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Introduction to Meta-Analyses of Structural Equation Models with {metaSEM}

Bringing together the best of two worlds
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Use  slo



MASEM Concepts

Meta-Analysis

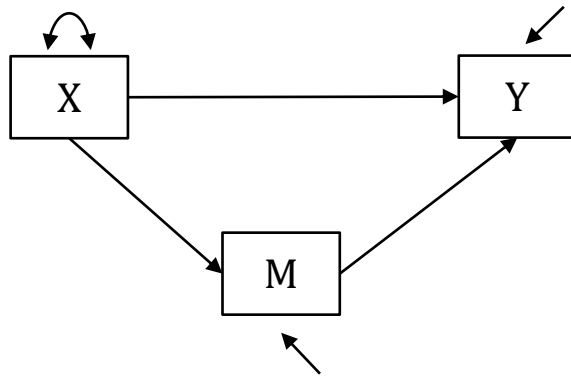
Interest in the average effects

Interest in differences between studies

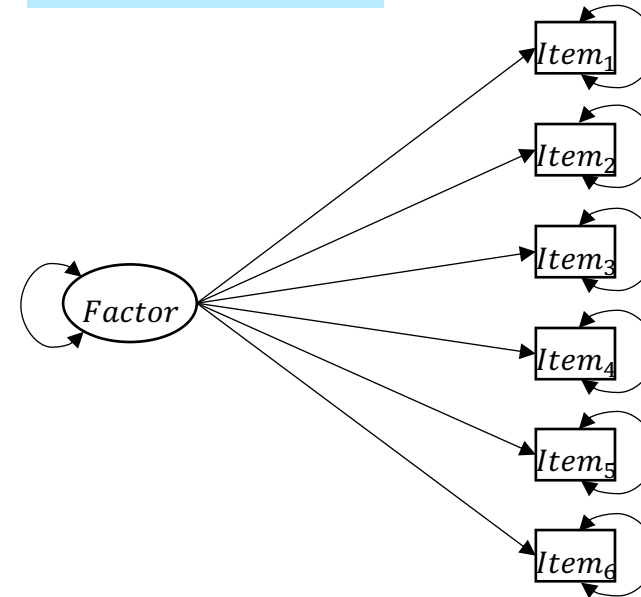
MASEM Concepts

Structural Equation Modeling

Path Model



Factor model



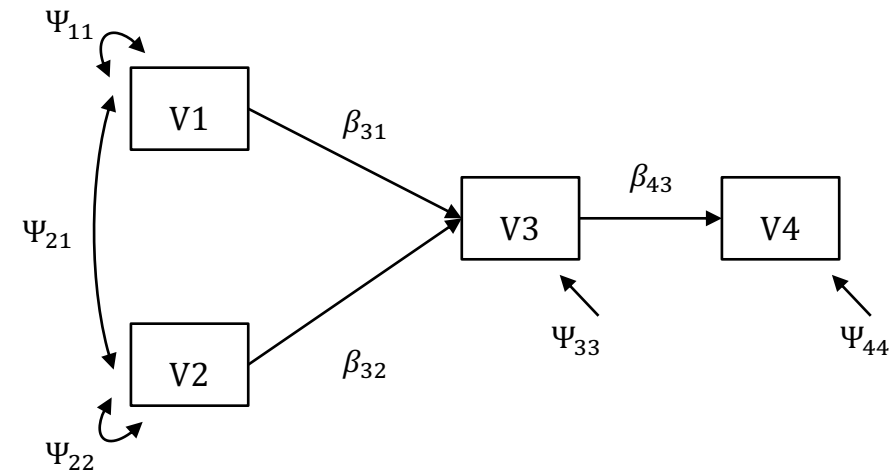
No raw data needed

MASEM Concepts

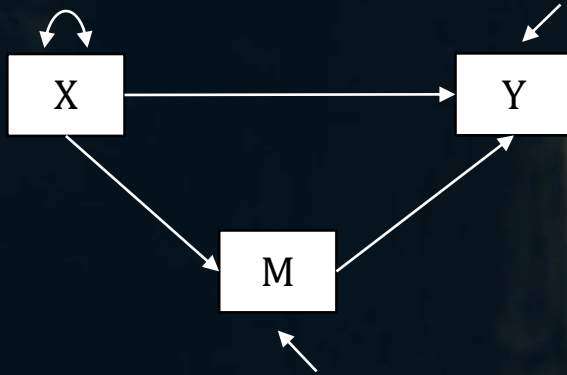
Structural Equation Modeling

- The covariance matrix between the observed variables is modeled as a function of SEM parameters.

