

# BIOSTAT 650 Project

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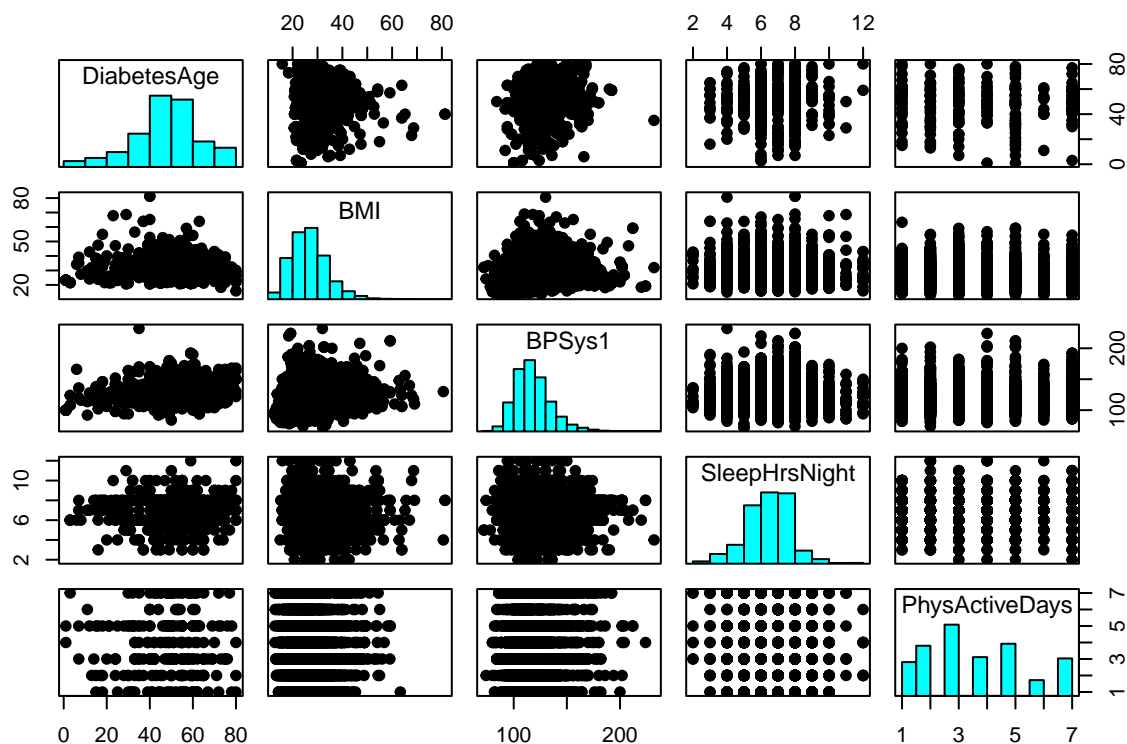
1 + 1

```
df = NHANES
#df = NHANES["DiabetesAge" > 20]
colnames(df)

## [1] "ID" "SurveyYr" "Gender" "Age"
## [5] "AgeDecade" "AgeMonths" "Race1" "Race3"
## [9] "Education" "MaritalStatus" "HHIncome" "HHIncomeMid"
## [13] "Poverty" "HomeRooms" "HomeOwn" "Work"
## [17] "Weight" "Length" "HeadCirc" "Height"
## [21] "BMI" "BMICatUnder20yrs" "BMI_WHO" "Pulse"
## [25] "BPSysAve" "BPDiaAve" "BPSys1" "BPDia1"
## [29] "BPSys2" "BPDia2" "BPSys3" "BPDia3"
## [33] "Testosterone" "DirectChol" "TotChol" "UrineVol1"
## [37] "UrineFlow1" "UrineVol2" "UrineFlow2" "Diabetes"
## [41] "DiabetesAge" "HealthGen" "DaysPhysHlthBad" "DaysMentHlthBad"
## [45] "LittleInterest" "Depressed" "nPregnancies" "nBabies"
## [49] "Age1stBaby" "SleepHrsNight" "SleepTrouble" "PhysActive"
## [53] "PhysActiveDays" "TVHrsDay" "CompHrsDay" "TVHrsDayChild"
