(residual_4)),
  Min = c(min(residual_1), min(residual_2), min(residual_3), min(residual_4)),
  Max = c(max(residual_1), max(residual_2), max(residual_3), max(residual_4))
)

# Display the summary
print(residual_summary)

```

	Residuals	Mean	SD	Min	Max
## 1	Regular	1.457066e-16	6.178303	-16.958137	36.356759
## 2	Standardized	2.759984e-05	1.001216	-2.750704	5.883818
## 3	Studentized	2.756080e-04	1.002353	-2.754968	5.930967
## 4	Externally Studentized	2.756080e-04	1.002353	-2.754968	5.930967

```

# Load necessary library
library(ggplot2)

# Assuming m_full is your linear model
# m_full <- lm(SleepMinNight ~ ., data = df3)

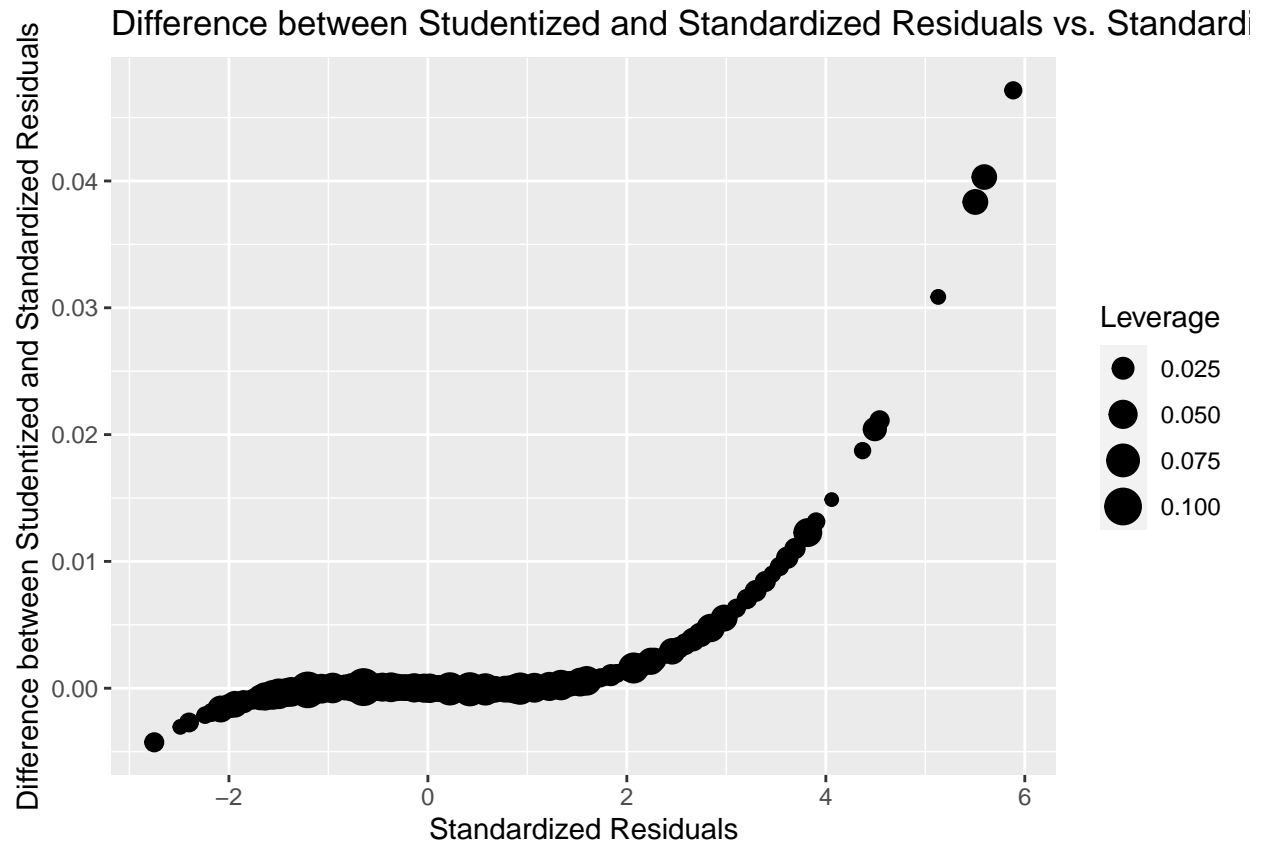
# Calculate standardized and studentized residuals
residual_2 <- rstandard(m_full)
residual_3 <- rstudent(m_full)

# Calculate leverage values
leverage_values <- hatvalues(m_full)

# Create a data frame for plotting
plot_data <- data.frame(
  Standardized_Residuals = residual_2,
  Difference = residual_3 - residual_2,
  Leverage = leverage_values
)