V1	V2	V3	V4
1.00			
ρ_{21}	1.00		
ρ_{31}	ρ_{32}	1.00	
ρ_{41}	ρ_{42}	ρ_{43}	1.00



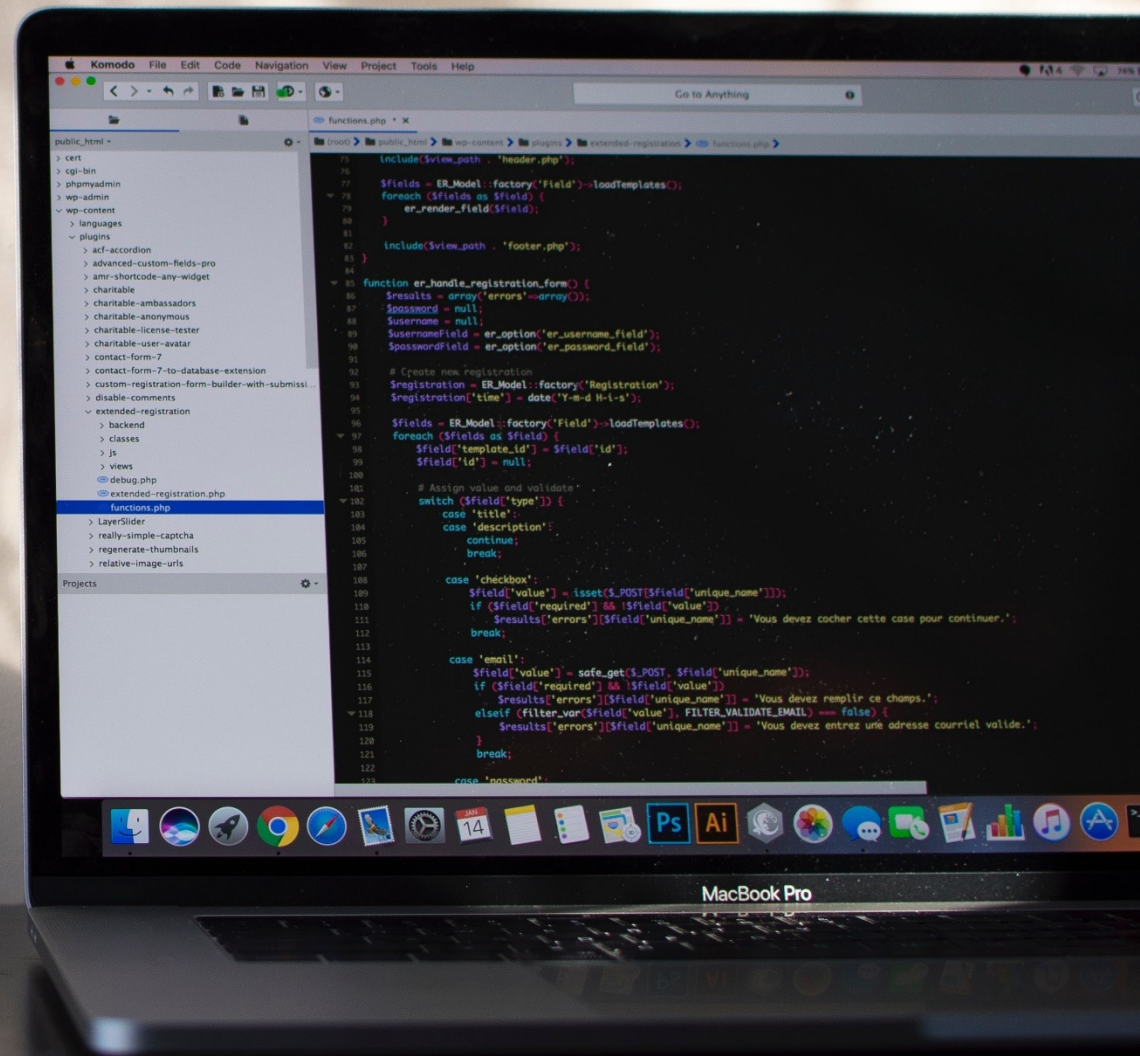
Why MASEM?