## [57] "CompHrsDayChild" "Alcohol12PlusYr" "AlcoholDay" "AlcoholYear"
## [61] "SmokeNow" "Smoke100" "Smoke100n" "SmokeAge"
## [65] "Marijuana" "AgeFirstMarij" "RegularMarij" "AgeRegMarij"
## [69] "HardDrugs" "SexEver" "SexAge" "SexNumPartnLife"
## [73] "SexNumPartYear" "SameSex" "SexOrientation" "PregnantNow"

scatmatrixData = df[,c("DiabetesAge", "BMI", "BPSys1", "SleepHrsNight", "PhysActiveDays")]
panel.hist <- function(x, ...)
{
  usr <- par("usr"); on.exit(par(usr))
  par(usr = c(usr[1:2], 0, 1.5) )
  h <- hist(x, plot = FALSE)
  breaks <- h$breaks; nB <- length(breaks)
  y <- h$counts; y <- y/max(y)
  rect(breaks[-nB], 0, breaks[-1], y, col = "cyan", ...)
}
pairs(scatmatrixData, pch = 19, diag.panel=panel.hist)

## Warning in par(usr): argument 1 does not name a graphical parameter
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```



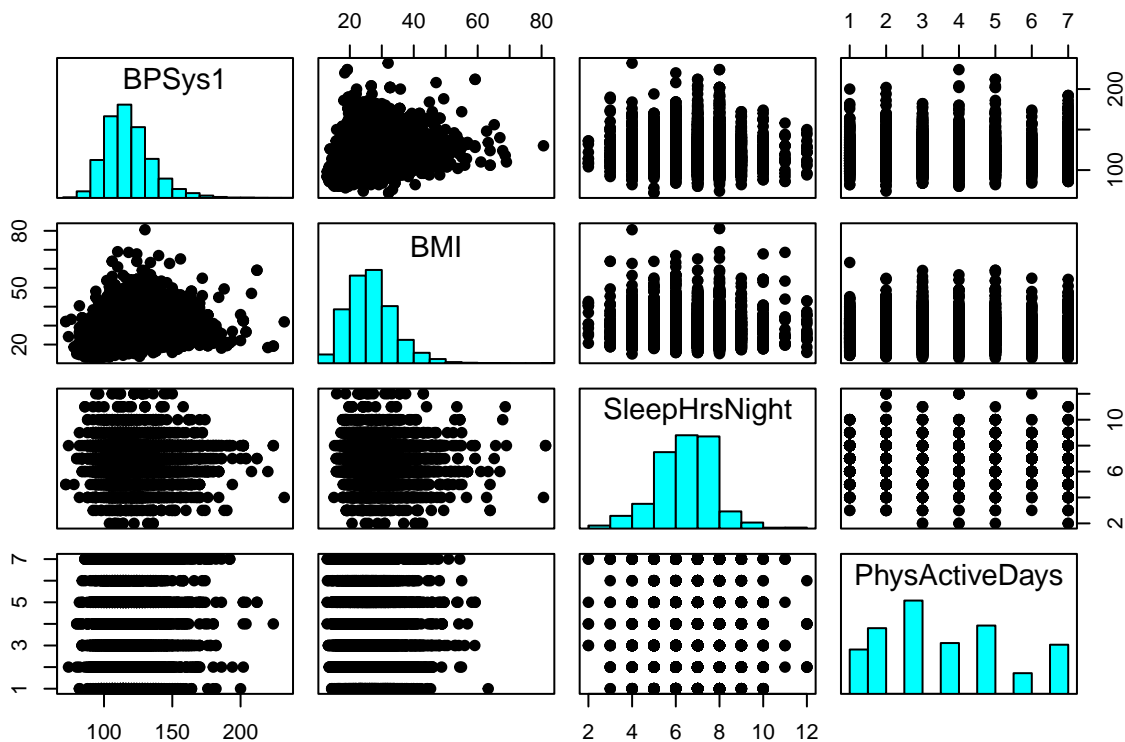
```
df = NHANES
#poverty gender useless?
colnames(df)
```

