

Example of a MASEM-analysis using the OC approach

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2017

Contents

Installing the metaSEM package	1
Loading metaSEM and the Stage1.OC() function	1
Loading the data	1
Fit the Stage 1 model	2
Fit the Stage 2 model	4
References	6

Installing the metaSEM package

- R can be downloaded at <http://www.r-project.org/>.
- We only need to install the metaSEM-package once.

```
install.packages("metaSEM")
```

Loading metaSEM and the Stage1.OC() function

```
library(metaSEM)
```

```
## Warning: package 'metaSEM' was built under R version 3.3.3
```

```
## Warning: package 'OpenMx' was built under R version 3.3.3
```

```
# read the Stage1.OC() function from an URL  
source("http://www.suzannejak.nl/Stage1.OC.R")
```

Loading the data

- The datafile should be saved in the working directory, it can be downloaded from <http://www.suzannejak.nl/exampledata.R>

```
load("exampledata.R")
```

```
data <- exampledat$data  
n <- exampledat$n
```

```
## show part of the data  
## NA's represent missing correlations  
head(data)
```

```
## $`1`  
##          v1          v2          v3          v4 v5  
## v1 1.0000000          NA 0.1337607 0.3790011 NA  
## v2          NA 1.0000000 0.3176261 0.5343438 NA  
## v3 0.1337607 0.3176261 1.0000000 0.2490913 NA
```

```

## v4 0.3790011 0.5343438 0.2490913 1.0000000 NA
## v5      NA      NA      NA      NA NA NA
##
## $`2`
##      v1      v2      v3      v4 v5
## v1 1.0000000 0.4277664 0.2104662 0.4189287 NA
## v2 0.4277664 1.0000000 0.2254605 0.5791790 NA
## v3 0.2104662 0.2254605 1.0000000 0.2434884 NA
## v4 0.4189287 0.5791790 0.2434884 1.0000000 NA
## v5      NA      NA      NA      NA NA
##
## $`3`
##      v1      v2      v3      v4      v5
## v1 1.0000000      NA 0.12120473 0.3869396 0.16440272
## v2      NA 1.0000000 0.28942770 0.5320153 0.22664944
## v3 0.1212047 0.2894277 1.00000000 0.2968476 0.09832017
## v4 0.3869396 0.5320153 0.29684761 1.0000000      NA
## v5 0.1644027 0.2266494 0.09832017      NA 1.00000000
##
## $`4`
##      v1      v2      v3      v4      v5
## v1 1.0000000      NA 0.1762705 0.3357207 0.1899715
## v2      NA 1.0000000 0.3682914 0.5684330 0.2579966
## v3 0.1762705 0.3682914 1.00000000 0.3413604 0.2251231
## v4 0.3357207 0.5684330 0.3413604 1.0000000 0.4965113
## v5 0.1899715 0.2579966 0.2251231 0.4965113 1.0000000
##
## $`5`
##      v1      v2      v3      v4 v5
## v1 1.0000000 0.4237114 0.1475889 0.3858164 NA
## v2 0.4237114 1.0000000 0.1648990 0.5951703 NA
## v3 0.1475889 0.1648990 1.00000000 0.2303143 NA
## v4 0.3858164 0.5951703 0.2303143 1.0000000 NA
## v5      NA      NA      NA      NA NA
##
## $`6`
##      v1      v2      v3      v4 v5
## v1 1.0000000      NA 0.1931003 0.3732731 NA
## v2      NA 1.0000000 0.1878165 0.4958514 NA
## v3 0.1931003 0.1878165 1.00000000 0.3248142 NA
## v4 0.3732731 0.4958514 0.3248142 1.0000000 NA
## v5      NA      NA      NA      NA NA

