Structural Equation Modeling

P.09 - Statistical Modeling of Panel Data

November 15, 2022 (13:43:19)

Lab Description

For this practical you will need the following packages: lavaan and semPlot. You can install and load these packages using the following code:

```
# Install packages.
install.packages(c("lavaan", "semPlot"))

# Load the packages.
library(lavaan)
library(semPlot)
```

Exercise 1

In this exercise you are going to investigate whether a repeated measurement conforms to the *simplex* or *quasi-simplex* correlation structure. Consider the dataset health.dat, which can be found in the folder for the current practical on Canvas. These data are derived from a national health survey with interviews of individuals aged 50 years and above conducted biannually. You are going to analyze the self-rated health question about overall health collected over six waves. Ratings from this question were from 1 (*poor*) to 5 (*excellent*). The repeated measurement variables are: srh1, srh2, srh3, srh4, srh5, srh6. To get you started, you can use the following code to load the data and set the variable names.

Set the working directory to the location where your data file has been downloaded and load the data.

```
# For example.
setwd("/Users/mihai/Downloads")

# Load data.
data_ex_1 <- read.table("health.dat")

# Inspect the data.
View(data_ex_1)</pre>
```

Set the variable names.

```
# Variable names.
variable_ex_1_names = c(
    "age", "srh1", "srh2", "srh3", "srh4", "srh5", "srh6", "bmi1",
    "bmi2", "bmi3", "bmi4", "bmi5", "bmi6", "cesdna1", "cesdpa1", "cesdso1",
    "cesdna2", "cesdpa2", "cesdso2", "cesdna3", "cesdpa3", "cesdso3",
    "cesdna4", "cesdpa4", "cesdso4", "cesdna5", "cesdpa5", "cesdso5",
    "cesdna6", "cesdpa6", "cesdso6", "diab1", "diab2", "diab3 ", "diab4", "diab5", "diab6"
)

# Set the names.
names(data_ex_1) <- variable_ex_1_names

# List variables.
str(data_ex_1)</pre>
```

- a. Estimate the perfect simplex model for these six repeated measurement. Request a standardized solution, and evaluate the fit of this model.
- b. Inspect the correlation of the first time (t_1) point measurement with later time points, and the standardized auto-regression coefficients. Does the pattern of these coefficients provide evidence that the perfect simplex model holds?
- c. Now estimate the quasi-simplex model. For identification purposes, set the measurement residual variances of the first and last measurement to 0. Again, request a standardized solution and evaluate the fit of this model.
- d. Perform a likelihood ratio test of the perfect simplex model against the quasi-simplex model. Interpret the result of this test.
- e. Obtain the estimated correlations among the latent variables, using the lavInspect() function. Note that you can check the documentation for this function by running ?lavInspect in R.
- f. Inspect the correlation of the second time point (t_2) measurement with later time points, and the standardized auto-regression coefficients. Does the pattern of these coefficients provide evidence that the data conform to the simplex structure, if measurement error is taken into account for the measurement at t_2 to t_5 ?

Exercise 2

Consider the Cross-Lagged Panel Model depicted in Figure 1.

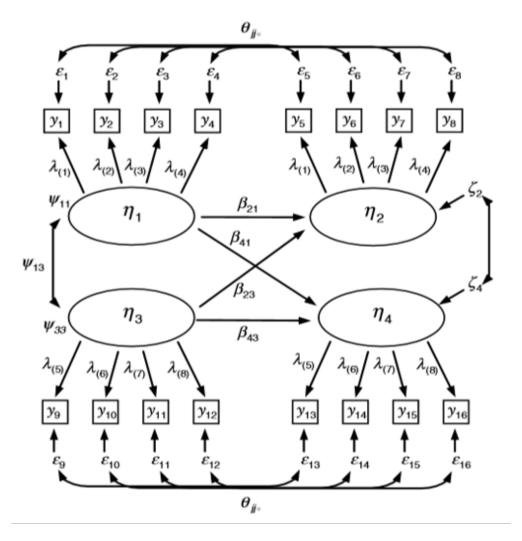


Figure 1: Example of Cross-Lagged Panel Model with four latent variables.

Using the socex1.dat data, estimate a similar model to investigate the bidirectional effects between positive affect (PA) and unwanted advice (UA). The specifications of the model are as follows:

- W1POSAFF (latent variable PA at t_1) with indicators: w1happy, w1enjoy, w1satis, w1joyful, and w1please
- W2POSAFF (latent variable PA at t_2) with indicators: w2happy, w2enjoy, w2satis, w2joyful, and w2please
- W1UNW (latent variable UA at t_1) with indicators: w1unw1, w1unw2, and w1unw3
- W2UNW (latent variable UA at t_2) with indicators: w2unw1, w2unw2, and w2unw3

In addition, in this model you should:

• Do not use the marker method to set the scale of the latent variables, but free the marker variables and

set exogenous factor variance for first occasion latent variables for identification purpose to 1.