Why combining meta-analysis and SEM?

What SEM adds to meta-analysis?

MASEM Example

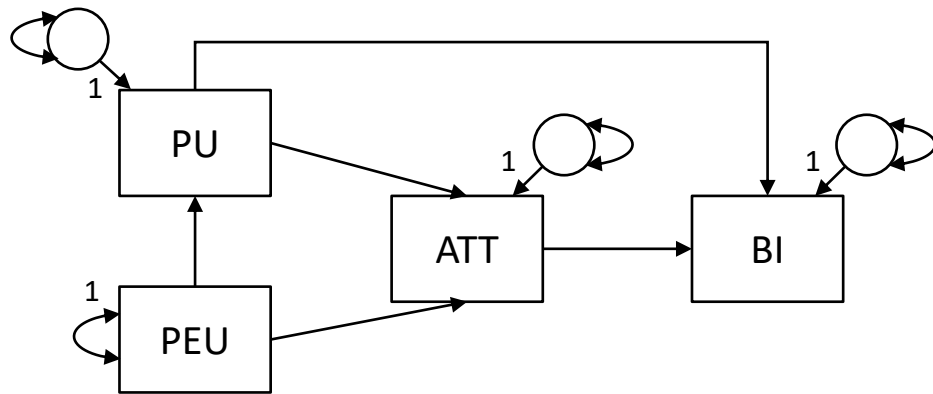


Technology Acceptance Model (TAM)

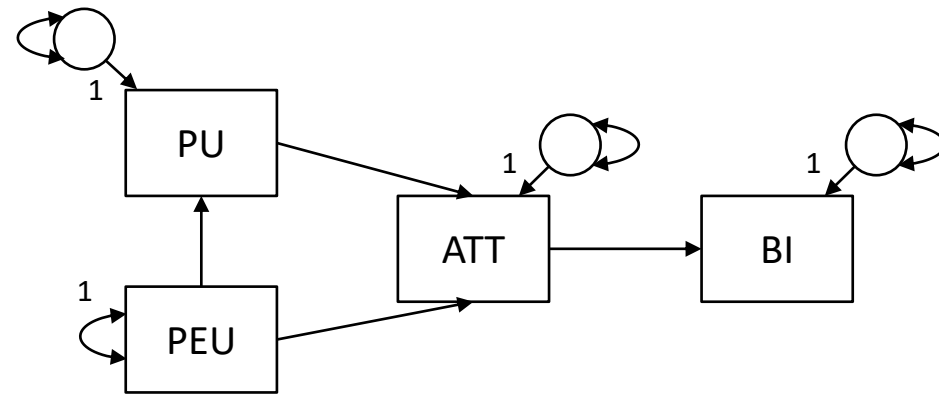
A Model Explaining Teachers' Intentions to Use Technology

Variables

- BI: Intentions to use technology, ATT: Attitudes toward technology
- PEU: Perceived ease of use, PU: Perceived usefulness



Model 1



Model 2

Technology Acceptance Model (TAM)

