

GC-3 Workshop: Measurement Invariance Testing in R

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Outline

- Brief overview of measurement invariance (MI)
 - Definition of MI, measurement noninvariance, partial invariance
 - Importance of validating the MI assumption
 - Levels of MI
- Illustrative example
 - Investigate MI of the mini-IPIP (International Personality Item Pool) scale across gender.
 - Small Exercise

Motivating Example

- The Positive and Negative Affect Schedule (PANAS) scale has been found to function differently for participants from different cultural backgrounds.

Table 1. Positive and Negative Affect Schedule (PANAS) Scorecard [27].

1	2	3	4	5
Very Slightly or Not at All	A Little	Moderately	Quite a Bit	Extremely
_____1. Interested			_____11. Irritable	
_____2. Distressed			_____12. Alert	
_____3. Excited			_____13. Ashamed	
_____4. Upset			_____14. Inspired	
_____5. Strong			_____15. Nervous	
_____6. Guilty			_____16. Determined	
_____7. Scared			_____17. Attentive	
_____8. Hostile			_____18. Jittery	
_____9. Enthusiastic			_____19. Active	
_____10. Proud			_____20. Afraid	

- Are the observed differences in scale scores caused by real differences in negative affect or the different cultural meanings of scale items?

What Is Measurement Invariance (MI)?

- Formal definition:
- Implied meaning: Using the same questionnaire in different groups (such as countries or at various points in time, or under different conditions) does measure the same construct in the same way.

What Is Measurement Noninvariance (MNI)?

- If there is a violation of MI \Rightarrow Measurement noninvariance/Item bias/Differential item functioning
 - The scale has different measurement properties across groups for individuals with the same latent construct level
 - E.g., Male consistently reported a higher score than female in mathematics self-efficacy scale.
- If MI only holds for a subset of items \Rightarrow Partial invariance

How To Test For MI?

- Multi-Group Confirmatory Factor Analysis (MGCFA)
 - Likelihood ratio test to compare nested models, which means one model with equality constraints of a particular parameter and the other without such constraints.
 - E.g., the invariance of loadings can be tested by comparing the model with freely estimated loadings and the model with has equality constraints of loadings across groups
- Item Response Theory

Multi-Group Confirmatory Factor Analysis (MGCFA)

- CFA assumes the observed items load on a latent factor that represents the construct
- A single-factor MG-CFA

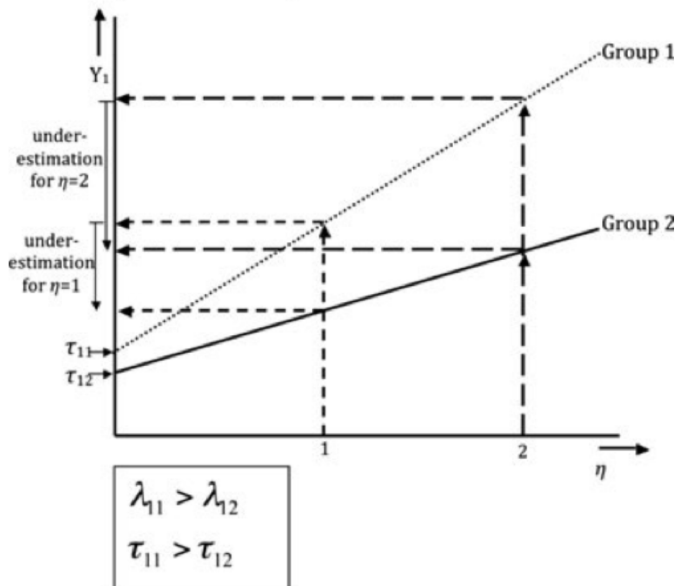
$$y_{ik} = \nu_k + \lambda_k \eta_{ik} + \epsilon_{ik}$$

- y_{ik} and η_{ik} are the observed continuous response and the latent construct score for the i th person in the k th group
- ν_k represents intercepts, λ_k represents factor loadings, and the unique factor variables represents ϵ_{ik} .