```

Fit the Stage 1 model

```
stage1.fit <- Stage1.OC(data,n)
```

- The parameters named 'varname_varname' (for example 'v1_v2') without an additional subscript are the estimated pooled correlations

```
stage1.fit$summary
```

```
## Summary of Stage1
##
```

```

## free parameters:
##      name      matrix row col Estimate Std.Error A
## 1      v2_v1_1 Study1.R  1  2 0.4082216 0.05677017
## 2      v3_v1 Study1.R  1  3 0.1723269 0.01866181
## 3      v3_v2 Study1.R  2  3 0.2500300 0.01803853
## 4      v4_v1 Study1.R  1  4 0.3844283 0.01677918
## 5      v4_v2 Study1.R  2  4 0.5612909 0.01347725
## 6      v4_v3 Study1.R  3  4 0.2733916 0.01809644
## 7 Study1.D[1,1] Study1.D  1  1 1.0003962 0.05404203
## 8 Study1.D[2,2] Study1.D  2  2 1.0033897 0.05150474
## 9 Study1.D[3,3] Study1.D  3  3 0.9961179 0.05461772
## 10 Study1.D[4,4] Study1.D  4  4 1.0079903 0.05136543
## 11      v2_v1 Study2.R  1  2 0.3961914 0.02066434
## 12 Study2.D[1,1] Study2.D  1  1 0.9876256 0.03758887
## 13 Study2.D[2,2] Study2.D  2  2 0.9889389 0.03593669
## 14 Study2.D[3,3] Study2.D  3  3 1.0014348 0.03946251
## 15 Study2.D[4,4] Study2.D  4  4 0.9899748 0.03596041
## 16      v2_v1_3 Study3.R  1  2 0.4085775 0.03422400
## 17      v5_v1 Study3.R  1  5 0.1923540 0.02384504
## 18      v5_v2 Study3.R  2  5 0.2570609 0.02319053
## 19      v5_v3 Study3.R  3  5 0.1610699 0.02400087
## 20      v5_v4_3 Study3.R  4  5 0.4798270 0.03186567
## 21 Study3.D[1,1] Study3.D  1  1 1.0022034 0.03225473
## 22 Study3.D[2,2] Study3.D  2  2 1.0075205 0.03105862
## 23 Study3.D[3,3] Study3.D  3  3 0.9973941 0.03248184
## 24 Study3.D[4,4] Study3.D  4  4 1.0106213 0.03075806
## 25 Study3.D[5,5] Study3.D  5  5 1.0061537 0.03306331
## 26      v2_v1_4 Study4.R  1  2 0.4200181 0.05127224
## 27      v5_v4 Study4.R  4  5 0.4614729 0.03055451
## 28 Study4.D[1,1] Study4.D  1  1 1.0048568 0.04935553
## 29 Study4.D[2,2] Study4.D  2  2 0.9900994 0.04598144
## 30 Study4.D[3,3] Study4.D  3  3 0.9817736 0.04843889
## 31 Study4.D[4,4] Study4.D  4  4 0.9862023 0.04370591
## 32 Study4.D[5,5] Study4.D  5  5 0.9858476 0.04769470
## 33 Study5.D[1,1] Study5.D  1  1 0.9944915 0.03173957
## 34 Study5.D[2,2] Study5.D  2  2 0.9892372 0.03014256
## 35 Study5.D[3,3] Study5.D  3  3 1.0090789 0.03332190
## 36 Study5.D[4,4] Study5.D  4  4 0.9910450 0.03018272
## 37      v2_v1_6 Study6.R  1  2 0.4173744 0.04101664
## 38 Study6.D[1,1] Study6.D  1  1 1.0042399 0.04006509
## 39 Study6.D[2,2] Study6.D  2  2 1.0252479 0.03925901
## 40 Study6.D[3,3] Study6.D  3  3 0.9979039 0.04023611
## 41 Study6.D[4,4] Study6.D  4  4 1.0176855 0.03851655
## 42      v5_v4_7 Study7.R  4  5 0.4689783 0.05024808
## 43 Study7.D[1,1] Study7.D  1  1 1.0215404 0.05340393
## 44 Study7.D[2,2] Study7.D  2  2 1.0056254 0.04957512
## 45 Study7.D[3,3] Study7.D  3  3 1.0220451 0.05497446
## 46 Study7.D[4,4] Study7.D  4  4 1.0227939 0.05026655
## 47 Study7.D[5,5] Study7.D  5  5 1.0040288 0.05367500
## 48 Study8.D[1,1] Study8.D  1  1 1.0000747 0.03697773
## 49 Study8.D[2,2] Study8.D  2  2 1.0043971 0.03550675
## 50 Study8.D[3,3] Study8.D  3  3 0.9959951 0.03762448
## 51 Study8.D[4,4] Study8.D  4  4 0.9992211 0.03451656
## 52 Study8.D[5,5] Study8.D  5  5 1.0024216 0.03775416

