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West

W 9

QA

```
# Read in packages and data
suppressPackageStartupMessages({
  library(tidyverse)
  library(NHANES)
  library(neuralnet)
})
```

1).

```
# Create indicator variable RM1 and Sex1
data <- NHANES %>%
  mutate(RM1 = ifelse(RegularMarij=="Yes",1,0), Sex1 = ifelse(Gender=="male",1,0))

# Summarize indicators
data %>%
  select(RM1, Sex1) %>%
  summary()
```

```
##           RM1           Sex1
##  Min.      :0.000   Min.      :0.000
##  1st Qu.:0.000   1st Qu.:0.000
##  Median :0.000   Median :0.000
##  Mean    :0.276   Mean    :0.498
##  3rd Qu.:1.000   3rd Qu.:1.000
##  Max.    :1.000   Max.    :1.000
##  NA's    :5059
```

2).

```
# Limit dataset to HardDrugs, RM1, Age, BMI and Sex1
# Scale Age and BMI to be between 0-1
data <- data %>%
  select(HardDrugs, RM1, Age, BMI, Sex1) %>%
  mutate_at(vars(Age, BMI), funs(scale(.)) %>% as.vector())

data %>% glimpse()
```

```
## Observations: 10,000
## Variables: 5
## $ HardDrugs <fct> Yes, Yes, Yes, NA, Yes, NA, NA, No, No, No, No, Yes,...
## $ RM1      <dbl> 0, 0, 0, NA, 0, NA, NA, 0, 0, 0, NA, 1, 1, NA, NA, 0...
```

```
## $ Age      <dbl> -0.1224285, -0.1224285, -0.1224285, -1.4618598, 0.54...
## $ BMI      <dbl> 0.753718485, 0.753718485, 0.753718485, -1.540027677,...
## $ Sex1     <dbl> 1, 1, 1, 1, 0, 1, 1, 0, 0, 0, 1, 1, 1, 0, 0, 1, 0, 1...
```

3).

```
# remove missing values
data <- data %>% na.omit()

# get length of remaining columns
data %>% lapply(length)
```

```
## $HardDrugs
## [1] 4907
##
## $RM1
## [1] 4907
##
## $Age
## [1] 4907
##
## $BMI
## [1] 4907
##
## $Sex1
## [1] 4907
```

4).

```
# Set random seed
set.seed(1847)
p <- 0.2
m <- 4907

# get random indices and split train and test data
train_ind <- sample.int(m, (1-p)*m)
traind <- data[train_ind,]
testd <- data[-train_ind,]

traind %>% glimpse()
```

```
## Observations: 3,925
## Variables: 5
## $ HardDrugs <fct> No, No, No, Yes, No, No, No, No, Yes, No, No, No, No...
## $ RM1      <dbl> 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0...
## $ Age      <dbl> 0.90446882, 0.41334403, -0.56890557, -0.47961016, -0...
## $ BMI      <dbl> -0.3687530, -0.5504091, 1.7270694, 0.3849839, -0.143...
## $ Sex1     <dbl> 0, 1, 0, 0, 1, 0, 1, 0, 1, 1, 0, 0, 0, 0, 0, 1, 0, 0...

testd %>% glimpse()
```

```
## Observations: 982
## Variables: 5
```

```
## $ HardDrugs <fct> Yes, No, No, Yes, No, No, No, No, Yes, Yes, No, No, ...
## $ RM1 <dbl> 0, 0, 0, 1, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1, 0...
## $ Age <dbl> -0.12242848, 0.36869632, 0.36869632, 0.77052570, 0.0...
## $ BMI <dbl> 0.75371848, 0.07860880, 0.07860880, -0.08542388, 1.2...
## $ Sex1 <dbl> 1, 0, 0, 1, 1, 1, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0, 1, 0...
```

