

# RECSM: Quantitative Methods in Social Research

Day 3 - 04 07 2025

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## Measurement of the Unobserved Variables and Scale Construction.

### Final Problem Set

**Step 1:** Open a new script and save it as day3. Clear your workspace.

**Step 2:** Load the General Social Survey (gss2016.RData) dataset and install “Psych” alongside “psychTools” package.

**Step 3:** Once you subset the ten variables related to confidence in institutions, you can explore the correlations among those variables in the dataset.

**Step 4:** Try to create a new variable that combines all 10 likert-scale variables (hint: use rowMeans function).

**Step 5:** Test the reliability of this scale through Cronbach’s alpha (hint: use “alpha” function).

**Step 6:** Test for the number of factors in your data using parallel analysis.

**Step 7:** Do the same analysis above with Principle Component Analysis (PCA).

**Step 8:** Use factor analysis with the data. Then compare the solution to a hierarchical cluster analysis using the “ICLUST” algorithm and function (Revelle, 1979).

### Solutions for problem sets

```
#Remove objects from the environment
rm(list=ls())
#Set your working directory
setwd("~/Downloads/RECSM workshop")
#Load the dataset
load("gss2016.RData")
#Install the package psych
install.packages( "psych" , repos = "http://cran.rstudio.com/" )

## Installing package into '/Users/buraksonmez/Library/R/arm64/4.3/library'
## (as 'lib' is unspecified)

##
## The downloaded binary packages are in
## /var/folders/f6/gb9b_pqd0yg5t19h5791zmjr0000gp/T//RtmpptS3Dbp downloaded_packages
```

```

install.packages( "psychTools" , repos = "http://cran.rstudio.com/" )

## Installing package into '/Users/buraksonmez/Library/R/arm64/4.3/library'
## (as 'lib' is unspecified)

##
## The downloaded binary packages are in
## /var/folders/f6/gb9b_pqd0yg5t19h5791zmjr0000gp/T//RtmpptS3Dbp downloaded_packages

install.packages( "corrplot" , repos = "http://cran.rstudio.com/" )

## Installing package into '/Users/buraksonmez/Library/R/arm64/4.3/library'
## (as 'lib' is unspecified)

##
## The downloaded binary packages are in
## /var/folders/f6/gb9b_pqd0yg5t19h5791zmjr0000gp/T//RtmpptS3Dbp downloaded_packages

library(psych)

## Warning: package 'psych' was built under R version 4.3.3

library(psychTools)

## Warning: package 'psychTools' was built under R version 4.3.3

library(corrplot)

## Warning: package 'corrplot' was built under R version 4.3.3

## corrplot 0.95 loaded

## Subsetting some variables related to confidence in institutions.
confvars <- c("confinan", "conbus", "conclerg", "coneduc",
"compress", "contv", "conjudge", "consci",
"conlegis", "conarmy")
confdata <- gss[confvars]
lowerCor(confdata)

##          cnfnn conbs cnclr condc cnprs contv cnjdg consc cnlgs cnrmy
## confinan 1.00
## conbus   0.35  1.00
## conclerg 0.28  0.29  1.00
## coneduc  0.25  0.21  0.23  1.00
## compress 0.15  0.15  0.11  0.21  1.00
## contv    0.26  0.22  0.14  0.24  0.37  1.00
## conjudge 0.22  0.25  0.13  0.26  0.22  0.20  1.00
## consci   0.07  0.15  0.02  0.13  0.13  0.10  0.31  1.00
## conlegis 0.33  0.24  0.21  0.29  0.21  0.22  0.34  0.13  1.00
## conarmy  0.31  0.22  0.20  0.20  0.12  0.15  0.20  0.17  0.21  1.00

