Reliability Testing

```
In [1]: # Install and load the required libraries
        install.packages("psych")
        library(psych)
        # Read the modified CSV file
        modified_data <- read.csv("Complete_Data_Modified.csv")</pre>
        # Extract relevant columns for reliability analysis
        # Replace "Scale1", "Scale2", etc., with the actual names of your scales
        scales <- modified_data[, c("Panas...", "Panas...1", "BFI..E.", "BFI..A.", "BFI..</pre>
                                     "EI..Self.A.", "EI..Self.M.", "EI..Social.A.", "EI..
        # Perform Cronbach's alpha analysis
        cronbach_result <- psych::alpha(scales, check.keys = TRUE)</pre>
        # Display the Cronbach's alpha results
        print(cronbach_result)
        # Extract reliability coefficients
        reliabilities <- cronbach_result$total$raw_alpha
        # Display reliability coefficients for each scale
        cat("\nReliability coefficients for each scale:\n")
        print(reliabilities)
        Updating HTML index of packages in '.Library'
        Making 'packages.html' ...
         done
        Warning message in psych::alpha(scales, check.keys = TRUE):
        "Some items were negatively correlated with the first principal component and w
        ere automatically reversed.
        This is indicated by a negative sign for the variable name."
```

```
Reliability analysis
Call: psych::alpha(x = scales, check.keys = TRUE)
```

raw_alpha std.alpha G6(smc) average_r S/N ase mean sd median_r 0.66 0.76 0.81 0.18 3.1 0.019 64 2.5 0.14

95% confidence boundaries

lower alpha upper

Feldt 0.59 0.66 0.73 Duhachek 0.62 0.66 0.70

Reliability if an item is dropped:

| | raw_alpha | std.alpha | G6(smc) | average_r | S/N | alpha se | var.r | med.r |
|-------------|-----------|-----------|---------|-----------|-----|----------|-------|-------|
| Panas | 0.66 | 0.76 | 0.81 | 0.20 | 3.2 | 0.019 | 0.033 | 0.16 |
| Panas1 | 0.66 | 0.77 | 0.81 | 0.20 | 3.3 | 0.019 | 0.033 | 0.16 |
| BFIE | 0.66 | 0.76 | 0.81 | 0.19 | 3.1 | 0.019 | 0.035 | 0.16 |
| BFIA | 0.66 | 0.73 | 0.78 | 0.17 | 2.7 | 0.019 | 0.034 | 0.13 |
| BFIC | 0.66 | 0.74 | 0.79 | 0.18 | 2.9 | 0.019 | 0.033 | 0.14 |
| BFIN. | 0.66 | 0.75 | 0.80 | 0.19 | 3.1 | 0.019 | 0.034 | 0.14 |
| BFIO | 0.66 | 0.74 | 0.80 | 0.18 | 2.9 | 0.019 | 0.035 | 0.14 |
| EISelf.A. | 0.60 | 0.74 | 0.79 | 0.18 | 2.8 | 0.019 | 0.032 | 0.14 |
| EISelf.M. | 0.59 | 0.71 | 0.76 | 0.16 | 2.5 | 0.020 | 0.027 | 0.14 |
| EISocial.A. | 0.56 | 0.72 | 0.77 | 0.17 | 2.6 | 0.023 | 0.027 | 0.14 |
| EIRM. | 0.55 | 0.72 | 0.77 | 0.17 | 2.6 | 0.024 | 0.027 | 0.14 |
| Total | 0.64 | 0.70 | 0.74 | 0.15 | 2.4 | 0.020 | 0.020 | 0.14 |
| PAQ- | 0.66 | 0.76 | 0.81 | 0.20 | 3.2 | 0.019 | 0.033 | 0.16 |
| CBCL | 0.66 | 0.75 | 0.81 | 0.19 | 3.1 | 0.019 | 0.035 | 0.14 |

