

# hsls\_els

1 = strongly disagree 2 = disagree 3 = agree 4 = strongly agree

i1 = Teen (9th / 11th grader) confident can do excellent job on (fall 2009 / spring 2012) math tests

i2 = Teen (9th / 11th grader) certain can understand (fall 2009 / spring 2012) math textbook

i3 = Can understand difficult math class (ELS ONLY!)

i4 = Teen confident can do an excellent job on math assignments

i5 = Teen certain can master skills in math course

Can understand difficult math class

```
library(dplyr)
```

Attaching package: 'dplyr'

The following objects are masked from 'package:stats':

filter, lag

The following objects are masked from 'package:base':

intersect, setdiff, setequal, union

```
library(lavaan)
```

This is lavaan 0.6-12

lavaan is FREE software! Please report any bugs.

```
library(ltm)
```

Warning: package 'ltm' was built under R version 4.3.2

Loading required package: MASS

Attaching package: 'MASS'

The following object is masked from 'package:dplyr':

```
select
```

Loading required package: msm

Warning: package 'msm' was built under R version 4.3.2

Loading required package: polycor

Warning: package 'polycor' was built under R version 4.3.2

```
library(sjlabelled)
```

Warning: package 'sjlabelled' was built under R version 4.3.3

Attaching package: 'sjlabelled'

The following object is masked from 'package:dplyr':

```
as_label
```

```
library(kableExtra)
```

Warning: package 'kableExtra' was built under R version 4.3.3

Attaching package: 'kableExtra'

The following object is masked from 'package:dplyr':

group\_rows

```
library(sirt)
```

Warning: package 'sirt' was built under R version 4.3.3

- sirt 4.1-15 (2024-02-06 00:05:40)

```
library(mirt)
```

Warning: package 'mirt' was built under R version 4.3.3

Loading required package: stats4

Loading required package: lattice

Attaching package: 'mirt'

The following object is masked from 'package:ltm':

Science

```
#source("code/download_data.R")  
source("F:/Users/alex/OneDrive/Documents/data/prepare_data.R")
```

Warning: package 'here' was built under R version 4.3.3

here() starts at F:/Users/alex/OneDrive/Documents/data

Joining with `by = join\_by(stu\_id, sch\_id, STRAT\_ID, psu, sex, i1, i2, i4, i5, i1\_2, i2\_2, i4\_2, i5\_2, sample)`

Warning in rm(dat\_full, els, els\_02\_12\_byf3pststu\_v1\_0, hsls, hsls\_17\_student\_pets\_sr\_v1\_0, : object 'no\_nas\_hsls5' not found

```

m_items <- paste0("i", 1:5)
m_items_2 <- paste0(m_items, "_2")
# get subset of relevant variables
dat <- dat[, c("stu_id", "sample", "sex", "dropout", m_items, m_items_2)]

dat$mean_score <- c(rowMeans(dat[dat$sample == "ELS", m_items], na.rm = TRUE),
                    rowMeans(dat[dat$sample == "HSLs", m_items[-3]], na.rm = TRUE))
dat$mean_score_2 <- c(rowMeans(dat[dat$sample == "ELS", m_items_2], na.rm = TRUE),
                      rowMeans(dat[dat$sample == "HSLs", m_items_2[-3]], na.rm = TRUE))

# Creating only HSLs
hsls <- subset(dat, sample == "HSLs")

hsls_1 <- hsls[, c("i1", "i2", "i4", "i5")]
head(hsls_1)

```

	i1	i2	i4	i5
16198	4	3	4	3
16199	3	3	4	3
16200	4	2	4	3
16201	3	3	3	3
16202	3	3	3	3
16203	4	4	4	3

```

hsls_2 <- hsls[, c("i1_2", "i2_2", "i4_2", "i5_2")]
head(hsls_2)

```

	i1_2	i2_2	i4_2	i5_2
16198	4	4	4	4
16199	2	2	3	2
16200	2	1	3	3
16201	NA	NA	NA	NA
16202	NA	NA	NA	NA
16203	3	3	3	3

```

hsls_1_noNA <- na.omit(hsls_1)
hsls_2_noNA <- na.omit(hsls_2)