# Create the plot
ggplot(plot_data, aes(x = Standardized_Residuals, y = Difference)) +
  geom_point(aes(size = Leverage)) +
  ggtitle("Difference between Studentized and Standardized Residuals vs. Standardized Residuals") +
  xlab("Standardized Residuals") +
  ylab("Difference between Studentized and Standardized Residuals")

```



```
# Display the plot
print(ggplot)
```

```
## function (data = NULL, mapping = aes(), ..., environment = parent.frame())
## {
##   UseMethod("ggplot")
## }
## <bytecode: 0x4c54e60>
## <environment: namespace:ggplot2>
```

```
# Load necessary library
library(ggplot2)
```

```
# Assuming m_full is your linear model
# m_full <- lm(SleepMinNight ~ ., data = df3)
```

```
# Calculate studentized and externally studentized residuals
```

```
residual_3 <- rstudent(m_full)
```

```
residual_4 <- rstudent(m_full) # Externally studentized residuals are typically the same as studentized
```

```
# Regular residuals
```

```
residual_1 <- m_full$residuals
```

```
# Create a data frame for plotting
```

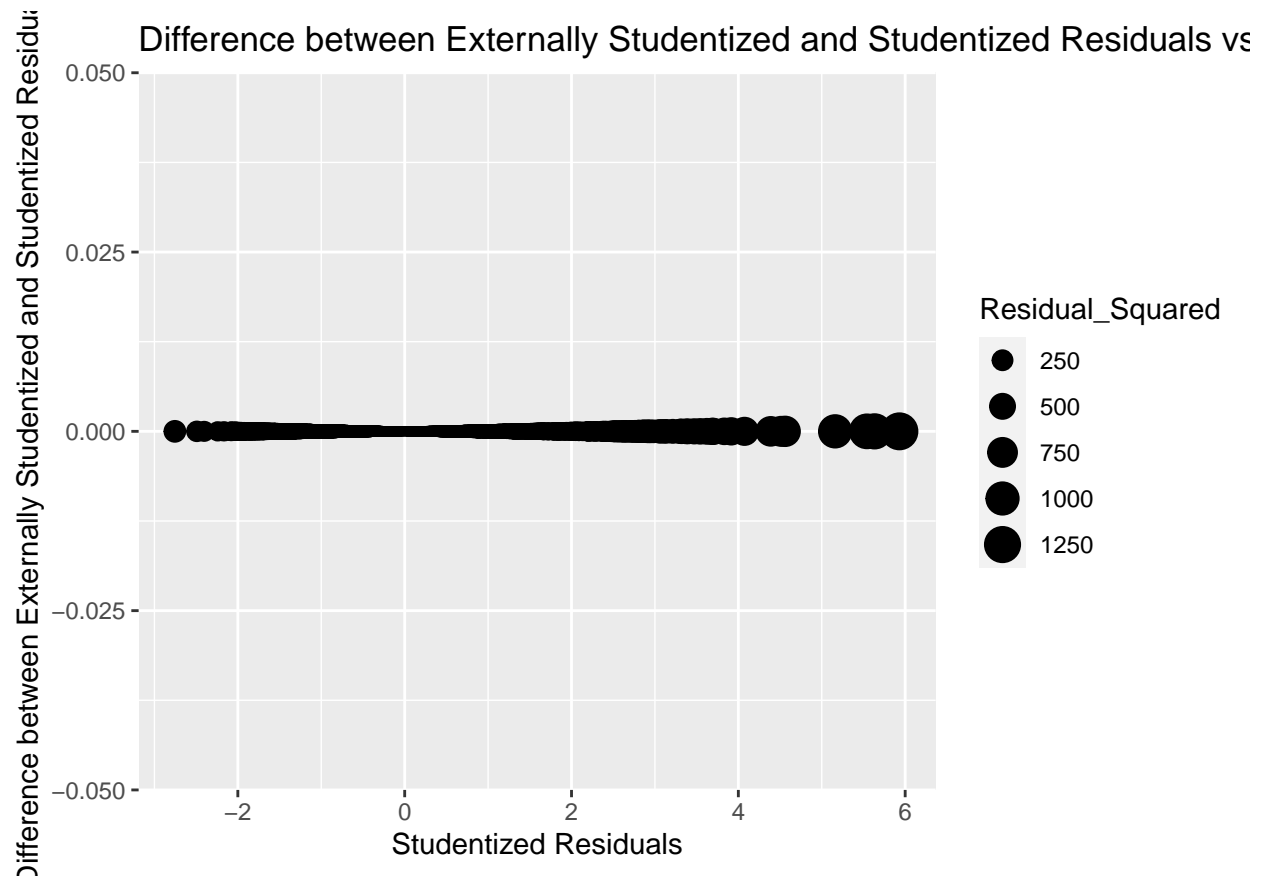
```
plot_data <- data.frame(
  Studentized_Residuals = residual_3,
  Difference = residual_4 - residual_3,
```

