Detection of misspecifications in Mplus

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THE PROBLEM AND A SOLUTION

EXAMPLE 2

CONCLUSIONS

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EXAMPLE 1: CORRELATIONS BETWEEN GENETIC POLYMORPHISMS

▶ Rioux et al., Nature Genetics 29, 223 - 228 (2001)

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ALTERNATIVE MODELS

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Chisquare: 32.4, *df*: 2, p: 0.0000





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► CFI: 0.912



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- ► TLI: 0.735



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- ► RMSEA: 0.616 (0.441, 0.812)

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► SRMR: 0.010



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- ► CFI: 0.912
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- ► RMSEA: 0.616 (0.441, 0.812)
- ► SRMR: 0.010
- Modification index (MI) of restricted-to-1-correlation:

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- ▶ Chisquare: 32.4, *df*: 2, p: 0.0000
- ► CFI: 0.912
- ► TLI: 0.735
- ► RMSEA: 0.616 (0.441, 0.812)
- ► SRMR: 0.010
- Modification index (MI) of restricted-to-1-correlation: MI: 4.589 (p = 0.032)

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- ► RMSEA: 0.616 (0.441, 0.812)
- ► SRMR: 0.010
- Modification index (MI) of restricted-to-1-correlation:

MI: 4.589 (p = 0.032)

- EPC: -0.014
 - (this would imply a correlation of 1 0.014 = 0.986)

The unrestricted model estimated with Mplus $% \mathcal{M} = \mathcal{M} = \mathcal{M} + \mathcal{M$



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The unrestricted model estimated with Mplus $% \mathcal{M} = \mathcal{M} = \mathcal{M} + \mathcal{M$



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The unrestricted model estimated with Mplus $% \mathcal{M} = \mathcal{M} = \mathcal{M} + \mathcal{M$



Chisquare: 27.4, *df*: 1, p: 0.0000

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- Chisquare: 27.4, *df*: 1, p: 0.0000
- ► ∆ Chisquare: 5.0, *df*: 1, p: 0.025

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- Chisquare: 27.4, *df*: 1, p: 0.0000
- ► ∆ Chisquare: 5.0, *df*: 1, p: 0.025

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► CFI: 0.923



- ► Chisquare: 27.4, *df*: 1, p: 0.0000
- ► ∆ Chisquare: 5.0, *df*: 1, p: 0.025

- ► CFI: 0.923
- ► TLI: 0.540



- Chisquare: 27.4, *df*: 1, p: 0.0000
- ► ∆ Chisquare: 5.0, *df*: 1, p: 0.025
- ► CFI: 0.923
- ► TLI: 0.540
- ► RMSEA: 0.812 (0.567, 1.087)

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- ► TLI: 0.540
- ► RMSEA: 0.812 (0.567, 1.087)

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► SRMR: 0.012

EXAMPLE 2

CONCLUSIONS

CAN WE CONSIDER ALL FOUR VARIABLES AS INDICATORS OF THE SAME THING?



► What shall we decide?

CAN WE CONSIDER ALL FOUR VARIABLES AS INDICATORS OF THE SAME THING?



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- ▶ But according to all criteria, the model should be rejected.
- This is not what we wanted!
- Whether we used χ², Δχ², or any of the fit measures, we would make a wrong decision.

EXAMPLE 2

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Restricted model:

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(this would imply a correlation of 1 - 0.014 = 0.986)

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WHAT IS THE PROBLEM?

► The misspecification of -0.014 is not substantively relevant

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- Conclusion: when the power of the test is high and the test statistics indicate the model should be rejected, the EPC must be inspected.
- If the misspecification (EPC) does not exceed some threshold of acceptability, the model is not misspecified
- On the other hand, if the EPC *does* exceed the threshold, the model is misspecified

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DECISION RULES

	High power	Low power
Significant MI	Inspect EPC	Misspecification
Nonsignificant MI	No misspecification	Inconclusive

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How do we obtain the power?

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► Exercise: guess the sample size in the previous example ...

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 - ► Hint: the power of the test was very high (close to 1.0)

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POWER

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Power

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- The correct answer is:
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- The high power is due to the very large loadings
- So power does not *just* depend on sample size. Things are not so simple.

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Power

The power of the modification index test to detect a certain misspecification (say, δ) can be determined just from the value of the MI and the EPC.

Saris, W.E., A. Satorra, & W. van der Veld (2009). *Testing Structural Equation Models or Detection of Misspecifications?*, Structural Equation Modeling, 16 pp. 561-582.