```

```

## Find out relatively stronger correlations.
round(cor(confdata, use='pairwise'), 2)

##          confinan conbus conclerg coneduc compress contv conjudge consci
## confinan    1.00   0.35    0.28   0.25    0.15   0.26    0.22   0.07
## conbus      0.35   1.00    0.29   0.21    0.15   0.22    0.25   0.15
## conclerg    0.28   0.29    1.00   0.23    0.11   0.14    0.13   0.02
## coneduc     0.25   0.21    0.23   1.00    0.21   0.24    0.26   0.13
## compress    0.15   0.15    0.11   0.21    1.00   0.37    0.22   0.13
## contv       0.26   0.22    0.14   0.24    0.37   1.00    0.20   0.10
## conjudge    0.22   0.25    0.13   0.26    0.22   0.20    1.00   0.31
## consci      0.07   0.15    0.02   0.13    0.13   0.10    0.31   1.00
## conlegis    0.33   0.24    0.21   0.29    0.21   0.22    0.34   0.13
## conarmy     0.31   0.22    0.20   0.20    0.12   0.15    0.20   0.17
##          conlegis conarmy
## confinan    0.33   0.31
## conbus      0.24   0.22
## conclerg    0.21   0.20
## coneduc     0.29   0.20
## compress    0.21   0.12
## contv       0.22   0.15
## conjudge    0.34   0.20
## consci      0.13   0.17
## conlegis    1.00   0.21
## conarmy     0.21   1.00

cor <- cor(confdata)

#At first glance, there appears to be some
#association among confidence in different institutions.

## Creating a variable combining 10 likert scale questions.
confdata$con.scale <- rowMeans(confdata, na.rm=FALSE)
range(confdata$con.scale, na.rm=T)

## [1] 1 3

#Find coefficient alpha as an estimate of reliability. This may be done for a single scale. Interpret the results.
alpha(confdata)

## Number of categories should be increased in order to count frequencies.

## 
## Reliability analysis
## Call: alpha(x = confdata)
## 
##   raw_alpha std.alpha G6(smc) average_r S/N      ase mean      sd median_r
##   0.77        0.8     0.96      0.27    4 0.0063  2.1 0.35      0.22
## 
##   95% confidence boundaries
##           lower alpha upper

```

```

## Feldt      0.76  0.77  0.79
## Duhacheck 0.76  0.77  0.79
##
## Reliability if an item is dropped:
##          raw_alpha std.alpha G6(smc) average_r S/N alpha se  var.r med.r
## confinan      0.75      0.78     0.89      0.26 3.6  0.0070 0.0223  0.22
## conbus        0.75      0.79     0.90      0.27 3.7  0.0069 0.0234  0.22
## conclerg      0.77      0.80     0.89      0.28 3.9  0.0065 0.0221  0.22
## coneduc       0.76      0.79     0.89      0.27 3.7  0.0068 0.0236  0.22
## compress      0.77      0.79     0.89      0.28 3.9  0.0066 0.0225  0.24
## contv         0.76      0.79     0.89      0.27 3.7  0.0067 0.0228  0.22
## conjudge      0.75      0.78     0.88      0.27 3.6  0.0069 0.0230  0.22
## consci        0.78      0.80     0.90      0.29 4.1  0.0063 0.0208  0.24
## conlegis      0.75      0.78     0.89      0.27 3.6  0.0069 0.0230  0.22
## conarmy       0.76      0.79     0.90      0.28 3.8  0.0066 0.0235  0.24
## con.scale     0.72      0.72     0.72      0.21 2.6  0.0077 0.0058  0.21
##
## Item statistics
##          n raw.r std.r r.cor r.drop mean    sd
## confinan 1946  0.60  0.60  0.59   0.47  2.2 0.65
## conbus   1926  0.57  0.57  0.56   0.44  2.0 0.59
## conclerg 1914  0.50  0.49  0.47   0.34  2.1 0.68
## coneduc  1948  0.57  0.56  0.55   0.43  1.9 0.66
## compress 1937  0.50  0.50  0.48   0.35  2.4 0.64
## contv    1938  0.54  0.54  0.53   0.40  2.3 0.65
## conjudge 1915  0.59  0.58  0.58   0.45  1.9 0.67
## consci   1884  0.41  0.41  0.39   0.26  1.6 0.60
## conlegis 1925  0.59  0.59  0.58   0.47  2.5 0.61
## conarmy  1937  0.51  0.51  0.50   0.37  1.5 0.63
## con.scale 1792  1.00  1.00  1.02   0.99  2.1 0.34

```

```

##Test for the number of factors in your data using parallel analysis
fa.parallel(confdata)

```

```

## Warning in fa.stats(r = r, f = f, phi = phi, n.obs = n.obs, np.obs = np.obs, :
## The estimated weights for the factor scores are probably incorrect. Try a
## different factor score estimation method.