Meta-analytic data set based on teacher samples

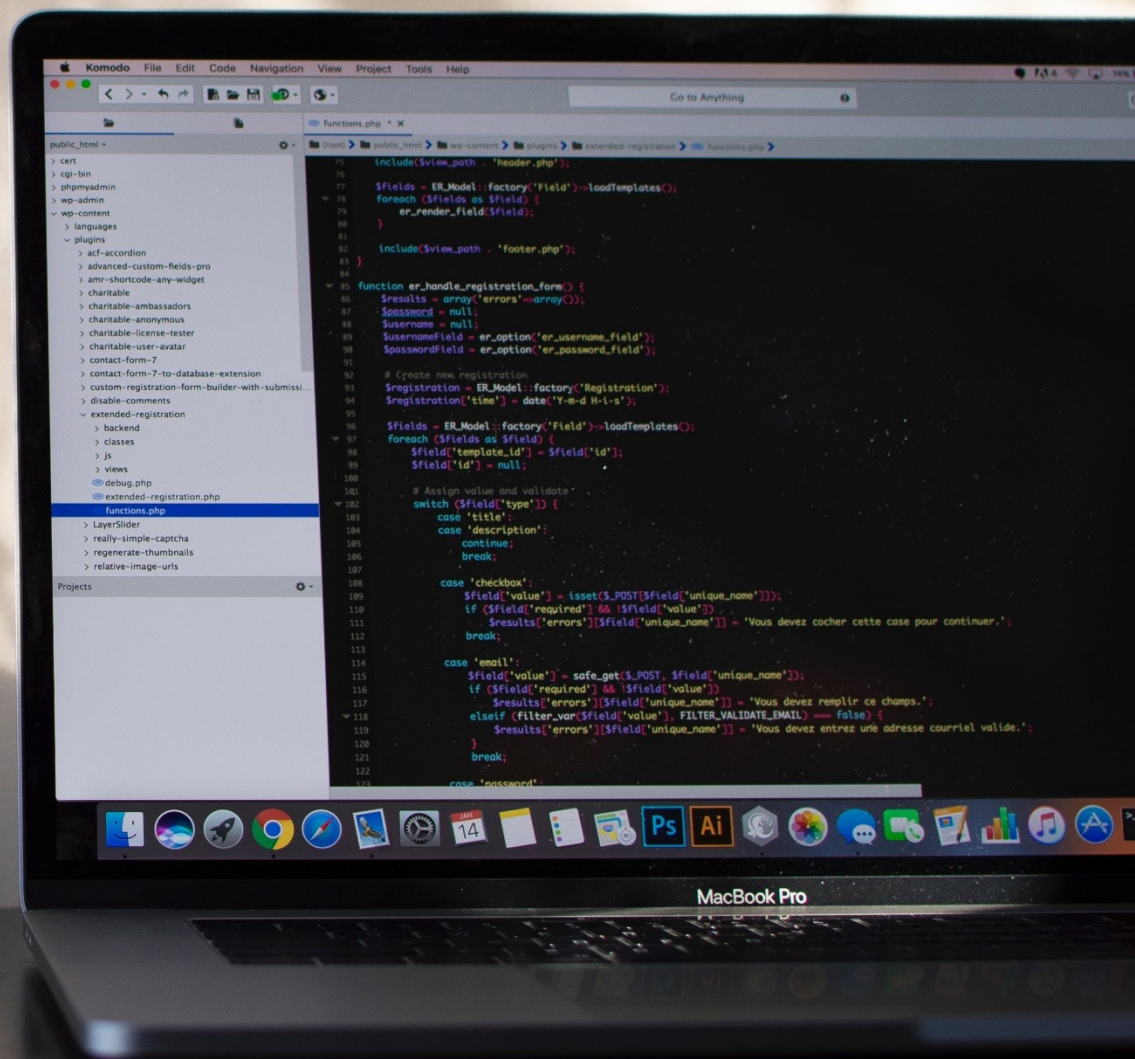
Data set

- 50 studies with independent samples
- 4 variables
- 300 correlations
- 50 correlation matrices
- Pre- and in-service teacher samples

Main RQs

- 1. Model evaluation:** To what extent does the TAM represent the teacher data across samples?
- 2. Model comparison:** To what extent does a direct effect exist?

R



Reading the Data

PU	PEU	ATT	BI
1.00			
ρ_{21}	1.00		
ρ_{31}	ρ_{32}	1.00	
ρ_{41}	ρ_{42}	ρ_{43}	1.00

