Correlation Matrices, Reliability Tests, and t-Tests

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<pre>library(mosaic) library(ggplot2) library(dplyr)</pre>		
<pre>#install.packages("] library(psych)</pre>	psych")	

Data inside R packages

To demonstrate the use of the alpha() fuction in the psych package from computing Cronbach's alpha, we will need a data set with scale items. Many packages have example datasets pre-loaded. The psych package has a dataset called bfi that has n = 2800 participants responding to 25 personality items for the Big Five Inventory.

```
#We can see a list of ALL the datasets current avaliable data()
```

If we see a dataset we like we can load it into our environment

```
data(bfi)
#?bfi
glimpse(bfi)
## Observations: 2 800
```

```
## $ A2
               <int> 4, 4, 4, 4, 3, 6, 5, 3, 3, 5, 4, 5, 5, 5, 5, 3, 6, 5...
## $ A3
               <int> 3, 5, 5, 6, 3, 5, 5, 1, 6, 6, 5, 5, 5, 5, 2, 6, 6, 5...
## $ A4
               <int> 4, 2, 4, 5, 4, 6, 3, 5, 3, 6, 6, 5, 6, 6, 2, 6, 2, 4...
## $ A5
               <int> 4, 5, 4, 5, 5, 5, 5, 1, 3, 5, 5, 5, 4, 6, 1, 3, 5, 5...
## $ C1
               <int> 2, 5, 4, 4, 4, 6, 5, 3, 6, 6, 4, 5, 5, 4, 5, 5, 4, 5...
## $ C2
               <int> 3, 4, 5, 4, 4, 6, 4, 2, 6, 5, 3, 4, 4, 4, 5, 5, 4, 5...
## $ C3
               <int> 3, 4, 4, 3, 5, 6, 4, 4, 3, 6, 5, 5, 3, 4, 5, 5, 4, 5...
## $ C4
               <int> 4, 3, 2, 5, 3, 1, 2, 2, 4, 2, 3, 4, 2, 2, 2, 3, 4, 4...
## $ C5
               <int> 4, 4, 5, 5, 2, 3, 3, 4, 5, 1, 2, 5, 2, 1, 2, 5, 4, 3...
## $ E1
               <int> 3, 1, 2, 5, 2, 2, 4, 3, 5, 2, 1, 3, 3, 2, 3, 1, 1,
## $ E2
               <int> 3, 1, 4, 3, 2, 1, 3, 6, 3, 2, 3, 3, 3, 2, 4, 1, 2, 2...
               <int> 3, 6, 4, 4, 5, 6, 4, 4, NA, 4, 2, 4, 3, 4, 3, 6, 5, ...
## $ E3
## $ E4
               <int> 4, 4, 4, 4, 4, 5, 5, 2, 4, 5, 5, 5, 2, 6, 6, 6, 5, 6...
## $ E5
               <int> 4, 3, 5, 4, 5, 6, 5, 1, 3, 5, 4, 4, 4, 5, 5, 4, 5, 6...
## $ N1
               <int> 3, 3, 4, 2, 2, 3, 1, 6, 5, 5, 3, 4, 1, 1, 2, 4, 4, 6...
## $ N2
               <int> 4, 3, 5, 5, 3, 5, 2, 3, 5, 5, 3, 5, 2, 1, 4, 5, 4, 5...
## $ N3
               <int> 2, 3, 4, 2, 4, 2, 2, 2, 5, 4, 3, 2, 1, 2, 4, 4, 5...
## $ N4
               <int> 2, 5, 2, 4, 4, 2, 1, 6, 3, 2, 2, 2, 2, 2, 2, 5, 4, 4...
## $ N5
               <int> 3, 5, 3, 1, 3, 3, 1, 4, 3, 4, 3, NA, 2, 1, 3, 5, 5, ...
               <int> 3, 4, 4, 3, 3, 4, 5, 3, 6, 5, 5, 4, 4, 5, 5, 6, 5, 5...
## $ 01
## $ 02
               <int> 6, 2, 2, 3, 3, 3, 2, 2, 6, 1, 3, 6, 2, 3, 2, 6, 1, 1...
## $ 03
               <int> 3, 4, 5, 4, 4, 5, 5, 4, 6, 5, 5, 4, 4, 4, 5, 6, 5, 4...
## $ 04
               <int> 4, 3, 5, 3, 3, 6, 6, 5, 6, 5, 6, 5, 5, 4, 5, 3, 6, 5...
## $ 05
               <int> 3, 3, 2, 5, 3, 1, 1, 3, 1, 2, 3, 4, 2, 4, 5, 2, 3, 4...
## $ gender
               <int> 1, 2, 2, 2, 1, 2, 1, 1, 1, 2, 1, 1, 2, 1, 1, 1, 2, 1...
## $ education <int> NA, NA, NA, NA, NA, 3, NA, 2, 1, NA, 1, NA, NA, NA, ...
## $ age
               <int> 16, 18, 17, 17, 17, 21, 18, 19, 19, 17, 21, 16, 16, ...
```