Item statistics

n raw.r std.r r.cor r.drop mean 206 0.13 0.29 0.18 0.111 99.4 0.59 Panas..-Panas...1 206 0.11 0.24 0.12 0.096 1.2 0.42 BFI..E.- 206 0.18 0.34 0.23 0.164 99.1 0.45 BFI..A.- 206 0.37 0.59 0.55 0.356 99.3 0.54 BFI..C.- 206 0.32 0.50 0.45 0.310 99.3 0.50 BFI..N. BFI..N. 206 0.19 0.38 0.30 0.180 2.2 0.53 BFI..O.- 206 0.34 0.49 0.42 0.327 99.0 0.38 EI..Self.A. 206 0.65 0.52 0.47 0.489 78.0 7.59 EI..Self.M. 206 0.81 0.75 0.78 0.588 69.9 13.30 EI..Social.A. 206 0.79 0.64 0.65 0.637 76.4 9.42 EI..RM. 206 0.82 0.66 0.67 0.651 70.9 11.44 206 0.93 0.81 0.88 0.928 3.0 0.95 Total PAO-206 0.17 0.28 0.17 0.151 99.5 CBCL 206 0.16 0.38 0.29 0.143 2.5 0.72

Non missing response frequency for each item

1 2 3 4 5 miss 0.44 0.50 0.05 0.00 0.00 Panas.. Panas...1 0.77 0.23 0.00 0.00 0.00 BFI..E. 0.16 0.79 0.06 0.00 0.00 BFI..A. 0.37 0.60 0.03 0.00 0.00 BFI..C. 0.35 0.64 0.01 0.00 0.00 BFI..N. 0.05 0.67 0.27 0.00 0.00 BFI..O. 0.08 0.85 0.06 0.00 0.00 Total 0.05 0.26 0.32 0.35 0.02 PAO 0.62 0.22 0.16 0.00 0.00 0 CBCL 0.14 0.27 0.59 0.00 0.00

Reliability coefficients for each scale:

[1] 0.6610361

```
In [2]: # Cronbach Alpha Tests

# Calculate Cronbach's alpha
alpha_result <- alpha(modified_data, check.keys = TRUE)
cronbach_alpha <- alpha_result$total$raw_alpha

# Print the Cronbach's alpha value
cat("The Results of Cronbach Analysis is as Follows\n\n")
print(cronbach_alpha)
print(alpha_result)

cat("\n\n\n\n")</pre>
```

Warning message in alpha(modified_data, check.keys = TRUE):
"Some items were negatively correlated with the first principal component and w ere automatically reversed.

This is indicated by a negative sign for the variable name."

The Results of Cronbach Analysis is as Follows

```
[1] 0.6610361
```

```
Reliability analysis
```

Call: alpha(x = modified_data, check.keys = TRUE)

95% confidence boundaries

lower alpha upper

Feldt 0.59 0.66 0.73 Duhachek 0.62 0.66 0.70

Reliability if an item is dropped:

| | raw_alpha | std.alpha | G6(smc) | average_r | S/N | alpha se | var.r | med.r |
|-------------|-----------|-----------|---------|-----------|-----|----------|-------|-------|
| Panas | 0.66 | 0.76 | 0.81 | 0.20 | 3.2 | 0.019 | 0.033 | 0.16 |
| Panas1 | 0.66 | 0.77 | 0.81 | 0.20 | 3.3 | 0.019 | 0.033 | 0.16 |
| BFIE | 0.66 | 0.76 | 0.81 | 0.19 | 3.1 | 0.019 | 0.035 | 0.16 |
| BFIA | 0.66 | 0.73 | 0.78 | 0.17 | 2.7 | 0.019 | 0.034 | 0.13 |
| BFIC | 0.66 | 0.74 | 0.79 | 0.18 | 2.9 | 0.019 | 0.033 | 0.14 |
| BFIN. | 0.66 | 0.75 | 0.80 | 0.19 | 3.1 | 0.019 | 0.034 | 0.14 |
| BFIO | 0.66 | 0.74 | 0.80 | 0.18 | 2.9 | 0.019 | 0.035 | 0.14 |
| EISelf.A. | 0.60 | 0.74 | 0.79 | 0.18 | 2.8 | 0.019 | 0.032 | 0.14 |
| EISelf.M. | 0.59 | 0.71 | 0.76 | 0.16 | 2.5 | 0.020 | 0.027 | 0.14 |
| EISocial.A. | 0.56 | 0.72 | 0.77 | 0.17 | 2.6 | 0.023 | 0.027 | 0.14 |
| EIRM. | 0.55 | 0.72 | 0.77 | 0.17 | 2.6 | 0.024 | 0.027 | 0.14 |
| Total | 0.64 | 0.70 | 0.74 | 0.15 | 2.4 | 0.020 | 0.020 | 0.14 |
| PAQ- | 0.66 | 0.76 | 0.81 | 0.20 | 3.2 | 0.019 | 0.033 | 0.16 |
| CBCL | 0.66 | 0.75 | 0.81 | 0.19 | 3.1 | 0.019 | 0.035 | 0.14 |