```

```

# Creating only ELS
els <- subset(dat, sample == "ELS")

els_1 <- els[, c("i1", "i2", "i3", "i4", "i5")]
head(els_1)

  i1 i2 i3 i4 i5
1  2  1  2  2  1
2  4  3  4  4  4
3  3  2  2  3  2
4  4  3  3  3  4
5  2  2  3  3  3
6  2  2  3  3 NA

els_2 <- els[, c("i1_2", "i2_2", "i3_2", "i4_2", "i5_2")]
head(els_2)

  i1_2 i2_2 i3_2 i4_2 i5_2
1     3     2     2     2     3
2     3     3     3     3     3
3     2     3     3     2     2
4     3     2     2     3     4
5     3     3     3     4     4
6    NA    NA    NA    NA    NA

els_1_noNA <- na.omit(els_1)
els_2_noNA <- na.omit(els_2)

cfa_config <- '
group: ELS
math =~ NA      * i1 +
        el2_1 * i2 +
        el3_1 * i3 +
        el4_1 * i4 +
        el5_1 * i5

# Naming the intercepts!
i1 ~ nu1_1 * 1
i2 ~ nu2_1 * 1

```

```

i3 ~ nu3_1 * 1
i4 ~ nu4_1 * 1
i5 ~ nu5_1 * 1

# Naming the residual variances!
i1 ~~ theta1_1 * i1
i2 ~~ theta2_1 * i2
i3 ~~ theta3_1 * i3
i4 ~~ theta4_1 * i4
i5 ~~ theta5_1 * i5

# Adding the covariances
i1 ~~ i2
i2 ~~ i3

# Fixing latent variance to 1, as we freed first factor loading
math ~~ 1 * math

# Fixing latent mean to 0 for identification?
math ~ 0 * 1

group: HSLS
math =~ NA      * i1 +
          hl2_2 * i2 +
          hl4_2 * i4 +
          hl5_2 * i5

# Naming the intercepts!
i1 ~ nu1_2 * 1
i2 ~ nu2_2 * 1
i4 ~ nu4_2 * 1
i5 ~ nu5_2 * 1

# Naming the residual variances!
i1 ~~ theta1_2 * i1
i2 ~~ theta2_2 * i2
i4 ~~ theta4_2 * i4
i5 ~~ theta5_2 * i5

# Adding the covariances
#i1 ~~ i2

```

```

    i2 ~~ i4

    # Fixing latent variance to 1, as we freed first factor loading
    math ~~ 1 * math

    # Fixing latent mean to 0 for identification?
    math ~ 0 * 1
  ,

fit_config <- cfa(cfa_config, data = dat, group = "sample",
                 estimator = "MLR", missing = "FIML", se = "robust.mlr")

```

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some observed variables are not in the model

24 32 33 34 35 36 37 38 39 40 44 45 47 50 51 56 57 65 66 67 69 72 73 77 78 87 99 101 110 111

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some observed variables are not in the model

16206 16211 16212 16222 16223 16226 16227 16229 16235 16241 16250 16252 16268 16273 16274 16275

```

s_config <- summary(fit_config, fit.measures = TRUE, standardized = TRUE)

mod_indices <- modindices(fit_config, sort. = TRUE, free.remove = FALSE)
head(mod_indices)

```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
35	i1	~~	i3	1	ELS	1	81.621	0.033	0.033	0.119	0.119
42	i4	~~	i5	1	ELS	1	81.621	0.038	0.038	0.218	0.218
37	i1	~~	i5	1	ELS	1	76.182	-0.027	-0.027	-0.113	-0.113
40	i3	~~	i4	1	ELS	1	27.064	-0.018	-0.018	-0.090	-0.090
39	i2	~~	i5	1	ELS	1	12.584	0.011	0.011	0.043	0.043
38	i2	~~	i4	1	ELS	1	12.584	-0.011	-0.011	-0.045	-0.045

```

# Just for ELS
mod_indices_els <- mod_indices[mod_indices$group == "ELS", ]
head(mod_indices_els)

```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
35	i1	~~	i3	1	ELS	1	81.621	0.033	0.033	0.119	0.119
42	i4	~~	i5	1	ELS	1	81.621	0.038	0.038	0.218	0.218

```

37 i1 ~~ i5      1  ELS      1 76.182 -0.027 -0.027 -0.113 -0.113
40 i3 ~~ i4      1  ELS      1 27.064 -0.018 -0.018 -0.090 -0.090
39 i2 ~~ i5      1  ELS      1 12.584  0.011  0.011  0.043  0.043
38 i2 ~~ i4      1  ELS      1 12.584 -0.011 -0.011 -0.045 -0.045

```

```

# Just for HSLs
mod_indices_hsls <- mod_indices[mod_indices$group == "HSLs", ]
head(mod_indices_hsls)