```

```

## 53      v5_v4_9  Study9.R  4  5 0.4469941 0.04817291
## 54 Study9.D[1,1] Study9.D  1  1 0.9877207 0.04649311
## 55 Study9.D[2,2] Study9.D  2  2 0.9808660 0.04375659
## 56 Study9.D[3,3] Study9.D  3  3 0.9848725 0.04759421
## 57 Study9.D[4,4] Study9.D  4  4 0.9795266 0.04310023
## 58 Study9.D[5,5] Study9.D  5  5 0.9896828 0.04807775
## 59      v2_v1_10 Study10.R  1  2 0.3708073 0.06741842
## 60 Study10.D[1,1] Study10.D  1  1 0.9831932 0.05474844
## 61 Study10.D[2,2] Study10.D  2  2 0.9748127 0.05317144
## 62 Study10.D[3,3] Study10.D  3  3 0.9835970 0.05459663
## 63 Study10.D[4,4] Study10.D  4  4 0.9819895 0.05437257
##
## Model Statistics:
##           | Parameters | Degrees of Freedom | Fit (-2lnL units)
## Model:           63                62                10026.812
## Saturated:       125                0                 9963.677
## Independence:    NA                 NA                 NA
## Number of observations/statistics: 2718/125
##
## chi-square: <U+03C7>2 ( df=62 ) = 63.13464, p = 0.4360062
## Information Criteria:
##           | df Penalty | Parameters Penalty | Sample-Size Adjusted
## AIC:      -60.86536                189.1346                NA
## BIC:      -427.13976                561.3167                361.1456
## CFI: NA
## TLI: NA (also known as NNFI)
## RMSEA: 0.002594831 [95% CI (0, 0.01313995)]
## Prob(RMSEA <= 0.05): 1
## To get additional fit indices, see help(mxRefModels)
## timestamp: 2017-10-10 11:52:50
## Wall clock time (HH:MM:SS.hh): 00:00:00.54
## optimizer: SLSQP
## OpenMx version number: 2.7.12
## Need help? See help(mxSummary)

```

Fit the Stage 2 model

- Create a matrix with direct effects (A) and a matrix with variances and covariances (S)
- For instruction regarding model specification see Jak (2015) or Cheung (2014).

```

nvar <- nrow(data[[1]])
varnames <- rownames(data[[1]])

Amatrix <- create.mxMatrix(
  c( 0,0,0,0,0,
      0,0,0,0,0,
      0,0,0,0,0,
      "0.2*b41", "0.5*b42", "0.1*b43", 0,0,
      0,0,0, "0.4*b54", 0),
  type = "Full",
  nrow = nvar,
  ncol = nvar,
  byrow = TRUE,

```

```

name = "A",
dimnames = list(varnames,varnames))

Smatrix <- create.mxMatrix(
  c("1",
    ".2*p21","1",
    ".2*p31",".2*p32","1",
    0,0,0,"1*p44",
    0,0,0,0,"1*p55"),
  type="Symm",
  byrow = TRUE,
  name="S",
  dimnames = list(varnames,varnames))

```

- Fitting the Stage 2 model on the pooled correlation matrix from the random effects Stage 1 analysis

```

Stage2.fit <- wls(Cov = stage1.fit$pooledR, asyCov = stage1.fit$acovR,
  n = sum(n), Amatrix = Amatrix, Smatrix = Smatrix)

```

```
summary(Stage2.fit)
```

```

##
## Call:
## wls(Cov = stage1.fit$pooledR, asyCov = stage1.fit$acovR, n = sum(n),
##     Amatrix = Amatrix, Smatrix = Smatrix)
##
## 95% confidence intervals: z statistic approximation
## Coefficients:
##      Estimate Std. Error  lbound  ubound z value Pr(>|z|)
## b41 0.181407  0.018181 0.145773 0.217042  9.9777 < 2.2e-16 ***
## b42 0.457195  0.016009 0.425818 0.488572 28.5584 < 2.2e-16 ***
## b43 0.128694  0.016314 0.096719 0.160669  7.8884 3.109e-15 ***
## b54 0.469241  0.027082 0.416161 0.522321 17.3265 < 2.2e-16 ***
## p21 0.396448  0.020666 0.355943 0.436952 19.1836 < 2.2e-16 ***
## p31 0.172007  0.018668 0.135418 0.208596  9.2139 < 2.2e-16 ***
## p32 0.250629  0.018047 0.215257 0.286001 13.8874 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Goodness-of-fit indices:
##
##                               Value
## Sample size                   2718.0000
## Chi-square of target model     2.8342
## DF of target model             3.0000
## p value of target model        0.4179
## Number of constraints imposed on "Smatrix" 0.0000
## DF manually adjusted           0.0000
## Chi-square of independence model 2135.4534
## DF of independence model      10.0000
## RMSEA                          0.0000
## RMSEA lower 95% CI             0.0000
## RMSEA upper 95% CI             0.0317
## SRMR                           0.0113

```

```
## TLI                1.0003
## CFI                1.0000
## AIC                -3.1658
## BIC                -20.8888
## OpenMx status1: 0 ("0" or "1": The optimization is considered fine.
## Other values indicate problems.)
```

References

- Jak, S. (2015). Meta-analytic structural equation modeling. Cham: Springer International Publishing
- Cheung, M.W.-L. (2014). MetaSEM: an R package for meta-analysis using structural equation modeling. *Frontiers in Psychology*, 5 (1521)