```

Residual_Squared = residual_1^2
)

# Create the plot
ggplot(plot_data, aes(x = Studentized_Residuals, y = Difference)) +
  geom_point(aes(size = Residual_Squared)) +
  ggtitle("Difference between Externally Studentized and Studentized Residuals vs. Studentized Residuals") +
  xlab("Studentized Residuals") +
  ylab("Difference between Externally Studentized and Studentized Residuals")

```



```

# Display the plot
print(ggplot)

```

```

## function (data = NULL, mapping = aes(), ..., environment = parent.frame())
## {
##   UseMethod("ggplot")
## }
## <bytecode: 0x4c54e60>
## <environment: namespace:ggplot2>

```

```

# Load necessary library
library(ggplot2)

# Assuming m_full is your linear model
# m_full <- lm(SleepMinNight ~ ., data = df3)

# Calculate regular residuals

```

```

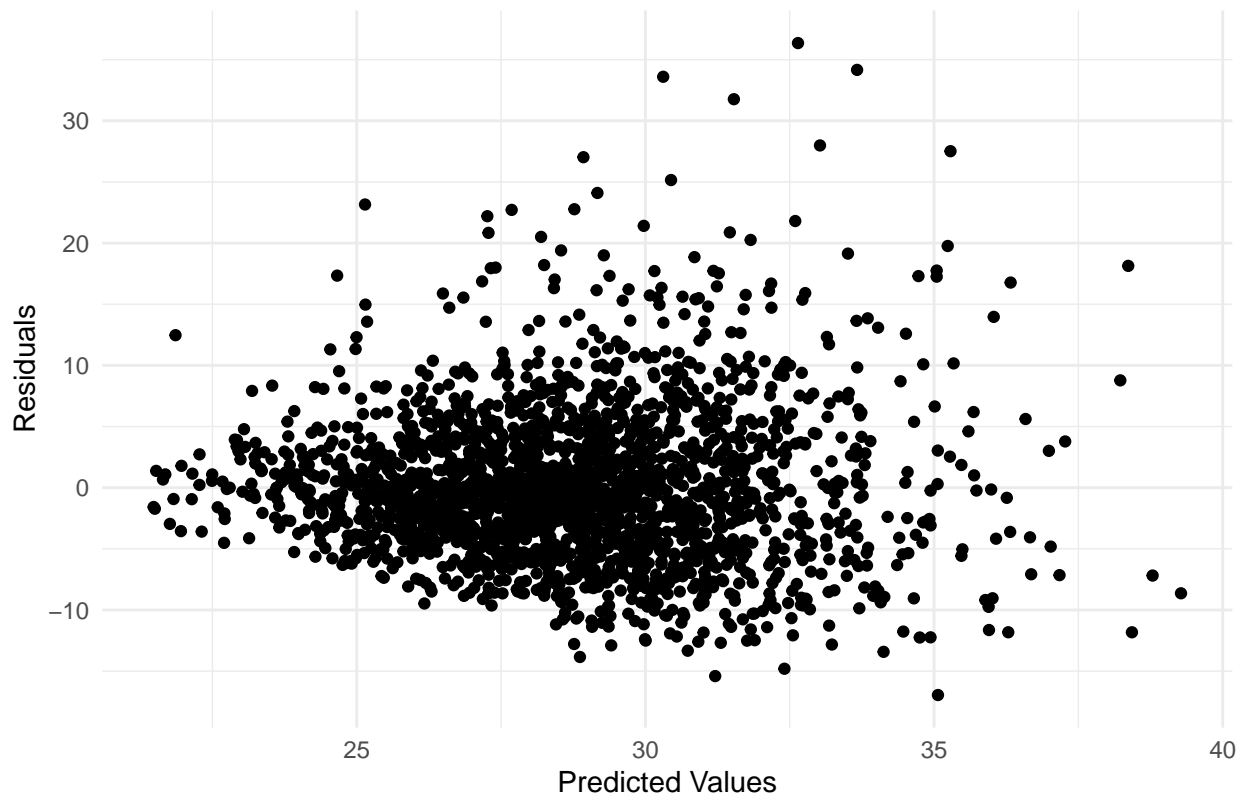
residual_1 <- m_full$residuals

# Get predicted values from the model
predicted_values <- predict(m_full)

# Create the plot
ggplot() +
  geom_point(aes(x = predicted_values, y = residual_1)) +
  ggtitle("Residuals vs. Predicted Values") +
  xlab("Predicted Values") +
  ylab("Residuals") +
  theme_minimal()

```

Residuals vs. Predicted Values