	A	X	Y	Z	AA	AB	AC	AD	AE	AF	AG	AH
1	StudyID	Sample size	PU-PU	PU-PEOU	PEOU	PU-ATT	PEOU-ATT	ATT-ATT	PU-BI	PEOU-BI	ATT-BI	BI-BI
2	217	446	1	0,61	1	0,6776	0,5138	1	0,599184	0,449892	0,860328	1
3	247	88	1	0,661	1	0,59	0,584	1	0,75	0,746	0,684	1
4	281	271	1	0,89	1	0,423	0,187	1	0,374	0,82	0,512	1
5	289	234	1	0,5	1	0,64	0,55	1	0,55	0,48	0,59	1
6	311	29	1	0,512	1	0,461	0,754	1	0,364	0,669	0,595	1
7	320	313	1	0,279	1	0,63	0,873	1	0,319	0,441	0,505	1
8	332A	169	1	0,68	1	0,68	0,73	1	0,34	0,3	0,3	1
9	332B	170	1	0,4	1	0,56	0,43	1	0,23	0,32	0,36	1
10	335A	151	1	0,329	1	0,565	0,349	1	0,523	0,327	0,462	1

1. A list of correlation/covariance matrices.

2. A vector of sample size.

Understanding and Exploring your Data

Test for positive definiteness.

```
# PD check
pd_check <- is.pd(TAMu$data)
sum(!is.na(pd_check))
```

```
## [1] 50
```

```
pd_check
```

```
## Okazaki & Renda dos Santos (2012)
##                                     TRUE
##                                     Wong (2015)
##                                     TRUE
```

Explore your data.

```
# Some description of the studies
# Available studies per correlation
pattern.na(TAMu$data, show.na=FALSE)
```
```

|     | PU | PEU | ATT | BI |
|-----|----|-----|-----|----|
| PU  | 50 | 50  | 50  | 50 |
| PEU | 50 | 50  | 50  | 50 |
| ATT | 50 | 50  | 50  | 50 |
| BI  | 50 | 50  | 50  | 50 |

```
Available sample sizes per correlation
pattern.n(TAMu$data, TAMu$n)
```
```

	PU	PEU	ATT	BI
PU	14918	14918	14918	14918
PEU	14918	14918	14918	14918
ATT	14918	14918	14918	14918
BI	14918	14918	14918	14918

Stage 1: Pooling Correlation Matrices

Pooling correlation matrices

- REM
- Diagonal constraints of the heterogeneity matrix

```
# Combine correlation matrices using a REM  
#  
rem <- tssem1(TAMu$data, TAMu$n,  
              method = "REM",  
              RE.type = "Diag")  
  
# Summarize the results  
summary(rem)
```

Stage 1: Pooling Correlation Matrices

Pooling correlation matrices

- REM
- Diagonal constraints of the heterogeneity matrix

```
## 95% confidence intervals: z statistic approximation (robust=FALSE)
## Coefficients:
##              Estimate Std. Error    lbound    ubound z value  Pr(>|z|)
## Intercept1  0.4828802  0.0217706  0.4402105  0.5255498 22.1803 < 2.2e-16 ***
## Intercept2  0.6068160  0.0174683  0.5725787  0.6410533 34.7381 < 2.2e-16 ***
## Intercept3  0.5153033  0.0254332  0.4654551  0.5651514 20.2611 < 2.2e-16 ***
## Intercept4  0.5276429  0.0209277  0.4866253  0.5686604 25.2127 < 2.2e-16 ***
## Intercept5  0.4089908  0.0220210  0.3658303  0.4521512 18.5727 < 2.2e-16 ***
## Intercept6  0.5187087  0.0259696  0.4678093  0.5696081 19.9737 < 2.2e-16 ***
## Tau2_1_1    0.0210148  0.0047131  0.0117772  0.0302524  4.4588 8.243e-06 ***
## Tau2_2_2    0.0134545  0.0030663  0.0074446  0.0194643  4.3878 1.145e-05 ***
## Tau2_3_3    0.0298527  0.0064754  0.0171612  0.0425443  4.6102 4.023e-06 ***
## Tau2_4_4    0.0195271  0.0044114  0.0108809  0.0281733  4.4265 9.578e-06 ***
## Tau2_5_5    0.0210140  0.0048452  0.0115176  0.0305103  4.3371 1.444e-05 ***
## Tau2_6_6    0.0312785  0.0066695  0.0182064  0.0443505  4.6898 2.735e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Stage 1: Pooling Correlation Matrices

Pooling correlation matrices

- REM
- Diagonal constraints of the heterogeneity matrix