```

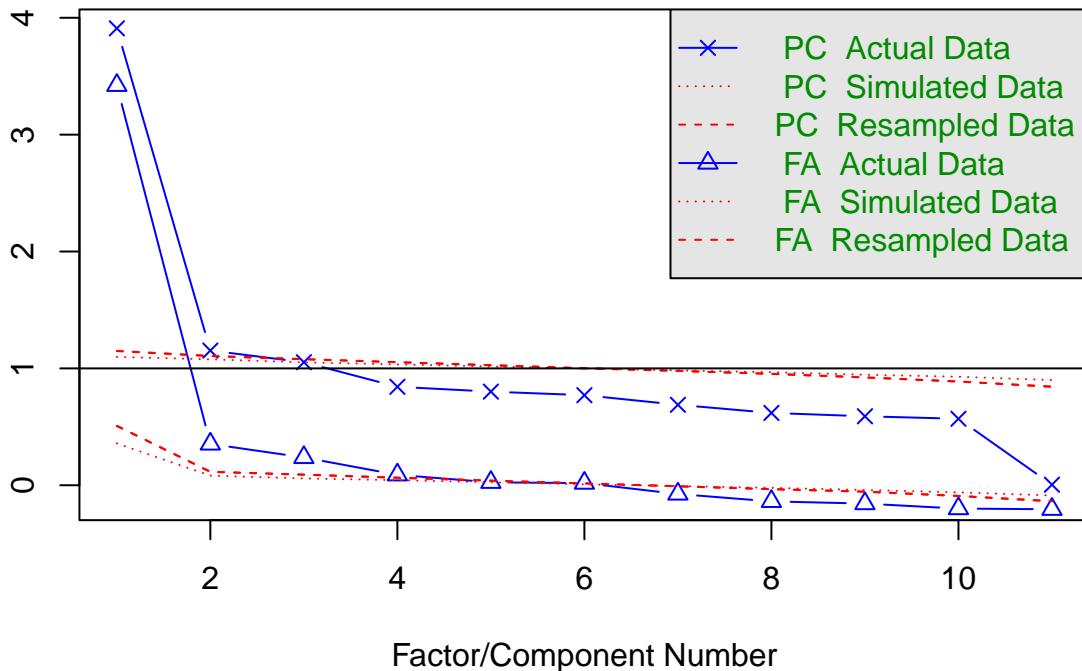
```

## Warning in fac(r = r, nfactors = nfactors, n.obs = n.obs, rotate = rotate, :
## ultra-Heywood case was detected. Examine the results carefully

```

eigenvalues of principal components and factor analysis

## Parallel Analysis Scree Plots



```
## Parallel analysis suggests that the number of factors = 4 and the number of components = 2
```

```
#Let's start with principal component analysis (PCA).
#The default function consists of one component.
#However, you can change this by using the argument.
#Please see all the arguments for PCA, using help function.
#In using PCA, the goal can be only data reduction,
#but the interpretation of components is frequently done
#in terms similar to those used when describing
#the latent variables estimated by FA.
#PCA reports the largest n eigen vectors
#rescaled by the square root of their eigen values.
principal(confdata)
```

```
## Principal Components Analysis
## Call: principal(r = confdata)
## Standardized loadings (pattern matrix) based upon correlation matrix
##          PC1    h2     u2 com
## confinan  0.62  0.38  0.6195   1
## conbus    0.58  0.34  0.6625   1
## conclerg  0.48  0.23  0.7696   1
## coneduc   0.57  0.32  0.6765   1
## compress  0.49  0.24  0.7644   1
## contv     0.54  0.29  0.7059   1
## conjudge  0.59  0.35  0.6518   1
```

```

## consci    0.37 0.14 0.8619   1
## conlegis  0.61 0.37 0.6289   1
## conarmy   0.51 0.26 0.7442   1
## con.scale 1.00 0.99 0.0053   1
##
##                               PC1
## SS loadings      3.91
## Proportion Var  0.36
##
## Mean item complexity =  1
## Test of the hypothesis that 1 component is sufficient.
##
## The root mean square of the residuals (RMSR) is  0.09
## with the empirical chi square  2403.31  with prob < 0
##
## Fit based upon off diagonal values = 0.92

## Retaining two factors instead of one. Check the results and identify the differences.
principal(confdata, nfactors = 2)