```

# Display the plot
print(ggplot)

## function (data = NULL, mapping = aes(), ..., environment = parent.frame())
## {
##   UseMethod("ggplot")
## }
## <bytecode: 0x4c54e60>
## <environment: namespace:ggplot2>

# Load necessary library
library(ggplot2)

# Assuming m_full is your linear model

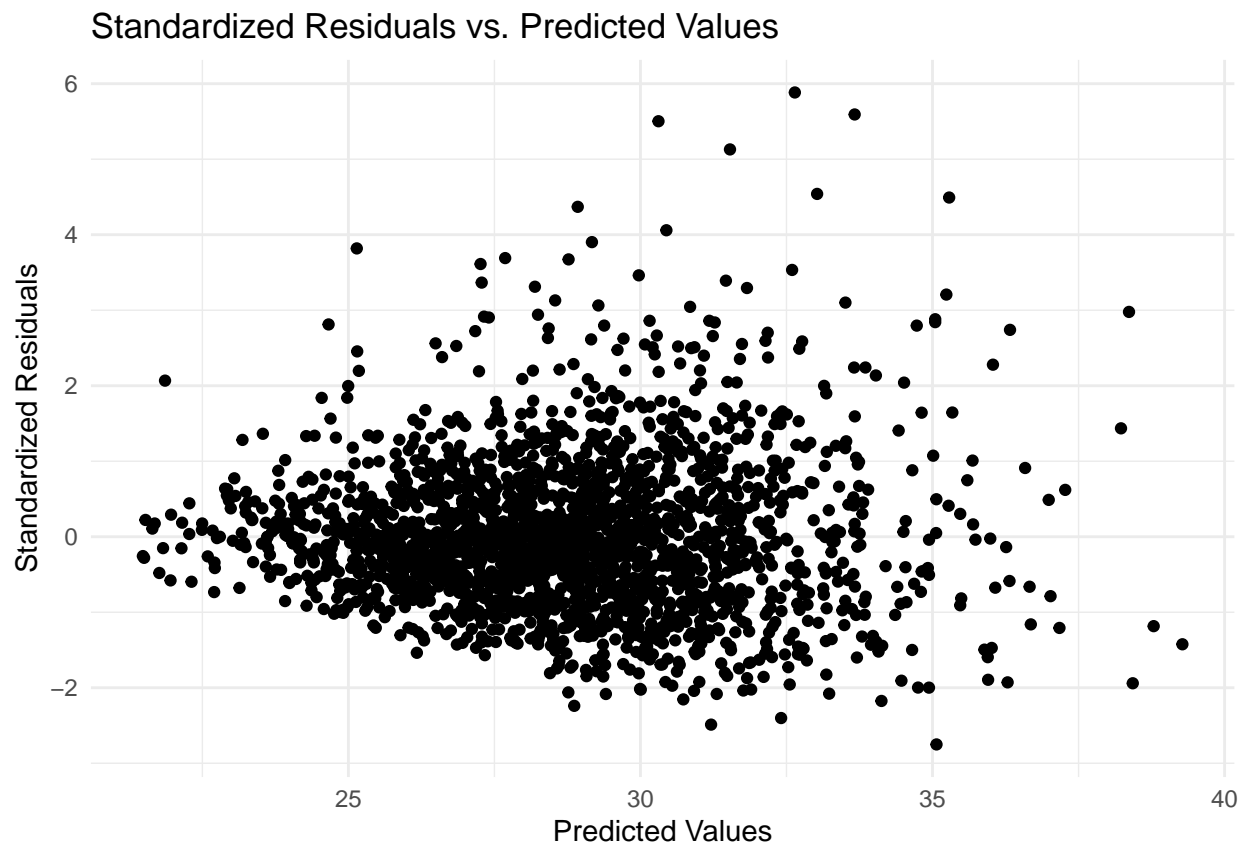
```

```
# m_full <- lm(SleepMinNight ~ ., data = df3)