Correlation Matrix

First, we can get descriptive statistics for individual personality scale items. Let's do the agreeableness scale first.

```
#Get descriptive statistics with mosaic or dplyr here. Get them for Age too.

#Also, get the frequencies for gender and education
```

In addition to descriptive statistics, we would probably also like to get a correlation matrix for items on a scale. I like the function corr.test() in the psych package. Notice that we are using the dplyr select() function inside of corr.test().

```
corr.test(select(bfi, A1, A2, A3, A4, A5))
## Call:corr.test(x = select(bfi, A1, A2, A3, A4, A5))
## Correlation matrix
```

```
1.00 -0.34 -0.27 -0.15
## A1
                              -0.18
## A2 -0.34
             1.00 0.49
                         0.34
                               0.39
## A3 -0.27
             0.49
                   1.00
                         0.36
                               0.50
## A4 -0.15
            0.34 0.36
                         1.00
                               0.31
## A5 -0.18 0.39 0.50 0.31
                               1.00
## Sample Size
##
        Α1
             A2
                  АЗ
                       A4
                            A5
## A1 2784 2757 2759 2767 2769
## A2 2757 2773 2751 2758 2757
## A3 2759 2751 2774 2759 2758
## A4 2767 2758 2759 2781 2765
## A5 2769 2757 2758 2765 2784
## Probability values (Entries above the diagonal are adjusted for multiple tests.)
      A1 A2 A3 A4 A5
##
## A1
       0
          0
             0
                0
## A2
       0
          0
             0
                0
                   0
## A3
       0
          0
             0
                0
                   0
## A4
          0
             0
                0
                   0
       0
## A5
       0 0
             0
                0
                  0
##
   To see confidence intervals of the correlations, print with the short=FALSE option
```

##

A1

A2

АЗ

A4

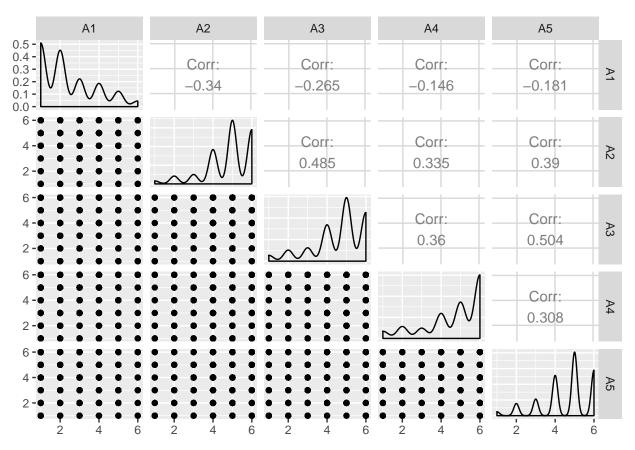
A5

```
#install.packages("GGally")
library(GGally)

ggpairs(select(bfi, A1, A2, A3, A4, A5))
```

There is also a pairs() function in Base R and the ggpairs() function in the GGally

package that is a ggplot correlation matrix/matrix scatterplot provider.



It is clear that we will need to reverse score A1 to assess reliability and to create our agreeableness scale scores. Hint: to reverse score an item use this formula: A1.r = (min(A1, na.rm = TRUE) + max(A1, na.rm = TRUE)) - A1.