5).

```
# Set random seed
set.seed(1847)

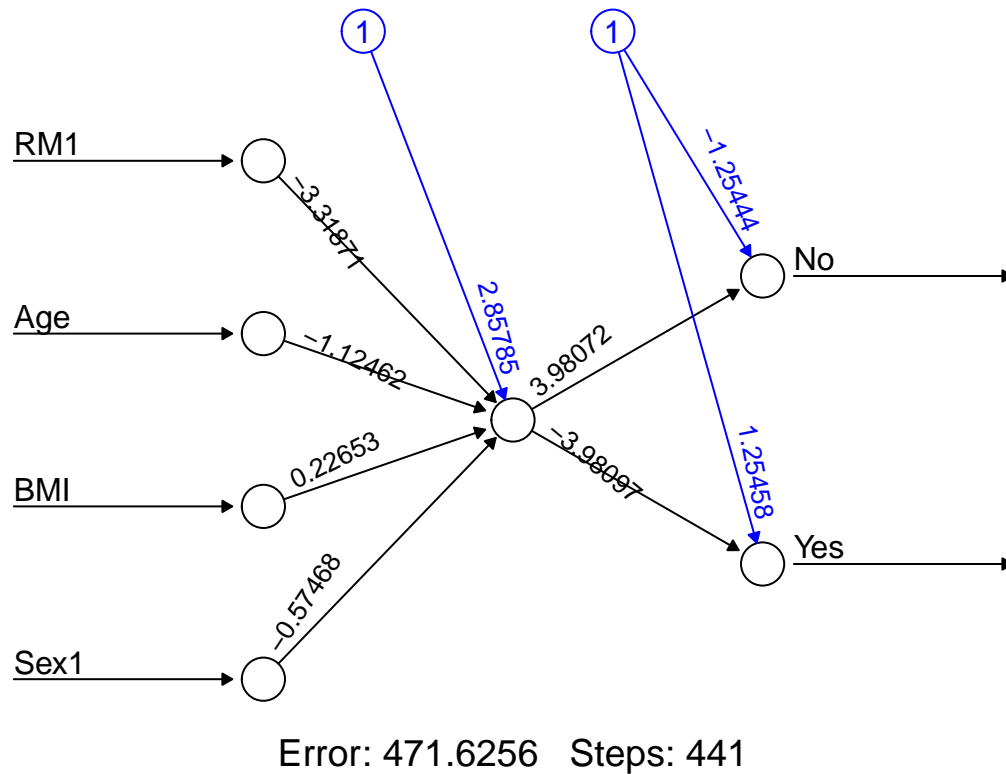
# create formula
f <- formula(HardDrugs ~ RM1 + Age + BMI + Sex1)

# fit the NN on the training data
mNN <- neuralnet(f, traintd, hidden=1, linear.output=F, rep=10, lifesign="minimal")
```

| | | | | | |
|--------------|--------------|------------|-------------|------------------|-----------------|
| ## hidden: 1 | thresh: 0.01 | rep: 1/10 | steps: 2657 | error: 471.62653 | time: 2.85 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 2/10 | steps: 1434 | error: 471.62676 | time: 1.54 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 3/10 | steps: 1166 | error: 471.62645 | time: 1.25 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 4/10 | steps: 567 | error: 471.62666 | time: 0.61 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 5/10 | steps: 1469 | error: 471.62667 | time: 1.59 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 6/10 | steps: 1481 | error: 471.62639 | time: 1.63 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 7/10 | steps: 441 | error: 471.6256 | time: 0.47 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 8/10 | steps: 909 | error: 471.62642 | time: 0.98 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 9/10 | steps: 602 | error: 471.62635 | time: 0.64 secs |
| ## hidden: 1 | thresh: 0.01 | rep: 10/10 | steps: 2242 | error: 471.62608 | time: 2.41 secs |

6).

```
# plot network form for result
plot(mNN, rep="best")
```



7).

```
# get predictions on test data for the best network result
p1 <- compute(mNN, testd, rep=7)

# create confusion matrix
table(round(p1$net.result[,2]), ifelse(testd$HardDrugs=="Yes",1,0))

##
##      0    1
## 0 717 113
## 1  59  93
```