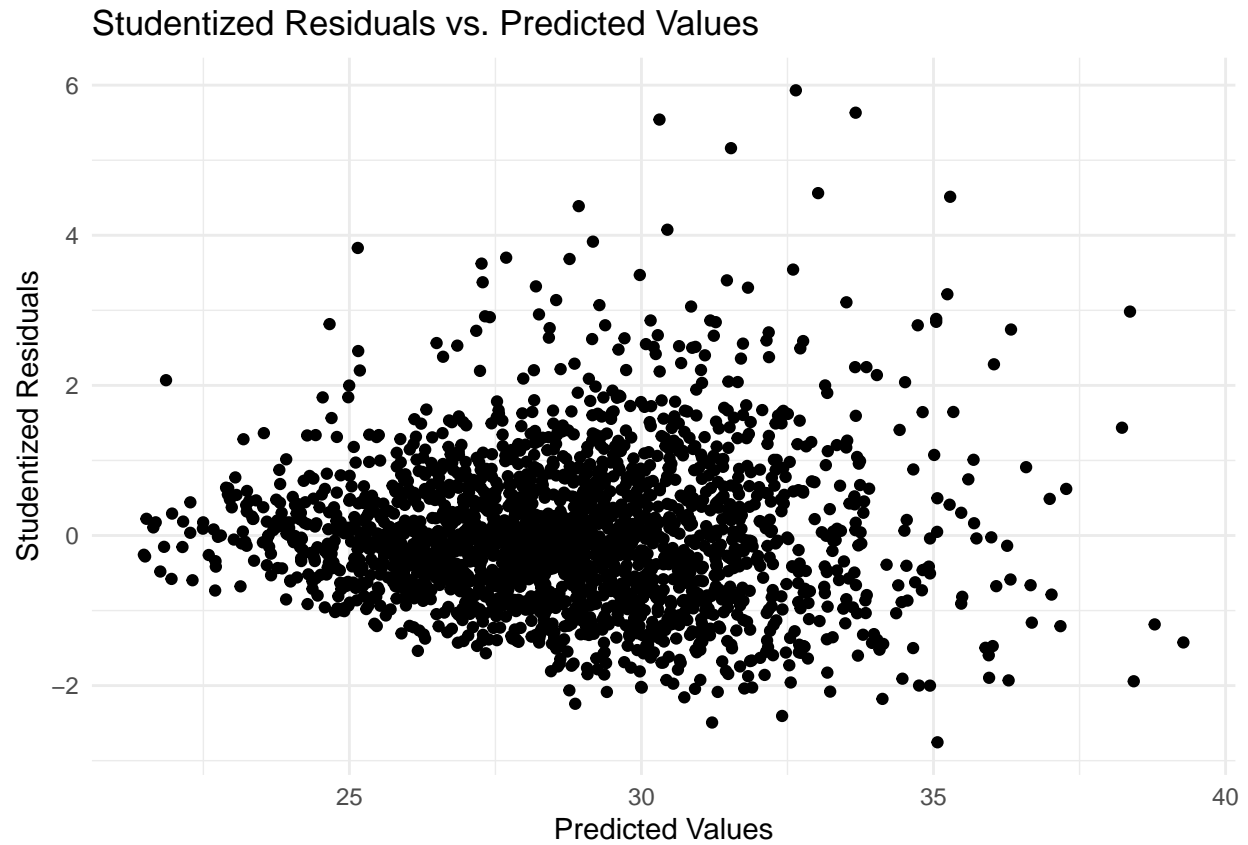
# Calculate different types of residuals
residual_2 <- rstandard(m_full)
residual_3 <- rstudent(m_full)
residual_4 <- rstudent(m_full) # Externally studentized residuals