- Impose equality constraints for the loadings between the different measurement moments for repeated indicators.
- Estimate the cross-lagged and auto-regressive effects.
- Include correlated measurement residuals.

To get you started, you can use the following R code to load the data and set the variable names.

```
# Load data.
data_ex_2 <- read.table("socex1.dat")

# Inspect the data.
View(data_ex_2)

# Variable names.
variable_ex_2_names <- c(
    "wivst1", "wivst2", "wivst3", "w2vst1", "w2vst2",
    "w2vst5", "w3vst1", "x3vst2", "w3vst3", "wiunv1", "wiunv2", "wiunw3",
    "w2unw1", "w2unw2", "w2unw3", "w3unw1", "w3unw2", "w3unw3", "widboth",
    "widsad", "widblues", "widdep", "w2dboth", "w2dsad", "w2ddep",
    "w3dsoth", "w3dsad", "w3dblues", "w3ddep", "wimarr2", "w1happy", "w1enjoy",
    "w1satis", "w1joyful", "w1please", "w2happy", "w2enjoy", "w2satis", "w2joyful",
    "w2please", "w3happy", "w3enjoy", "w3satis", "w3joyful", "w3please", "w1lea",
    "w2lea", "w3lea"
)

# Set the names.
names(data_ex_2) <- variable_ex_2_names

# List variables.
str(data_ex_2)</pre>
```

Answer the following:

- a. Estimate the parameters of this model (i.e., including standardized parameters) and evaluate the model fit.
- b. What is the size of the auto-regressive and cross-lagged standardized effects in this model?

Exercise 3

Using the dataset health.dat:

a. Estimate for the six repeated measurements the trait-state-error model, as proposed by Kenny & Zautra (1995). You can use as starting point the model depicted in *Figure 2*, but note that it should be extended to include y1 to y6.

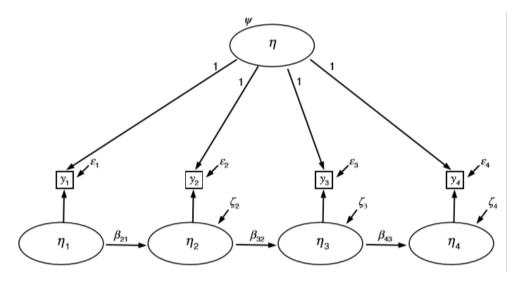


Figure 2: Example of repeated measures model.

To construct a single observed measurement per wave, calculate the average score of the three indicators per wave, as follows:

```
# Assuming you data for this exercise is called `data_ex_3`, add the average
# scores per wave as follows.
data_ex_3$cesd1 = (cesdna1 + cesdpa1 + cesdso1) / 3
data_ex_3$cesd2 = (cesdna2 + cesdpa2 + cesdso2) / 3
data_ex_3$cesd3 = (cesdna3 + cesdpa3 + cesdso3) / 3
data_ex_3$cesd4 = (cesdna4 + cesdpa4 + cesdso4) / 3
data_ex_3$cesd5 = (cesdna5 + cesdpa5 + cesdso5) / 3
data_ex_3$cesd6 = (cesdna6 + cesdpa6 + cesdso6) / 3
```

The model follows that shown in $Figure\ 2$ and, specifies a single trait factor with the loading for each indicator set equal to 1 and a latent state factor with a single loading set equal to 1 at each occasion. Each state factor is regressed on the state factor from the prior time point. Several longitudinal equality constraints should be imposed (i.e., stationarity):

- all auto-regressive coefficients are set equal
- all state factor residuals except for the first state factor are set equal
- all measurement residuals are set equal

The trait variance and the state factor variance at t_1 are free to be estimated, and η and η_1 are assumed independent.

- b. What is the fit of this model?
- c. Verify the calculations reported by Newsom for this model:
 - proportion of trait variance equal to .43
 - proportion of state variance equal to .10
 - proportion of error variance equal to .47

Note. The latent trait-state-error model is equivalent to the univariate STARTS model discussed during the lecture. In a nutshell, the STARTS model consists of:

- stable trait (i.e., the "trait" in the trait-state-error terminology)
- auto-regressive trait (i.e., the "state" in the trait-state-error terminology)
- state (i.e., the "error" in the *trait-state-error* terminology), which, in the univariate *STARTS* model, also contains error as mentioned in the lecture