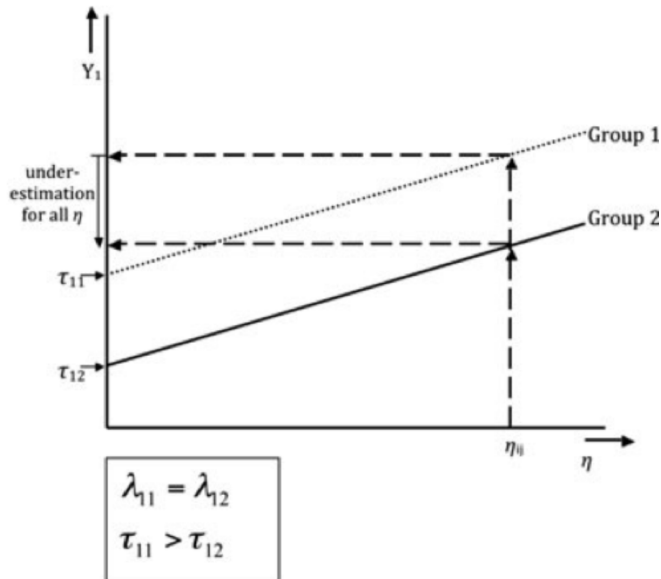
Levels Of MI Testing

- Configural Invariance: the same model holds for all the groups
- Metric/Weak Invariance: factor loadings (slopes) are the same across the groups
- Scalar/Strong Invariance: intercepts and loadings are the same across the groups
- Strict Invariance: unique factor invariance, intercepts and loadings are the same across the groups

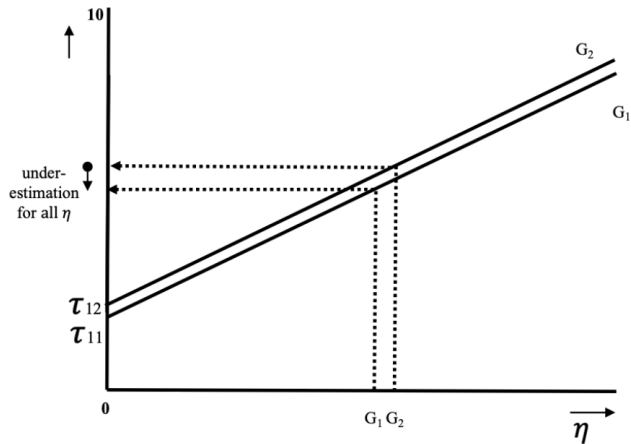
Configural Invariance



Metric Invariance



Scalar Invariance



Illustrative Example

- Goal: Examine measurement invariance of the mini-IPIP across gender.
- Data: Collected from 1994 Spring to 1996 Fall (Ock et al., 2020)
 - 564 participants (239 males, 325 females)
 - 20 items in total, four items per dimension (Agreeableness, Conscientiousness, Extraversion, Neuroticism, Openness to Experience)
 - 5-point Likert-type scale from 1 (*very inaccurate*) to 5 (*very accurate*).