# Get predicted values from the model
predicted_values <- predict(m_full)

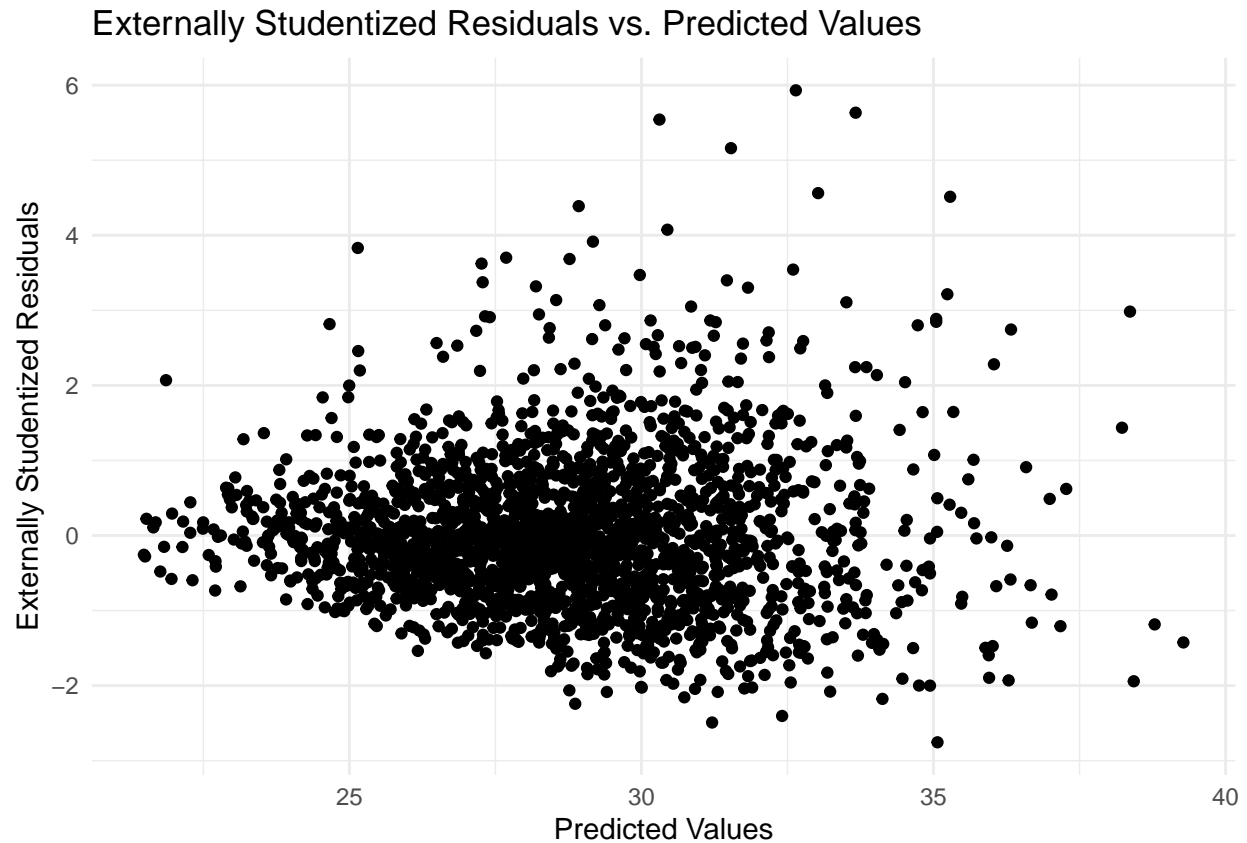
# Plot for Standardized Residuals
ggplot() +
  geom_point(aes(x = predicted_values, y = residual_2)) +
  ggtitle("Standardized Residuals vs. Predicted Values") +
  xlab("Predicted Values") +
  ylab("Standardized Residuals") +
  theme_minimal()
```



```
# Plot for Studentized Residuals
ggplot() +
  geom_point(aes(x = predicted_values, y = residual_3)) +
  ggtitle("Studentized Residuals vs. Predicted Values") +
  xlab("Predicted Values") +
  ylab("Studentized Residuals") +
  theme_minimal()
```



```
# Plot for Externally Studentized Residuals  
ggplot() +  
  geom_point(aes(x = predicted_values, y = residual_4)) +  
  ggtitle("Externally Studentized Residuals vs. Predicted Values") +  
  xlab("Predicted Values") +  
  ylab("Externally Studentized Residuals") +  
  theme_minimal()
```



## (5) Model Selection

```
step(m_full)
```

```
## Start:  AIC=7892.77
## BMI ~ SleepHrsNight + Age + Gender + Race1 + Poverty + TotChol +
##       BPDiaAve + BPSysAve + AlcoholYear + Smoke100 + UrineFlow1 +
##       DaysMentHlthBad + DaysPhysHlthBad + HealthGen + PhysActive +
##       SleepHrsNight * Age + SleepHrsNight * Gender + SleepHrsNight *
##       factor(Race1)
##
##
## Step:  AIC=7892.77
## BMI ~ SleepHrsNight + Age + Gender + Poverty + TotChol + BPDiaAve +
##       BPSysAve + AlcoholYear + Smoke100 + UrineFlow1 + DaysMentHlthBad +
##       DaysPhysHlthBad + HealthGen + PhysActive + factor(Race1) +
##       SleepHrsNight:Age + SleepHrsNight:Gender + SleepHrsNight:factor(Race1)
##
##
```

	Df	Sum of Sq	RSS	AIC
## - TotChol	1	0.4	82107	7890.8
## - Poverty	1	13.4	82120	7891.1
## - UrineFlow1	1	15.1	82122	7891.2
## - DaysPhysHlthBad	1	19.9	82127	7891.3
## <none>			82107	7892.8