```
## Q statistic on the homogeneity of effect sizes: 3622.948
## Degrees of freedom of the Q statistic: 294
## P value of the Q statistic: 0
##
## Heterogeneity indices (based on the estimated Tau2):
##                                     Estimate
## Intercept1: I2 (Q statistic)      0.9178
## Intercept2: I2 (Q statistic)      0.9146
## Intercept3: I2 (Q statistic)      0.9460
## Intercept4: I2 (Q statistic)      0.9217
## Intercept5: I2 (Q statistic)      0.9015
## Intercept6: I2 (Q statistic)      0.9493
```

Stage 1: Pooling Correlation Matrices

Pooling correlation matrices

```
# Extract the pooled correlation matrix
corr.rem <- vec2symMat(coef(rem, select="fixed"), diag=FALSE)
colnames(corr.rem) <- c("PU", "PEU", "ATT", "BI")
rownames(corr.rem) <- c("PU", "PEU", "ATT", "BI")
corr.rem
```

```
##              PU              PEU              ATT              BI
## PU   1.0000000  0.4828802  0.6068160  0.5153033
## PEU  0.4828802  1.0000000  0.5276429  0.4089908
## ATT  0.6068160  0.5276429  1.0000000  0.5187087
## BI   0.5153033  0.4089908  0.5187087  1.0000000
```

Model Estimation

Model estimation

- Lavaan syntax possible
- Translate into the RAM framework

```
# Model specification
Model1 <- " # Structural parameters
           BI ~ ATT + PU
           ATT ~ PEU + PU
           PU ~ PEU

           # Variance of independent variables
           PEU ~~ 1*PEU

           "
```

```
# Convert the model into the RAM language
RAM <- lavaan2RAM(Model1,
                  obs.variables = c("PU", "PEU", "ATT", "BI"))
```

Model Estimation

Model Estimation

```
# Model estimation
# Confidence intervals are likelihood-ratio based (LBCIs)
TS.Model1 <- tssem2(rem, RAM = RAM,
  intervals.type="LB",
  diag.constraints = TRUE,
  model.name = "Model 1 with the direct effect")
```

```
# Summarize the model parameters
summary(TS.Model1)
```

95% confidence intervals: Likelihood-based statistic

Coefficients:

##	Estimate	Std.Error	lbound	ubound	z	value	Pr(> z)
## ATTONPEU	0.32668	NA	0.26635	0.38442		NA	NA
## ATTONPU	0.43668	NA	0.38557	0.48754		NA	NA
## BIONATT	0.35455	NA	0.26355	0.44470		NA	NA
## BIONPU	0.32253	NA	0.23013	0.41347		NA	NA
## PUONPEU	0.49973	NA	0.45797	0.54132		NA	NA
## ATTWITHATT	0.56001	NA	0.51662	0.60096		NA	NA
## BIWITHBI	0.63306	NA	0.58739	0.67590		NA	NA
## PUWITHPU	0.75027	NA	0.70697	0.79026		NA	NA

Model Estimation

Model Estimation