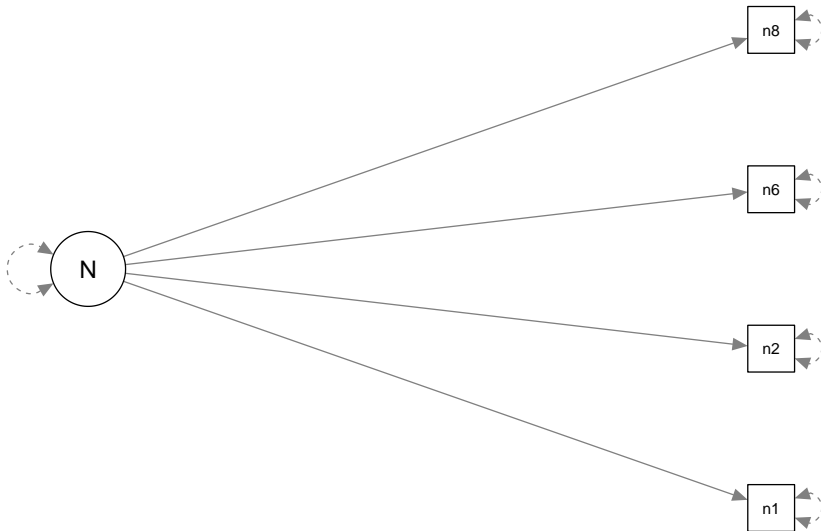
Model Specification

```
data <- read.table(here::here("example", "IPIPFFM.dat"),  
                  header = TRUE)%>%  
  select("sex", "e1", "e4", "e6", "e7", "n1", "n2", "n6", "n8")  
head(data)
```

```
##   sex e1 e4 e6 e7 n1 n2 n6 n8  
## 1   1  4  3  4  4  4  3  3  3  
## 2   1  3  2  2  2  2  4  4  4  
## 3   1  4  5  4  4  1  1  2  2  
## 4   1  2  4  4  3  3  4  4  4  
## 5   1  2  4  4  3  3  2  3  2  
## 6   1  2  4  4  4  2  2  2  1
```

```
mod_n <- 'N =~ n1 + n2 + n6 + n8'  
# check model fit  
fit_n <- cfa(mod = mod_n, data = data)  
# summary(fit_e, fit.measures = TRUE)
```

```
semPlot::semPaths(lavaanify(mod_n), rotation = 2, intercepts = TRUE)
```



Testing Metric Invariance

- Metric Invariance add equality constraints on factor loadings across gender
- The Likelihood Ratio Test (LRT) suggests the metric invariance model has similar fit as the configural invariance model

```
# fit configural invariance model
fit_con <- cfa(mod = mod_n, data = data,
              group = "sex", std.lv = TRUE)
# fit metric invariance model (loadings are constrained to be equal across groups)
fit_met <- cfa(mod = mod_n, data = data, group = "sex",
              group.equal = "loadings", std.lv = TRUE)
# Likelihood Ratio Test
lavTestLRT(fit_con, fit_met)
```

```
## Chi-Squared Difference Test
##
##           Df      AIC      BIC  Chisq Chisq diff Df diff Pr(>Chisq)
## fit_con    4 6547.7 6651.8 5.9861
## fit_met    7 6542.5 6633.6 6.8001    0.81397      3    0.8461
```

Testing Scalar Invariance

- Scalar Invariance add equality constraints on factor loadings and intercepts across groups
- The LRT suggests the scalar invariance model fit significantly worse than the metric invariance model

```
# fit scalar invariance model
fit_sca <- cfa(mod = mod_n, data = data, group = "sex",
              group.equal = c("loadings", "intercepts"),
              std.lv = TRUE)
# Likelihood Ratio test of scalar invariance model and metric invariance model
lavTestLRT(fit_met, fit_sca)
```

```
## Chi-Squared Difference Test
##
##           Df      AIC      BIC   Chisq Chisq diff Df diff Pr(>Chisq)
## fit_met    7 6542.5 6633.6   6.8001
## fit_sca   10 6553.1 6631.1  23.3668      16.567      3 0.0008676 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```


Option 1: pinSearch Function From pinsearch Package

- This function automates the iterative search for noninvariant parameters (loadings, intercepts, residuals)
- Item n1 has noninvariant intercepts across groups

```
res <- pinsearch::pinSearch(config_mod = mod_n, data = data, effect_size = TRUE,  
                           group = "sex", type = "intercepts")  
res$`Non-Invariant Items`
```

```
##   lhs rhs group      type  
## 1  n1      2 intercepts
```

Option 2: Sequential MI Testing By Scale Items

- Manually searching for noninvariant parameters (loadings, intercepts, residuals)
- Find the item associated with the largest change in $\chi^2 \Rightarrow$ item n1

```
# fit scalar invariance model
fit_sca1 <- cfa(mod_n, data = data, group = "sex",
               group.equal = c("loadings", "intercepts"),
               group.partial = "n1 ~ 1", std.lv = TRUE)
lavTestLRT(fit_sca1, fit_sca)
```