Item statistics

 n raw.r std.r r.cor r.drop mean
 sd

 Panas...
 206
 0.13
 0.29
 0.18
 0.111
 99.4
 0.59

 Panas...
 206
 0.11
 0.24
 0.12
 0.096
 1.2
 0.42

 BFI..E..
 206
 0.18
 0.34
 0.23
 0.164
 99.1
 0.45

 BFI..A..
 206
 0.37
 0.59
 0.55
 0.356
 99.3
 0.54

 BFI..O..
 206
 0.32
 0.50
 0.45
 0.310
 99.3
 0.50

 BFI..O..
 206
 0.19
 0.38
 0.30
 0.180
 2.2
 0.53

 BFI..O..
 206
 0.34
 0.49
 0.42
 0.327
 99.0
 0.38

 EI..Self.A.
 206
 0.65
 0.52
 0.47
 0.489
 78.0
 7.59

 EI..Self.M.
 206
 0.81
 0.75
 0.78
 0.588
 69.9
 13.30

 EI..Scocial.A.
 206
 0.82
 0.66
 0.67
 0.651
 70.9
 11.44

 Total
 206
 0.93
 0.81
 <

Non missing response frequency for each item

```
5 miss
            1
                2
                     3 4
         0.44 0.50 0.05 0.00 0.00
Panas...1 0.77 0.23 0.00 0.00 0.00
BFI..E. 0.16 0.79 0.06 0.00 0.00
BFI..A. 0.37 0.60 0.03 0.00 0.00
BFI..C. 0.35 0.64 0.01 0.00 0.00
BFI..N. 0.05 0.67 0.27 0.00 0.00
                                   0
BFI..O.
         0.08 0.85 0.06 0.00 0.00
         0.05 0.26 0.32 0.35 0.02
Total
PAO
         0.62 0.22 0.16 0.00 0.00
                                   0
```

In [4]: # Install and load the required libraries

```
install.packages("psych")
library(psych)
# Read the modified CSV file
modified data <- read.csv("Complete Data Modified.csv")</pre>
# Define your variables
independent_vars <- c('BFI..E.', 'BFI..A.', 'BFI..C.', 'BFI..N.', 'BFI..O.',</pre>
                        'EI..Self.A.', 'EI..Self.M.', 'EI..Social.A.', 'EI..RM.',
dependent_vars <- c('Panas...', 'Panas...1', 'CBCL')</pre>
mediator var <- 'PAQ'
# Function to calculate Cronbach's alpha for a group of variables or a single va
calculate_alpha <- function(variables) {</pre>
  if (length(variables) > 1) {
    scale_subset <- modified_data[, variables]</pre>
    cronbach_result <- psych::alpha(scale_subset, check.keys = TRUE)</pre>
    return(cronbach result$total$raw alpha)
  } else {
    # If it's a single variable, just return the variable name
    return(variables)
  }
}
# Calculate Cronbach's alpha for each group of variables
alpha_independent <- calculate_alpha(independent_vars)</pre>
alpha_dependent <- calculate_alpha(dependent_vars)</pre>
alpha_mediator <- calculate_alpha(mediator_var)</pre>
# Display the results
cat("Cronbach's alpha for Independent Variables:\n")
print(alpha_independent)
cat("\nCronbach's alpha for Dependent Variables:\n")
print(alpha_dependent)
cat("\nCronbach's alpha for Mediator Variable:\n")
print(alpha mediator)
Updating HTML index of packages in '.Library'
Making 'packages.html' ...
 done
Warning message in psych::alpha(scale_subset, check.keys = TRUE):
"Some items were negatively correlated with the first principal component and w
ere automatically reversed.
This is indicated by a negative sign for the variable name."
Warning message in psych::alpha(scale_subset, check.keys = TRUE):
"Some items were negatively correlated with the first principal component and w
ere automatically reversed.
This is indicated by a negative sign for the variable name."
```

```
Cronbach's alpha for Independent Variables:
[1] 0.6748856

Cronbach's alpha for Dependent Variables:
[1] 0.2910346

Cronbach's alpha for Mediator Variable:
[1] "PAQ"

In []:
```