INTRODUCTION	Example 1	The problem and a solution	Example 2	CONCLUSIONS

 The program Jrule for Mplus (Oberski 2010) helps you make decisions about misspecifications
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- ► It reads in your Mplus output file and gives information about MI, EPC, the power of the MI test, and the recommended decision based on your own criteria

INTRODUCTION	Example 1	The problem and a solution	EXAMPLE 2	CONCLUSIONS	

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It can be downloaded for free from http://wiki.github.com/daob/JruleMplus/

Image: Second						
Output rie to read:						
Filter by parameter Filter by decision Filter by group					Filter by group	
	•		•			•
Parameter 🔺	Decision •	Group 4 MI	• EPC •	Power 4	NCP	•
GENE1	Not misspecified (EPC < delta)	1 4.58	9 0.029	1.000	54.566	
GENE2	Not misspecified (EPC < delta)	1 4.58	9 0.029	1.000	54.566	
GENE2 WITH GENE1	Not misspecified (EPC < delta)	1 4.58	9 -0.014	1.000	234.133	
IRG1144 WITH IRG1143	Not misspecified (EPC < delta)	1 4.58	9 0.026	1.000	67.885	
IRG1218 WITH IRG1143	Not misspecified (EPC < delta)	1 4.58	9 0.026	1.000	67.885	
IRG1218 WITH IRG1144	Not misspecified (EPC < delta)	1 17.8	02 -0.052	1.000	65.836	
IRG1219 WITH IRG1143	Not misspecified (EPC < delta)	1 17.8	02 -0.051	1.000	68.443	
IRG1219 WITH IRG1144	Not misspecified (EPC < delta)	1 4.58	9 0.027	1.000	62.949	
IRG1219 WITH IRG1218	Not misspecified (EPC < delta)	1 4.58	9 0.027	1.000	62.949	

The current output file is 'Z:\home\daob\work\Presentations\Jrule\ld-1fac.out'.

EXAMPLE 2: PERSONALITY TRAITS AND VOTING

 "Big Five" personality traits: Openness, Conscientiousness, Extraversion, Agreeableness, and Neuroticism

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EXAMPLE 2: PERSONALITY TRAITS AND VOTING

- "Big Five" personality traits: Openness, Conscientiousness, Extraversion, Agreeableness, and Neuroticism
- Correlated with voting
- Hypothesized to affect voting only *indirectly*, through things like "a sense that voting is a duty", "political efficacy" (Gallego & Oberski, frth)

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HIGHLY SIMPLIFIED PATH MODEL



(all regression equations are also controlled for age, sex, and education -- not shown)

HIGHLY SIMPLIFIED PATH MODEL



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Should we introduce a path from Openness/Conscientiousness/Extraversion directly to Voting?

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Should we introduce a path from Openness/Conscientiousness/Extraversion directly to Voting?

I will conclude we should if the effect is bigger than 0.05.



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► Chisquare: 12.3, df = 4*, p = 0.0152



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▶ Chisquare: 12.3, *df* = 4^{*},

p = 0.0152 ► CFI: 0.965

The mediation model estimated with Mplus



(all regression equations are also controlled for age, sex, and education -- not shown)

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► Chisquare: 12.3, df = 4*, p = 0.0152

- ► CFI: 0.965
- ► TLI: 0.948



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► Chisquare: 12.3, df = 4*, p = 0.0152

- ► CFI: 0.965
- ► TLI: 0.948
- ► RMSEA: 0.026



(all regression equations are also controlled for age, sex, and education -- not shown)

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- ► TLI: 0.948
- ► RMSEA: 0.026
- ► WRMR: 0.885



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► MI's and EPC's: VOTE ON CONS MI: 1.349, EPC: 0.062



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► MI's and EPC's: VOTE ON CONS MI: 1.349, EPC: 0.062 VOTE ON EXTR MI: 7.259**, EPC: 0.072



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*df calculated for model with categorical variables (WLSMV estimator)

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Output file to read:							
vote.out							1 🖻
Parameters and misspecifications Misspecification plots Change decision rules							
Filter by paramete	r Filter	by decision	by decision			Filter by group	
VOTEEU09 O	•			•			•
Parameter 🔹	Decision 🔹	Group 4	MI 🖣	EPC 4	Power •	NCP	•
VOTEEU09 ON OPEN	Inconclusive	1	1.419	-0.025	0.479	3.633	
VOTEEU09 ON EXTR	Misspecified (EPC >= delta)	1	7.259	0.040	0.769	7.259	
VOTEEU09 ON CONS	Inconclusive	1	1.349	0.043	0.191	1.167	
	2	•					

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- **CONCLUSIONS**
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- This means we need better measures or a better model or a bigger sample or a combination

OVERALL CONCLUSIONS

 Chi square, fit measures, and MI are *all* affected by the power of the test

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- The power is not only a function of the sample size but can surprise you
- To make a correct decision, one must take into account the power of the test
- ► Saris & a. (2009) suggest one method for doing this
- ► That method is implemented in the free software Jrule for Mplus (Oberski 2010)
Thank you very much for your attention!

http://wiki.github.com/daob/JruleMplus/

daniel.oberski@upf.edu

This presentation: http://daob.org/