Now compute correlation matrices for the other 4 personality subscales and create any reverse scored items the need to be created.

```
#correlation matrices and reverse scored items
```

For more practice, find another dataset that sounds interesting and run some correlation matricies.

data()

Cronbach's alpha

Now that we have our reverse scored items and we have a sense of how the items relate, the next step is to test the scale reliability. We do this with Cronbach's alpha. There is a function called alpha() in the psych package.

```
alpha(select(bfi, A1.r, A2, A3, A4, A5))
```

##

```
## Reliability analysis
  Call: alpha(x = select(bfi, A1.r, A2, A3, A4, A5))
##
##
     raw_alpha std.alpha G6(smc) average_r S/N
                                                    ase mean sd
##
          0.7
                    0.71
                            0.68
                                       0.33 2.5 0.009
                                                       4.7 0.9
##
##
    lower alpha upper
                           95% confidence boundaries
## 0.69 0.7 0.72
##
##
    Reliability if an item is dropped:
##
        raw_alpha std.alpha G6(smc) average_r S/N alpha se
## A1.r
             0.72
                        0.73
                                0.67
                                           0.40 2.6
                                                       0.0087
## A2
             0.62
                        0.63
                                0.58
                                           0.29 1.7
                                                      0.0119
                                           0.28 1.6
## A3
             0.60
                        0.61
                                0.56
                                                       0.0124
## A4
             0.69
                        0.69
                                0.65
                                           0.36 2.3
                                                      0.0098
## A5
             0.64
                        0.66
                                0.61
                                           0.32 1.9
                                                      0.0111
##
##
    Item statistics
##
           n raw.r std.r r.cor r.drop mean
## A1.r 2784 0.58
                                  0.31
                     0.57
                           0.38
                                         4.6 1.4
## A2
        2773
                                  0.56
              0.73
                     0.75
                           0.67
                                         4.8 1.2
        2774 0.76
## A3
                     0.77
                           0.71
                                  0.59
                                         4.6 1.3
## A4
        2781
              0.65
                     0.63
                           0.47
                                  0.39
                                         4.7 1.5
## A5
        2784 0.69
                     0.70
                           0.60
                                  0.49
                                         4.6 1.3
##
## Non missing response frequency for each item
           1
                 2
                      3
                           4
##
                                5
## A1.r 0.03 0.08 0.12 0.14 0.29 0.33 0.01
        0.02 0.05 0.05 0.20 0.37 0.31 0.01
## A2
        0.03 0.06 0.07 0.20 0.36 0.27 0.01
## A3
## A4
        0.05 0.08 0.07 0.16 0.24 0.41 0.01
        0.02 0.07 0.09 0.22 0.35 0.25 0.01
## A5
```

With an alphas of .70 we have a reliabile agreeableness scale. The next step would be to create agreeableness scale scores for participants, the average of these 5 items. We can do this with mutate(). Note that we cannot use mean() here because mean() is going to get the mean of a column (a variable). What we want it the mean across columns.

```
bfi <- bfi %>%
  mutate(agreeable = (A1.r + A2 + A3 + A4 + A5)/5)
```

Let's take a look at it with a visualization. Also try facetting by gender.

```
#make some histograms, boxplots, and density plots of agreeableness
```

Now calculate the reliabilities for each of the other 4 scales. Be sure to include the reverse scored items you created where relevant.

#calculate reliabilities

Create scale scores and make visualizations of those scale scores.

```
#scale scores and visualizations
```

t-Tests

Independent Samples t-Test

Next we can ask, are men and women different on agreeableness? To answer this question we need to run an independent samples t-test. We'll use the function t.test() in the mosaic package. Recall that mosaic first needs the formula, y ~ x, then the data, data = dataName.

```
t.test(agreeable ~ gender, data = bfi)
```

```
##
## Welch Two Sample t-test
##
## data: agreeable by gender
## t = -10.725, df = 1654.5, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.4698058 -0.3245337
## sample estimates:
## mean in group 1 mean in group 2
## 4.377679 4.774848</pre>
```

There is a statistically significant difference between women and men on agreeableness, t(1654.50) = -10.73, p < .001, with women (M = 4.77, SD = 0.86) scoring higher than men (M = 4.38, SD = 0.93). It is possible to code in these numbers such that if the data were updated, the text would update as well.

```
tmod <- t.test(agreeable ~ gender, data = bfi)
ds <- favstats(agreeable ~ gender, data = bfi)</pre>
```

There is a statistically significant difference between women and men on agreeableness, t(1654.47) = -10.72, p < 0.001, with women (M = 4.77, SD = 0.86) scoring higher than men (M = 4.38, SD = 0.93).