```
## Chi-Squared Difference Test
##
##           Df      AIC      BIC  Chisq Chisq diff Df diff Pr(>Chisq)
## fit_sca1   9 6545.5 6627.8 13.736
## fit_sca  10 6553.1 6631.1 23.367      9.6311      1  0.001913 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```

fit_sca2 <- cfa(mod_n, data = data, group = "sex",
               group.equal = c("loadings", "intercepts"),
               group.partial = "n2 ~1", std.lv = TRUE)
lavTestLRT(fit_sca2, fit_sca)

## Chi-Squared Difference Test
##
##           Df      AIC      BIC  Chisq Chisq diff Df diff Pr(>Chisq)
## fit_sca2   9 6552.5 6634.9 20.753
## fit_sca  10 6553.1 6631.1 23.367      2.6133      1      0.106
fit_sca6 <- cfa(mod_n, data = data, group = "sex",
               group.equal = c("loadings", "intercepts"),
               group.partial = "n6 ~1", std.lv = TRUE)
lavTestLRT(fit_sca6, fit_sca)

```

```

## Chi-Squared Difference Test
##
##           Df      AIC      BIC  Chisq Chisq diff Df diff Pr(>Chisq)
## fit_sca6   9 6554.0 6636.4 22.308
## fit_sca  10 6553.1 6631.1 23.367      1.0589      1      0.3035

```

```
fit_sca8 <- cfa(mod_n, data = data, group = "sex",
               group.equal = c("loadings", "intercepts"),
               group.partial = "n8 ~1", std.lv = TRUE)
lavTestLRT(fit_sca8, fit_sca)
```

```
## Chi-Squared Difference Test
```

```
##
```

```
##           Df      AIC      BIC  Chisq  Chisq diff Df diff Pr(>Chisq)
```

```
## fit_sca8   9 6546.4 6628.7 14.629
```

```
## fit_sca  10 6553.1 6631.1 23.367      8.7383      1  0.003116 **
```

```
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Testing Strict Invariance

- The partial strict invariance model fit significantly worse than the partial scalar invariance model

```
fit_str <- cfa(mod_n, data = data, group = "sex",  
              group.equal = c("loadings", "intercepts", "residuals"),  
              group.partial = "n1 ~ 1", std.lv = TRUE)  
lavTestLRT(fit_sca1, fit_str)
```

```
## Chi-Squared Difference Test  
##  
##           Df      AIC      BIC  Chisq Chisq diff Df diff Pr(>Chisq)  
## fit_sca1   9 6545.5 6627.8 13.736  
## fit_str  13 6555.8 6620.9 32.101      18.365      4  0.001047 **  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Testing Strict Invariance

- Items n1,n2 has noninvariant unique factor variances across gender
- Effect size (Standardized mean difference) is 0.239

```
res_str <- pinsearch::pinSearch(config_mod = mod_n, data = data, effect_size = TRUE,  
                               group = "sex", type = "residuals")  
res_str$`Non-Invariant Items`
```

```
##   lhs rhs group      type  
## 1  n1      2 intercepts  
## 2  n2 n2     1 residuals  
## 3  n1 n1     1 residuals  
res_str$effect_size
```

```
##           n1-N  
## dmacs 0.2389943
```

Partial Strict Invariance Model

- Conclusion: items n6, n8 are strict invariant, item n2 is strong/scalar invariant, item n1 is weak/metric invariant
- Final model: partial strict invariance model with freely estimated intercept for item n1 and unique factor variances for items n1 and n2.

```
final_fit <- cfa(mod_n, data = data, group = "sex",  
  group.equal = c("loadings", "intercepts", "residuals"),  
  group.partial = c("n1 ~ 1", "n1 ~~ n1", "n2 ~~ n2"), std.lv = TRUE)
```

Exercise time

Please complete the file “exercise.RMD”.

Additional Topic:

- MI testing for categorical data
- Approximate invariance for many groups
- Practical significance of MI