

# Week11

Mathi Manavalan

4/6/2020

## Libraries

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.0 --

## v ggplot2 3.3.0      v purrr  0.3.3
## v tibble  2.1.3      v dplyr  0.8.4
## v tidyr   1.0.0      v stringr 1.4.0
## v readr   1.3.1      v forcats 0.4.0

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

```
library(Hmisc)
```

```
## Loading required package: lattice

## Loading required package: survival

## Loading required package: Formula

##
## Attaching package: 'Hmisc'

## The following objects are masked from 'package:dplyr':
##
##      src, summarize

## The following objects are masked from 'package:base':
##
##      format.pval, units
```

## Data Import and Cleaning

First, I am importing the entire dataset.

```
data <- as_tibble(spss.get("../data/GSS2006.sav", use.value.labels=TRUE))
```

Now, we only want the variables relevant to the personality inventory and the respondent's self-reported health. For the personality inventory, I am interpreting this to mean the variables (according to the code book for this dataset) from BIG5A1 to BIG5E2 from *The 2006 Module: Personality Traits*. For the respondent's self-reported health, I am interpreting this to mean the HEALTH variable.

```
clean <- data %>%
  select(BIG5A1, BIG5A2, BIG5B1, BIG5B2, BIG5C1, BIG5C2, BIG5D1, BIG5D2, BIG5E1, BIG5E2, HEALTH) %>%
  na.omit() %>%
  mutate(HEALTH = HEALTH %>%
    factor() %>%
    #fct_rev() %>%
    as.numeric()
  )
```

Responses of 'don't know', 'inapplicable', or other unclearly answered items are appropriately marked as NA according to R, so I removed all rows of data that contain NA (in any column(s)).

All of the variables are in their appropriate data type (factors), with corresponding levels.

## Analysis

```
set.seed(1)

# Shuffle row indices: rows
rows <- sample(nrow(clean))

# Randomly order data
shuffled <- clean[rows, ]

holdout <- clean[1:250, ]

train <- clean[251:nrow(clean), ]

olsr <- lm(HEALTH ~ ., holdout)
olsr
```

```
##
## Call:
## lm(formula = HEALTH ~ ., data = holdout)
##
## Coefficients:
##                (Intercept)                BIG5A1Agree
##                2.144651                -0.070409
## BIG5A1Neither agree nor disagree                BIG5A1Disagree
##                0.000958                -0.043901
##                BIG5A1Strongly disagree                BIG5A2Agree
##                -0.224848                0.173522
```

## BIG5A2Neither agree nor disagree	BIG5A2Disagree
## 0.400288	-0.021829
## BIG5A2Strongly disagree	BIG5B1Agree
## 0.362036	-0.264212
## BIG5B1Neither agree nor disagree	BIG5B1Disagree
## -0.307978	0.139656
## BIG5B1Strongly disagree	BIG5B2Agree
## -0.652772	0.509861
## BIG5B2Neither agree nor disagree	BIG5B2Disagree
## 0.469689	0.421304
## BIG5B2Strongly disagree	BIG5C1Agree
## 0.654539	0.193824
## BIG5C1Neither agree nor disagree	BIG5C2Agree
## 0.731099	-0.884251
## BIG5C2Neither agree nor disagree	BIG5C2Disagree
## -0.935055	-0.666508
## BIG5C2Strongly disagree	BIG5D1Agree
## -0.704198	0.021551
## BIG5D1Neither agree nor disagree	BIG5D1Disagree
## 0.189085	0.243709
## BIG5D1Strongly disagree	BIG5D2Agree
## 1.321893	0.401722
## BIG5D2Neither agree nor disagree	BIG5D2Disagree
## 0.210145	0.133441
## BIG5D2Strongly disagree	BIG5E1Agree
## 0.219446	0.085454
## BIG5E1Neither agree nor disagree	BIG5E1Disagree
## -0.104056	0.216474
## BIG5E1Strongly disagree	BIG5E2Agree
## -0.340226	-0.402309
## BIG5E2Neither agree nor disagree	BIG5E2Disagree
## -0.268693	-0.338218
## BIG5E2Strongly disagree	
## -0.167624	

## Visualization