```

## - SleepHrsNight:Age          1      78.0 82185 7892.8
## - DaysMentHlthBad           1     127.1 82234 7894.1
## - SleepHrsNight:factor(Race1) 4     430.5 82537 7896.0
## - SleepHrsNight:Gender       1     231.3 82338 7896.8
## - Smoke100                   1     336.7 82444 7899.6
## - PhysActive                 1     354.1 82461 7900.0
## - BPDiaAve                   1     689.3 82796 7908.8
## - BPSysAve                   1     758.2 82865 7910.6
## - AlcoholYear                1    1375.8 83483 7926.5
## - HealthGen                  4    4622.1 86729 8002.6
##
## Step: AIC=7890.78
## BMI ~ SleepHrsNight + Age + Gender + Poverty + BPDiaAve + BPSysAve +
##       AlcoholYear + Smoke100 + UrineFlow1 + DaysMentHlthBad + DaysPhysHlthBad +
##       HealthGen + PhysActive + factor(Race1) + SleepHrsNight:Age +
##       SleepHrsNight:Gender + SleepHrsNight:factor(Race1)
##
##              Df Sum of Sq  RSS    AIC
## - Poverty          1      13.4 82120 7889.1
## - UrineFlow1        1      15.0 82122 7889.2
## - DaysPhysHlthBad   1      19.9 82127 7889.3
## <none>              82107 7890.8
## - SleepHrsNight:Age  1      78.5 82186 7890.8
## - DaysMentHlthBad   1     127.4 82235 7892.1
## - SleepHrsNight:factor(Race1) 4     430.5 82538 7894.0
## - SleepHrsNight:Gender  1     231.6 82339 7894.8
## - Smoke100          1     338.1 82445 7897.6
## - PhysActive        1     354.6 82462 7898.1
## - BPDiaAve          1     698.7 82806 7907.0
## - BPSysAve          1     759.9 82867 7908.6
## - AlcoholYear       1    1377.1 83484 7924.6
## - HealthGen         4    4628.8 86736 8000.8
##
## Step: AIC=7889.13
## BMI ~ SleepHrsNight + Age + Gender + BPDiaAve + BPSysAve + AlcoholYear +
##       Smoke100 + UrineFlow1 + DaysMentHlthBad + DaysPhysHlthBad +
##       HealthGen + PhysActive + factor(Race1) + SleepHrsNight:Age +
##       SleepHrsNight:Gender + SleepHrsNight:factor(Race1)
##
##              Df Sum of Sq  RSS    AIC
## - UrineFlow1        1      13.2 82134 7887.5
## - DaysPhysHlthBad   1      19.3 82140 7887.6
## <none>              82120 7889.1
## - SleepHrsNight:Age  1      81.0 82201 7889.3
## - DaysMentHlthBad   1     133.0 82253 7890.6
## - SleepHrsNight:factor(Race1) 4     431.2 82552 7892.4
## - SleepHrsNight:Gender  1     228.7 82349 7893.1
## - PhysActive        1     342.9 82463 7896.1
## - Smoke100          1     373.7 82494 7896.9
## - BPDiaAve          1     702.9 82823 7905.5
## - BPSysAve          1     751.3 82872 7906.7
## - AlcoholYear       1    1363.8 83484 7922.6
## - HealthGen         4    4694.1 86815 8000.8
##

```



```

## Step: AIC=7887.48
## BMI ~ SleepHrsNight + Age + Gender + BPDiaAve + BPSysAve + AlcoholYear +
##      Smoke100 + DaysMentHlthBad + DaysPhysHlthBad + HealthGen +
##      PhysActive + factor(Race1) + SleepHrsNight:Age + SleepHrsNight:Gender +
##      SleepHrsNight:factor(Race1)
##
##              Df Sum of Sq  RSS    AIC
## - DaysPhysHlthBad      1      19.3 82153 7886.0
## <none>                                82134 7887.5
## - SleepHrsNight:Age      1      83.3 82217 7887.7
## - DaysMentHlthBad      1     134.4 82268 7889.0
## - SleepHrsNight:factor(Race1)  4     433.9 82568 7890.8
## - SleepHrsNight:Gender    1     229.6 82363 7891.5
## - PhysActive            1     352.7 82486 7894.7
## - Smoke100              1     372.5 82506 7895.2
## - BPDiaAve              1     705.1 82839 7903.9
## - BPSysAve              1     748.9 82883 7905.0
## - AlcoholYear           1    1388.6 83522 7921.6
## - HealthGen             4    4725.3 86859 7999.9
##
## Step: AIC=7885.98
## BMI ~ SleepHrsNight + Age + Gender + BPDiaAve + BPSysAve + AlcoholYear +
##      Smoke100 + DaysMentHlthBad + HealthGen + PhysActive + factor(Race1) +
##      SleepHrsNight:Age + SleepHrsNight:Gender + SleepHrsNight:factor(Race1)
##
##              Df Sum of Sq  RSS    AIC
## <none>                                82153 7886.0
## - SleepHrsNight:Age      1      82.2 82235 7886.1
## - DaysMentHlthBad      1     120.8 82274 7887.1
## - SleepHrsNight:factor(Race1)  4     432.0 82585 7889.3
## - SleepHrsNight:Gender    1     230.0 82383 7890.0
## - PhysActive            1     363.3 82516 7893.5
## - Smoke100              1     366.2 82519 7893.6
## - BPDiaAve              1     696.3 82849 7902.1
## - BPSysAve              1     750.4 82903 7903.5
## - AlcoholYear           1    1403.8 83557 7920.4
## - HealthGen             4    5179.9 87333 8009.6
##
## Call:
## lm(formula = BMI ~ SleepHrsNight + Age + Gender + BPDiaAve +
##      BPSysAve + AlcoholYear + Smoke100 + DaysMentHlthBad + HealthGen +
##      PhysActive + factor(Race1) + SleepHrsNight:Age + SleepHrsNight:Gender +
##      SleepHrsNight:factor(Race1), data = df3)
##
## Coefficients:
##              (Intercept)              SleepHrsNight
##              22.630757                -0.683663
##              Age              Gender
##             -0.079622              3.952743
##              BPDiaAve              BPSysAve
##              0.057703              0.051854
##             AlcoholYear             Smoke100Yes
##             -0.009057             -0.868816

```

##	DaysMentHlthBad	HealthGenVgood
##	-0.031240	1.870626
##	HealthGenGood	HealthGenFair
##	3.611653	5.361278
##	HealthGenPoor	PhysActiveYes
##	7.674279	-0.892385
##	factor(Race1)2	factor(Race1)3
##	-0.830286	-3.287026
##	factor(Race1)4	factor(Race1)5
##	-5.098631	1.043670
##	SleepHrsNight:Age	SleepHrsNight:Gender
##	0.013291	-0.506819
##	SleepHrsNight:factor(Race1)2	SleepHrsNight:factor(Race1)3
##	-0.169941	0.312389
##	SleepHrsNight:factor(Race1)4	SleepHrsNight:factor(Race1)5
##	0.545229	-0.627600