```

```

## Principal Components Analysis
## Call: principal(r = confdata, nfactors = 2)
## Standardized loadings (pattern matrix) based upon correlation matrix
##          RC1    RC2    h2    u2 com
## confinan  0.72  0.10 0.53 0.4740 1.0
## conbus    0.61  0.18 0.40 0.5984 1.2
## conclerg  0.71 -0.09 0.51 0.4924 1.0
## coneduc   0.44  0.36 0.32 0.6764 1.9
## compress  0.11  0.62 0.40 0.6008 1.1
## contv     0.29  0.50 0.33 0.6651 1.6
## conjudge  0.22  0.66 0.48 0.5222 1.2
## consci    -0.07 0.66 0.43 0.5667 1.0
## conlegis  0.47  0.38 0.37 0.6288 1.9
## conarmy   0.51  0.18 0.29 0.7083 1.2
## con.scale  0.75  0.66 1.00 0.0048 2.0
##
##          RC1    RC2
## SS loadings      2.77 2.29
## Proportion Var   0.25 0.21
## Cumulative Var  0.25 0.46
## Proportion Explained 0.55 0.45
## Cumulative Proportion 0.55 1.00
##
## Mean item complexity =  1.4
## Test of the hypothesis that 2 components are sufficient.
##
## The root mean square of the residuals (RMSR) is  0.1
## with the empirical chi square  2881.53  with prob < 0
##
## Fit based upon off diagonal values = 0.9

```

```

#The parallel factors technique compares
#the observed eigen values of a correlation matrix

```

```

#with those from random data.
#We could fit a one-factor model to these data
#in which all ten indicators are thought to reflect a common
#latent factor. Here, we estimate this factor using maximum likelihood.
conf1 <- fa(confdata, nfactors=1, fm="ml")
conf1

## Factor Analysis using method = ml
## Call: fa(r = confdata, nfactors = 1, fm = "ml")
## Standardized loadings (pattern matrix) based upon correlation matrix
##          ML1    h2    u2 com
## confinan 0.60 0.36 0.6399  1
## conbus   0.56 0.32 0.6814  1
## conclerg 0.49 0.24 0.7641  1
## coneduc  0.56 0.32 0.6843  1
## compress 0.49 0.24 0.7579  1
## contv    0.54 0.29 0.7063  1
## conjudge 0.59 0.35 0.6490  1
## consci   0.39 0.16 0.8442  1
## conlegis 0.59 0.35 0.6468  1
## conarmy  0.50 0.25 0.7487  1
## con.scale 1.00 1.00 0.0049  1
##
##          ML1
## SS loadings 3.87
## Proportion Var 0.35
##
## Mean item complexity = 1
## Test of the hypothesis that 1 factor is sufficient.
##
## df null model = 55 with the objective function = 6.38 with Chi Square = 18270.06
## df of the model are 44 and the objective function was 2.96
##
## The root mean square of the residuals (RMSR) is 0.09
## The df corrected root mean square of the residuals is 0.1
##
## The harmonic n.obs is 1885 with the empirical chi square 1535.68 with prob < 2.7e-293
## The total n.obs was 2867 with Likelihood Chi Square = 8454.79 with prob < 0
##
## Tucker Lewis Index of factoring reliability = 0.423
## RMSEA index = 0.258 and the 90 % confidence intervals are 0.254 0.263
## BIC = 8104.51
## Fit based upon off diagonal values = 0.92
## Measures of factor score adequacy
##          ML1
## Correlation of (regression) scores with factors 1.00
## Multiple R square of scores with factors       1.00
## Minimum correlation of possible factor scores 0.99

##We see that much of the variation in finance, business, judge,
##legislation, and education is explained by the latent factor.
##However, only 28% and 39% variance in science and press is explained
##with low communality and high uniquenesses. This suggests