```
# Summarize the model parameters  
summary(TS.Model1)
```

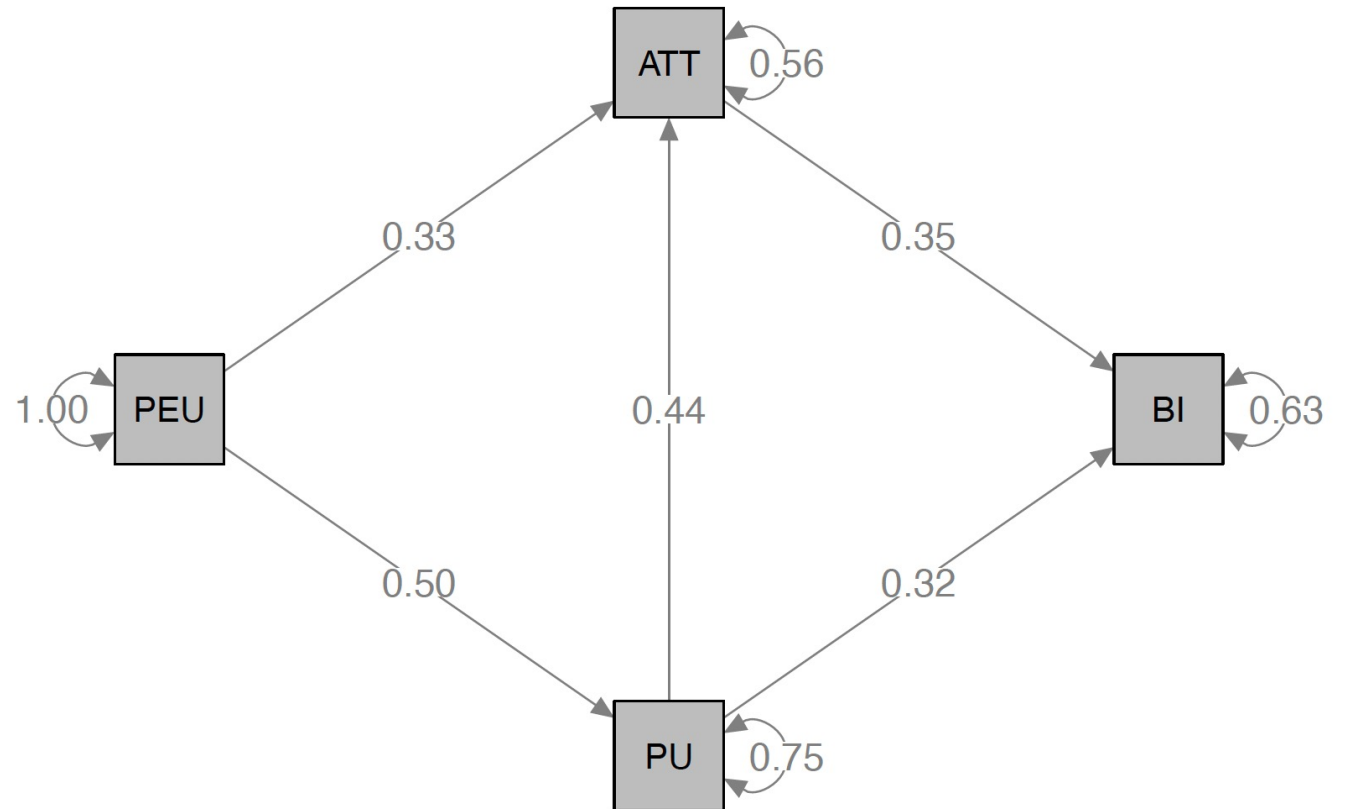
```
## Goodness-of-fit indices:  
##  
## Sample size 14918.0000  
## Chi-square of target model 10.0900  
## DF of target model 1.0000  
## p value of target model 0.0015  
## Number of constraints imposed on "Smatrix" 3.0000  
## DF manually adjusted 0.0000  
## Chi-square of independence model 3002.8440  
## DF of independence model 6.0000  
## RMSEA 0.0247  
## RMSEA lower 95% CI 0.0125  
## RMSEA upper 95% CI 0.0395  
## SRMR 0.0285  
## TLI 0.9818  
## CFI 0.9970  
## AIC 8.0900  
## BIC 0.4797
```

Model Estimation

Model Estimation

- Path diagram