```
## [1] "ID" "SurveyYr" "Gender" "Age"
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```

```
scatmatrixData = df[,c("BPSys1", "BMI", "SleepHrsNight",
  "PhysActiveDays")]
panel.hist <- function(x, ...)
```

```
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par(usr = c(usr[1:2], 0, 1.5) )
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breaks <- h$breaks; nB <- length(breaks)
y <- h$counts; y <- y/max(y)
rect(breaks[-nB], 0, breaks[-1], y, col = "cyan", ...)
}
pairs(scatmatrixData, pch = 19, diag.panel=panel.hist)
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```



```
model <- lm(DiabetesAge ~ Gender+Poverty+BMI+BPSys1+SleepHrsNight+PhysActiveDays, df)
summary(model)
```

```
##
## Call:
## lm(formula = DiabetesAge ~ Gender + Poverty + BMI + BPSys1 +
##     SleepHrsNight + PhysActiveDays, data = df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -44.087  -7.907   2.062   8.861  29.318
```

```
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)  32.96048   10.92836   3.016  0.00287 **
## Gendermale   -2.46465    2.11661  -1.164  0.24553
## Poverty      0.46344    0.62309   0.744  0.45781
## BMI          -0.09236    0.14055  -0.657  0.51180
## BPSys1       0.13469    0.05758   2.339  0.02024 *
## SleepHrsNight 0.25571    0.73547   0.348  0.72841
## PhysActiveDays -0.19888    0.53308  -0.373  0.70945
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 15.09 on 217 degrees of freedom
## (9776 observations deleted due to missingness)
## Multiple R-squared:  0.04008, Adjusted R-squared:  0.01354
## F-statistic:  1.51 on 6 and 217 DF, p-value: 0.176

model <- lm(BPSys1 ~ Age+Gender+Poverty+BMI+SleepHrsNight+PhysActiveDays+SmokeNow+AlcoholYear+HardDrugs
summary(model)

##
## Call:
## lm(formula = BPSys1 ~ Age + Gender + Poverty + BMI + SleepHrsNight +
##     PhysActiveDays + SmokeNow + AlcoholYear + HardDrugs, data = df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -39.397  -8.387  -0.997   7.730  69.906
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept)  89.959564   3.820975  23.544 < 2e-16 ***
## Age          0.413402   0.035437  11.666 < 2e-16 ***
## Gendermale    5.382522   0.903317   5.959 3.48e-09 ***
## Poverty      -0.843665   0.283924  -2.971  0.00303 **
## BMI          0.345235   0.075337   4.583 5.15e-06 ***
## SleepHrsNight 0.247155   0.331007   0.747  0.45543
## PhysActiveDays -0.021275   0.244823  -0.087  0.93077
## SmokeNowYes   1.325291   0.957252   1.384  0.16651
## AlcoholYear   0.002536   0.004169   0.608  0.54318
## HardDrugsYes  0.141125   0.964282   0.146  0.88367
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 14.18 on 1038 degrees of freedom
## (8952 observations deleted due to missingness)
## Multiple R-squared:  0.1709, Adjusted R-squared:  0.1637
## F-statistic: 23.78 on 9 and 1038 DF, p-value: < 2.2e-16

model <- lm(SexAge ~ HardDrugs, df)
summary(model)

##
## Call:
```

```

## lm(formula = SexAge ~ HardDrugs, data = df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -8.804 -1.804 -0.804  1.196 32.196
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  17.80354    0.05435  327.59  <2e-16 ***
## HardDrugsYes -1.91152    0.12383  -15.44  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.631 on 5527 degrees of freedom
## (4471 observations deleted due to missingness)
## Multiple R-squared:  0.04133,    Adjusted R-squared:  0.04116
## F-statistic: 238.3 on 1 and 5527 DF,  p-value: < 2.2e-16

```