```

```

##a poor factor solution. Let's compare against a 3-factor solution:
confactor <- fa(confdata, nfactors=3, fm="ml", rotate="oblimin")

## Loading required namespace: GPArotation

confactor

## Factor Analysis using method = ml
## Call: fa(r = confdata, nfactors = 3, rotate = "oblimin", fm = "ml")
## Standardized loadings (pattern matrix) based upon correlation matrix
##          ML1    ML3    ML2     h2     u2 com
## confinan  0.71 -0.10 -0.12  0.43  0.5713 1.1
## conbus    0.62 -0.08 -0.01  0.35  0.6531 1.0
## conclerg   0.59 -0.10 -0.13  0.30  0.7021 1.2
## coneduc    0.59  0.00 -0.04  0.33  0.6672 1.0
## compress   0.01  1.00 -0.01  1.00  0.0050 1.0
## contv      0.47  0.21 -0.06  0.32  0.6794 1.4
## conjudge   0.54  0.00  0.16  0.36  0.6373 1.2
## consci     0.01 -0.02  1.00  1.00  0.0050 1.0
## conlegis   0.63 -0.02 -0.04  0.38  0.6230 1.0
## conarmy    0.54 -0.09  0.03  0.28  0.7243 1.1
## con.scale  0.89  0.14  0.12  1.00  0.0049 1.1
##
##          ML1    ML3    ML2
## SS loadings   3.58  1.08  1.07
## Proportion Var 0.33  0.10  0.10
## Cumulative Var 0.33  0.42  0.52
## Proportion Explained 0.62  0.19  0.19
## Cumulative Proportion 0.62  0.81  1.00
##
## With factor correlations of
##          ML1    ML3    ML2
## ML1  1.00  0.37  0.28
## ML3  0.37  1.00  0.16
## ML2  0.28  0.16  1.00
##
## Mean item complexity = 1.1
## Test of the hypothesis that 3 factors are sufficient.
##
## df null model = 55 with the objective function = 6.38 with Chi Square = 18270.06
## df of the model are 25 and the objective function was 2.55
##
## The root mean square of the residuals (RMSR) is 0.07
## The df corrected root mean square of the residuals is 0.1
##
## The harmonic n.obs is 1885 with the empirical chi square 960.32 with prob < 1.5e-186
## The total n.obs was 2867 with Likelihood Chi Square = 7304.54 with prob < 0
##
## Tucker Lewis Index of factoring reliability = 0.12
## RMSEA index = 0.319 and the 90 % confidence intervals are 0.313 0.325
## BIC = 7105.51
## Fit based upon off diagonal values = 0.95
## Measures of factor score adequacy