```
library(olsrr)
```

```
##
## Attaching package: 'olsrr'
## The following object is masked from 'package:datasets':
##
##     rivers
```

```
ols_step_forward_p(m_full, penter = 0.1, details = F)
```

[illegible]

```
## Warning in b * sx: longer object length is not a multiple of shorter object
## length
```

```
## Note: model has aliased coefficients
##      sums of squares computed by model comparison
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##
##
##                                     Selection Summary

```

##	Variable		Adj.				
##	Step	Entered	R-Square	R-Square	C(p)	AIC	RMSE
##	1	HealthGen	0.0826	0.0809	180.3262	14153.5414	6.4747
##	2	BPDiaAve	0.1066	0.1045	121.3681	14098.4490	6.3908
##	3	AlcoholYear	0.1226	0.1201	82.8326	14061.6280	6.3349
##	4	factor(Race1)	0.1381	0.1341	45.3890	14031.1683	6.2844
##	5	BPSysAve	0.1450	0.1406	29.8339	14015.8275	6.2606
##	6	Smoke100	0.1483	0.1436	23.4714	14009.5178	6.2500
##	7	PhysActive	0.1519	0.1467	16.5415	14002.6086	6.2386
##	8	SleepHrsNight:factor(Race1)	0.1567	0.1496	6.1651	14000.1995	6.2279
##	9	SleepHrsNight	0.1567	0.1496	8.1651	14000.1995	6.2279
##	10	Gender	0.1578	0.1503	7.3501	13999.3671	6.2252
##	11	SleepHrsNight:Gender	0.1602	0.1523	3.3336	13995.3009	6.2179
##	12	Poverty	0.1606	0.1523	4.4769	13996.4356	6.2181
##	13	Race1	0.1606	0.1523	4.4769	13998.4356	6.2181
##	14	DaysPhysHlthBad	0.1606	0.1520	6.2801	14000.2368	6.2193
##	15	Age	0.1608	0.1518	7.7645	14001.7160	6.2200
##	16	TotChol	0.1608	0.1514	9.7324	14003.6835	6.2214
##	17	UrineFlow1	0.1610	0.1512	11.2234	14005.1691	6.2222
##	18	DaysMentHlthBad	0.1623	0.1521	10.0165	14003.9255	6.2189
##							

```
ols_step_forward_p(m_full, penter = 0.05, details = F)
```

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##                               Selection Summary
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##      Variable                               Adj.

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##	4	factor(Race1)	0.1381	0.1341	45.3890	14031.1683	6.2844
##	5	BPSysAve	0.1450	0.1406	29.8339	14015.8275	6.2606
##	6	Smoke100	0.1483	0.1436	23.4714	14009.5178	6.2500
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##	18	DaysMentHlthBad	0.1623	0.1521	10.0165	14003.9255	6.2189

```
## -----
ols_mallows_cp(model = m_3, fullmodel = m_full) # Mallows' Cp
```

```
## [1] 36.32895
```