We can also ask for equal variance assumed.

```
t.test(agreeable ~ gender, data = bfi, var.equal = TRUE)
```

##

```
## Two Sample t-test
##
## data: agreeable by gender
## t = -11.038, df = 2707, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.4677230 -0.3266165
## sample estimates:
## mean in group 1 mean in group 2
## 4.377679 4.774848</pre>
```

Test for gender differences for the other 4 scales. Test for gender differences between those who graduated college, and those who did not. Hint: use mutate() to create a dichotomous variable as we did for wise_hus.

```
#more t-Tests
```

Paired Samples t-Test

To demonstrate paired samples t-test I'd like to use yet another built in dataset. The sat.act dataset has information for 700 people on their SAT verbal, SAT quantitative, and ACT scores.

```
data(sat.act)
#?sat.act
glimpse(sat.act)
## Observations: 700
## Variables: 6
## $ gender
               <int> 2, 2, 2, 1, 1, 1, 2, 1, 2, 2, 1, 2, 1, 2, 2, 2, 2, 2...
## $ education <int> 3, 3, 3, 4, 2, 5, 5, 3, 4, 5, 3, 4, 4, 4, 3, 4, 3, 4...
## $ age
               <int> 19, 23, 20, 27, 33, 26, 30, 19, 23, 40, 23, 34, 32, ...
## $ ACT
               <int> 24, 35, 21, 26, 31, 28, 36, 22, 22, 35, 32, 29, 21, ...
## $ SATV
               <int> 500, 600, 480, 550, 600, 640, 610, 520, 400, 730, 76...
## $ SATQ
               <int> 500, 500, 470, 520, 550, 640, 500, 560, 600, 800, 71...
```

Let's compare poeple's scores on the SAT verbal to their scores on the SAT quantitative using the t.test() function. This time instead of using the formula in the beginning, we add the two variables to be compared separated by a \tt . We also need to add paired = TRUE.

```
t.test(SATV, SATQ, data = sat.act, paired = TRUE)
```

```
##
## Paired t-test
##
## data: SATV and SATQ
## t = 0.57483, df = 686, p-value = 0.5656
```

```
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -5.116109 9.351917
## sample estimates:
## mean of the differences
## 2.117904
```

There is no staistically significant difference between SAT verbal and SAT quantitative scores, t(686) = 0.57, p = .566. But perhaps there is a difference for boys only, or for girls only. We can create a dataset of just boys using filter() and then another one of just girls. This would be completely fine, but we could also pipe directly into the t.test() function. There is a catch: The pipe, %>%, puts the resulting dataframe into the first position in the destination function, but we want it in the third position. We can use the . symbol to pipe the dataframe into whichever position we'd like.

```
sat.act %>%
  filter(gender == 1) %>%
  t.test(SATV, SATQ, data = ., paired = TRUE)
##
##
   Paired t-test
##
## data: SATV and SATQ
## t = -3.4934, df = 244, p-value = 0.0005661
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -32.08114 -8.94743
## sample estimates:
## mean of the differences
##
                 -20.51429
sat.act %>%
  filter(gender == 2) %>%
  t.test(SATV, SATQ, data = ., paired = TRUE)
##
##
   Paired t-test
##
## data: SATV and SATQ
## t = 3.1813, df = 441, p-value = 0.00157
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
     5.604249 23.721543
##
## sample estimates:
## mean of the differences
##
                   14.6629
```

Explore the differences in ${\tt SATV}$ and ${\tt SATQ}$ for each level of the ${\tt education}$ variable.

 $\#differences\ for\ each\ level\ of\ education$