```

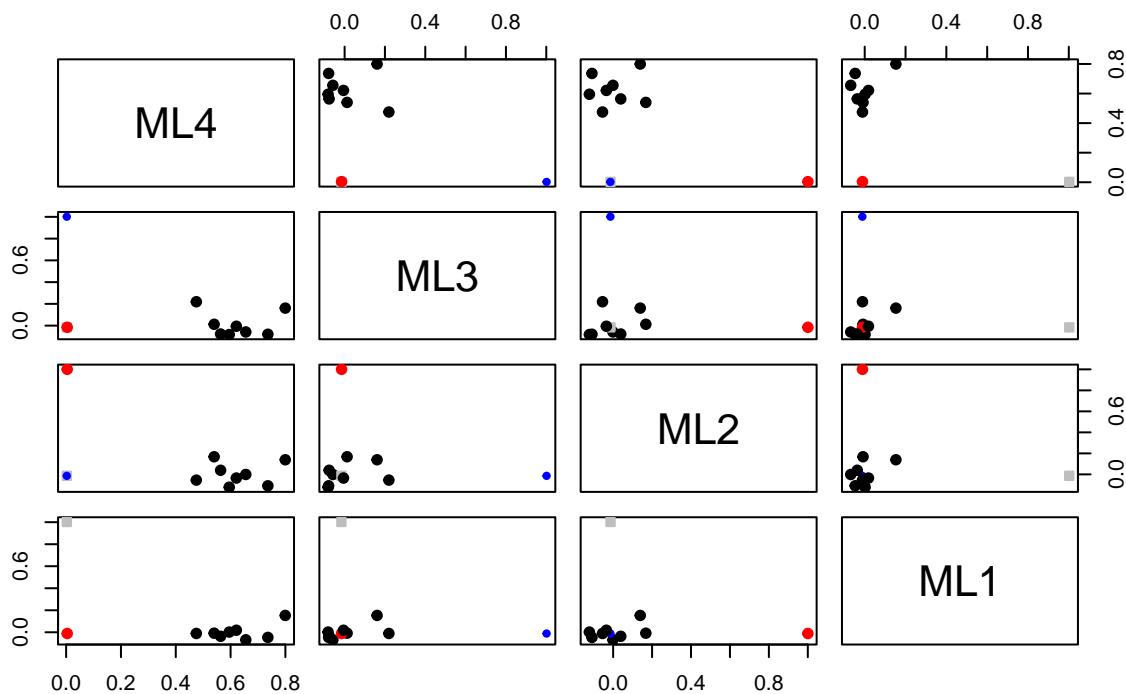
```

##                                     ML1  ML3  ML2
## Correlation of (regression) scores with factors   1.00 1.00 1.00
## Multiple R square of scores with factors        0.99 0.99 0.99
## Minimum correlation of possible factor scores  0.99 0.99 0.99

##The parallel analysis suggests a 4-factor solution.
confactor <- fa(confdata, nfactors=4, fm="ml", rotate="oblimin")
##fa.plot will plot the loading from a factor, principal components,
##or cluster analysis. If there are more than two factors, then a SPLOM
##of the loadings is generated.
fa.plot(confactor)

```

## Factor Analysis

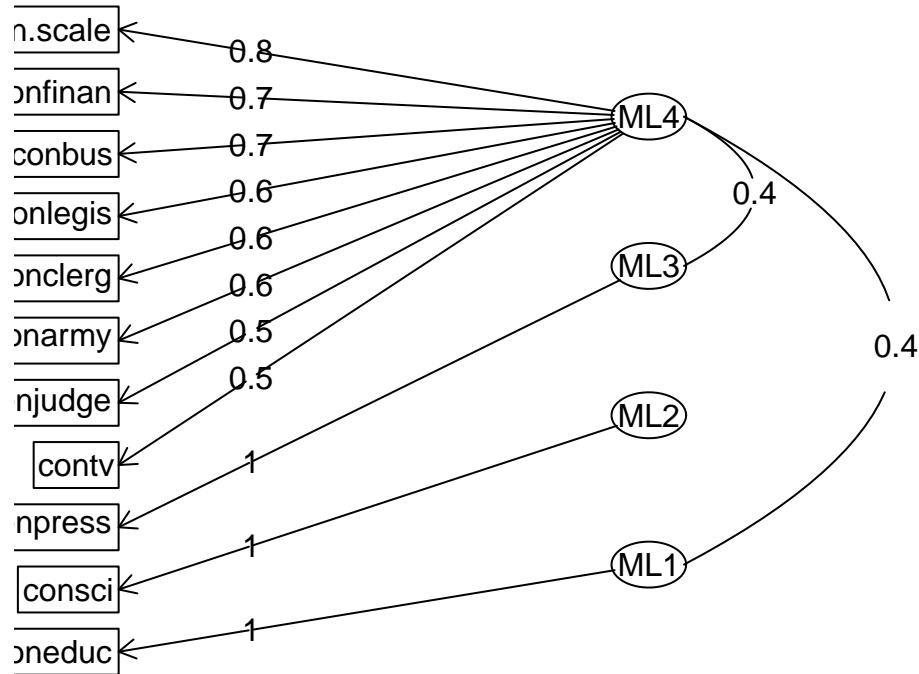


```

##fa.diagram replaces fa.graph and will draw a path diagram representing
##the factor structure. It does not require Rgraphviz and
##thus is probably preferred.
fa.diagram(confactor)