8).

```
# set random seed
set.seed(1847)

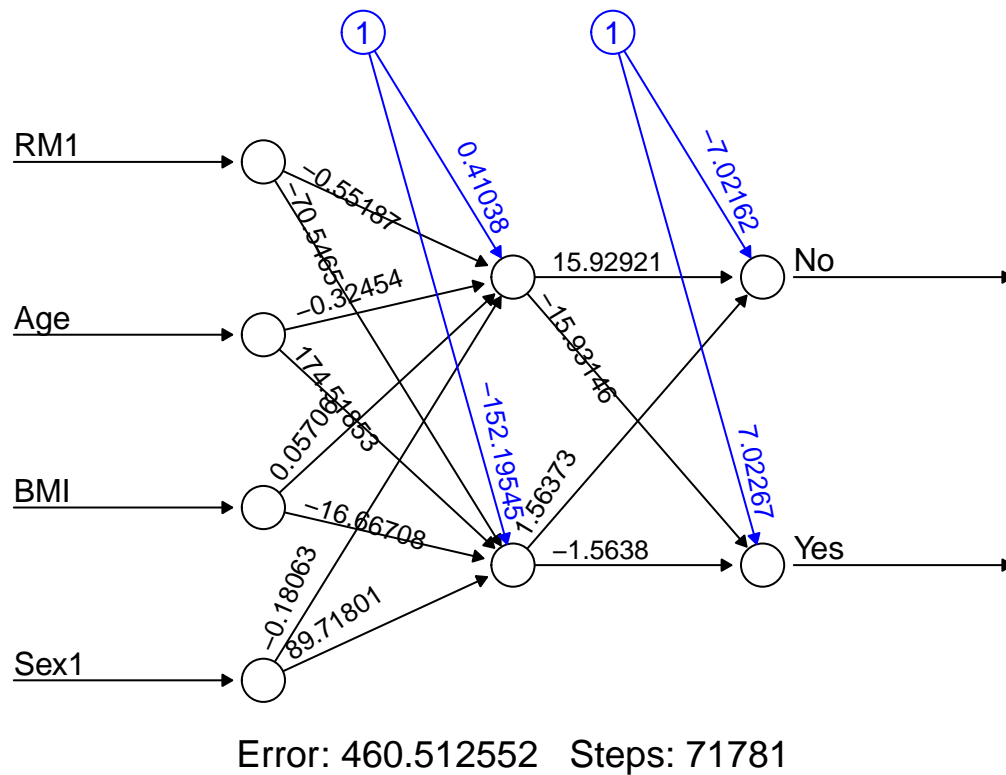
# fit the NN on the training data with 2 hidden layers
mNN2 <- neuralnet(f, traind, hidden=2, linear.output=F, rep=10, lifesign="minimal")

## hidden: 2    thresh: 0.01    rep: 1/10    steps: 16242    error: 462.56932    time: 23.32 secs
## hidden: 2    thresh: 0.01    rep: 2/10    steps: 3332     error: 467.32647    time: 4.64 secs
## hidden: 2    thresh: 0.01    rep: 3/10    steps: 16335    error: 462.54016    time: 23 secs
```

| hidden | thresh | rep | steps | error | time |
|--------|--------|-------|-------|-----------|------------|
| 2 | 0.01 | 4/10 | 71781 | 460.51255 | 1.74 mins |
| 2 | 0.01 | 5/10 | 17179 | 462.54186 | 24.37 secs |
| 2 | 0.01 | 6/10 | 4714 | 465.14661 | 6.56 secs |
| 2 | 0.01 | 7/10 | 4571 | 465.05289 | 6.35 secs |
| 2 | 0.01 | 8/10 | 12005 | 467.35928 | 16.89 secs |
| 2 | 0.01 | 9/10 | 1257 | 471.63031 | 1.79 secs |
| 2 | 0.01 | 10/10 | 5176 | 467.32646 | 7.46 secs |

9).

```
# plot network form for result
plot(mNN2, rep="best")
```



10).

```
# get predictions on test data for the best network result
p2 <- compute(mNN2, testd, rep=4)

# create confusion matrix
table(round(p2$net.result[,2]), ifelse(testd$HardDrugs=="Yes",1,0))
```

```
##
##      0      1
## 0 720 126
```

1 56 80