```
# Visualize the model and add its parameters
TS.Model1.plot <- meta2semPlot(TS.Model1)
semPaths(TS.Model1.plot, whatLabels="est",
  rotation=2,
  edge.label.cex = 1.25,
  sizeMan = 8,
  color = "grey",
  layout = "tree2")
```

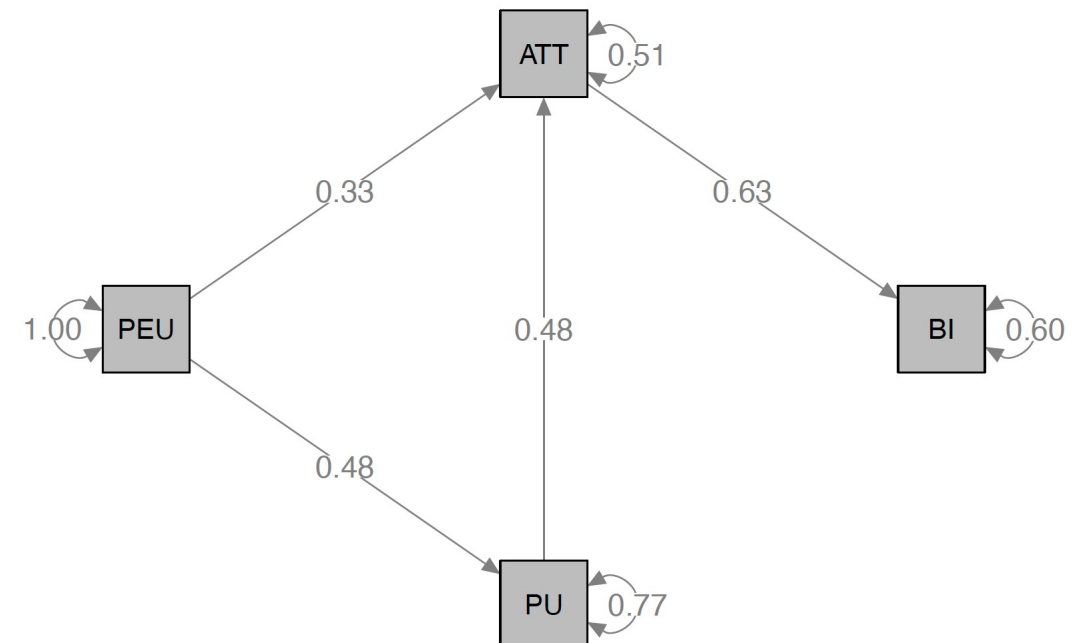
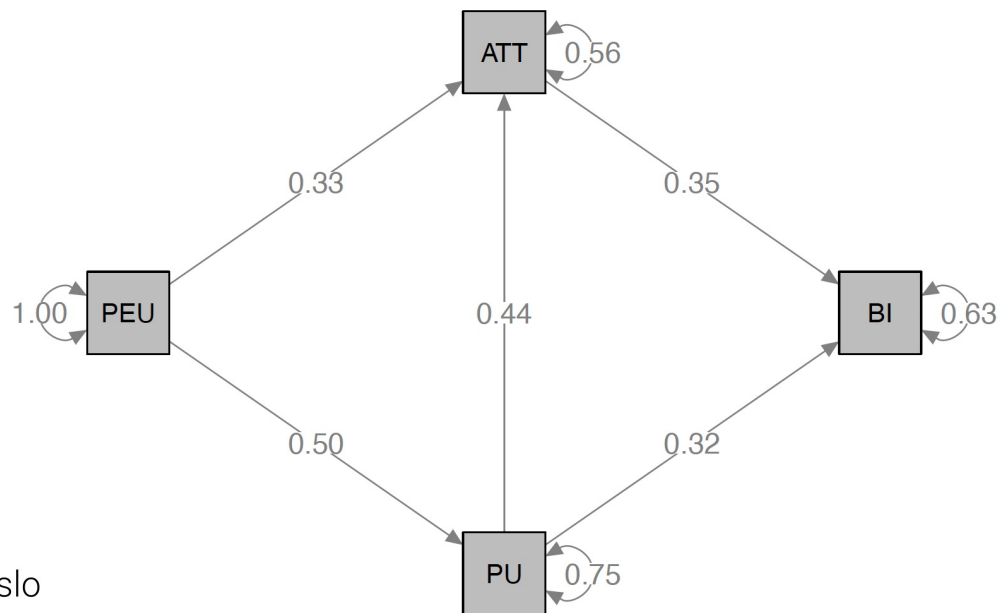


Model Comparison

Model Comparison

```
# Model comparison  
anova(TS.Model1, TS.Model2)
```

Likelihood-ratio test: $\chi^2(1) = 42.1$, $p < .001$



Errors and Warnings

- Check whether the database was organized properly.
- Check Lavaan syntax.
- Rerun the model.

MASEM

Key potential

- Test and compare models representing substantive theory
- Examine heterogeneity in model parameters

Key challenges

- Multilevel structure of the primary and secondary data
- Heterogeneity





The End.

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References

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