```

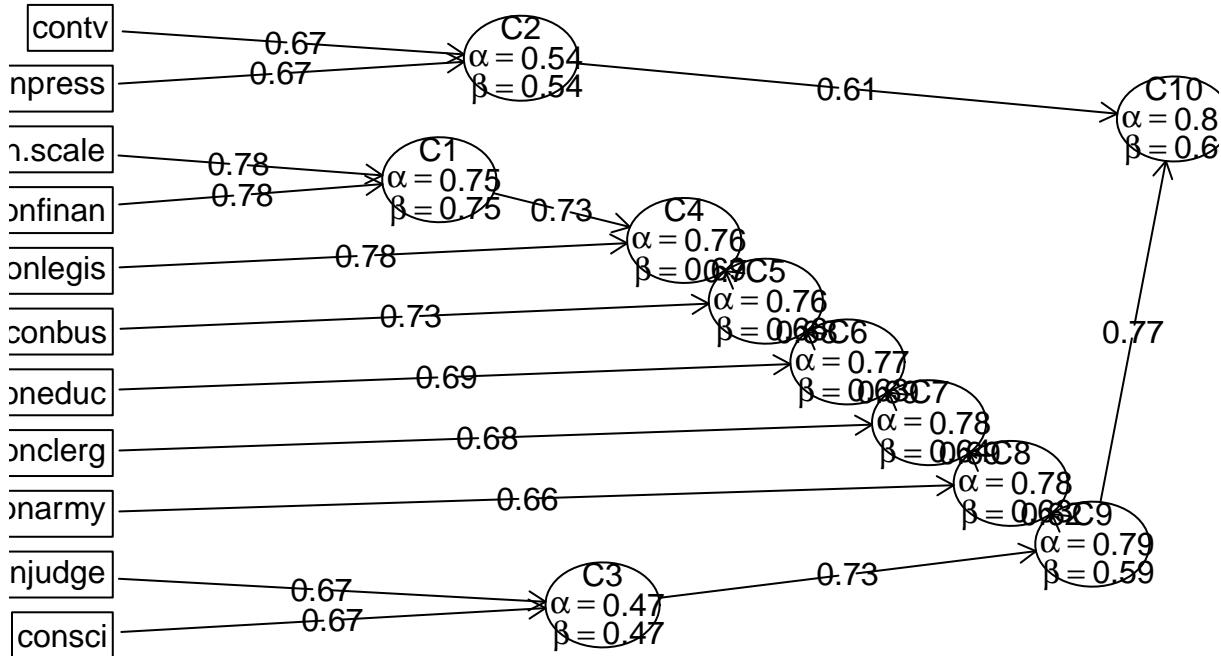
## Factor Analysis



```
#Item Cluster Analysis  
iclust(confdata)
```

```
## Warning in fa.stats(r = r, f = f, phi = phi, n.obs = n.obs, np.obs = np.obs, :  
## The estimated weights for the factor scores are probably incorrect. Try a  
## different factor score estimation method.
```

# ICLUST



```

## ICLUST (Item Cluster Analysis)
## Call: iclust(r.mat = confdata)
##
## Purified Alpha:
## C10
## 0.8
##
## G6* reliability:
## C10
## 1
##
## Original Beta:
## C10
## 0.64
##
## Cluster size:
## C10
## 11
##
## Item by Cluster Structure matrix:
##          0   P   C10
## confinan C10 C10 0.59
## conbus    C10 C10 0.56
## conclerg  C10 C10 0.47
## coneduc   C10 C10 0.55
## conpress  C10 C10 0.48

```

```
## contv      C10 C10 0.53
## conjudge   C10 C10 0.58
## consci     C10 C10 0.39
## conlegis   C10 C10 0.58
## conarmy    C10 C10 0.50
## con.scale  C10 C10 1.02
##
## With Sums of squares of:
## C10
## 3.8
##
## Purified scale intercorrelations
## reliabilities on diagonal
## correlations corrected for attenuation above diagonal:
##      C10
## C10 0.8
##
## Cluster fit =  0.72  Pattern fit =  0.97  RMSR =  0.08
```