```

```

      lhs op rhs block group level      mi      epc sepc.lv sepc.all sepc.nox
47  i4 ~~ i5      2  HSLs      1 0.044 -0.001 -0.001 -0.004 -0.004
43  i1 ~~ i2      2  HSLs      1 0.044 -0.001 -0.001 -0.003 -0.003
46  i2 ~~ i5      2  HSLs      1 0.044  0.001  0.001  0.003  0.003
44  i1 ~~ i4      2  HSLs      1 0.044  0.001  0.001  0.004  0.004
25  i2 ~1          2  HSLs      1 0.000  0.000  0.000  0.000  0.000
23 math =~ i5      2  HSLs      1 0.000  0.000  0.000  0.000  0.000

```

#CONFIG ELS + HSLs

```

config_comb <- '
# ELS
group: 1

# Time point 1
math_t1 =~ NA      * i1 +
            el2_1 * i2 +
            el3_1 * i3 +
            el4_1 * i4 +
            el5_1 * i5

# Naming the intercepts!
i1 ~ enu1_1 * 1
i2 ~ enu2_1 * 1
i3 ~ enu3_1 * 1
i4 ~ enu4_1 * 1
i5 ~ enu5_1 * 1

# Naming the residual variances!
i1 ~~ etheta1_1 * i1
i2 ~~ etheta2_1 * i2
i3 ~~ etheta3_1 * i3

```



```

i4 ~~ etheta4_1 * i4
i5 ~~ etheta5_1 * i5

# Fixing latent variance to 1, as we freed first factor loading
math_t1 ~~ 1 * math_t1

# Fixing latent mean to 0 for identification?
math_t1 ~ 0 * 1

# Time point 2
math_t2 =~ NA      * i1_2 +
            e12_2 * i2_2 +
            e13_2 * i3_2 +
            e14_2 * i4_2 +
            e15_2 * i5_2

# Naming the intercepts!
i1_2 ~ enu1_2 * 1
i2_2 ~ enu2_2 * 1
i3_2 ~ enu3_2 * 1
i4_2 ~ enu4_2 * 1
i5_2 ~ enu5_2 * 1

# Naming the residual variances!
i1_2 ~~ etheta1_2 * i1_2
i2_2 ~~ etheta2_2 * i2_2
i3_2 ~~ etheta3_2 * i3_2
i4_2 ~~ etheta4_2 * i4_2
i5_2 ~~ etheta5_2 * i5_2

## Adding the covariances ##
i1 ~~ i2
i1_2 ~~ i2_2

i1 ~~ i3
i1_2 ~~ i3_2

i2 ~~ i3
i2_2 ~~ i3_2

i4 ~~ i5

```

```

i4_2 ~~ i5_2

# Fixing latent variance to 1, as we freed first factor loading
math_t2 ~~ 1 * math_t2

# Fixing latent mean to 0 for identification
math_t2 ~ 0 * 1

# Correlations across time
math_t1 ~~ math_t2
i1 ~~ i1_2
i2 ~~ i2_2
i3 ~~ i3_2
i4 ~~ i4_2
i5 ~~ i5_2

# HSLS
group: 2
# Time Point 1
math_t1 =~ NA    * i1 +
           hl2_1 * i2 +
           #hl3_1 * i3 +
           hl4_1 * i4 +
           hl5_1 * i5

# Naming the intercepts!
i1 ~ hnu1_1 * 1
i2 ~ hnu2_1 * 1
#i3 ~ hnu3_1 * 1
i4 ~ hnu4_1 * 1
i5 ~ hnu5_1 * 1

# Naming the residual variances!
i1 ~~ htheta1_1 * i1
i2 ~~ htheta2_1 * i2
#i3 ~~ htheta3_1 * i3
i4 ~~ htheta4_1 * i4
i5 ~~ htheta5_1 * i5

```

```

# Fixing latent variance to 1, as we freed first factor loading
math_t1 ~~ 1 * math_t1

# Fixing latent mean to 0 for identification
math_t1 ~ 0 * 1

# Time Point 2
math_t2 =~ NA      * i1_2 +
              hl2_2 * i2_2 +
              #hl3_2 * i3_2 +
              hl4_2 * i4_2 +
              hl5_2 * i5_2

# Naming the intercepts!
i1_2 ~ hnu1_2 * 1
i2_2 ~ hnu2_2 * 1
#i3_2 ~ hnu3_2 * 1
i4_2 ~ hnu4_2 * 1
i5_2 ~ hnu5_2 * 1

# Naming the residual variances!
i1_2 ~~ htheta1_2 * i1_2
i2_2 ~~ htheta2_2 * i2_2
#i3_2 ~~ htheta3_1 * i3_2
i4_2 ~~ htheta4_2 * i4_2
i5_2 ~~ htheta5_2 * i5_2

## Adding the covariances ##
i1 ~~ i2
i1_2 ~~ i2_2

i1 ~~ i4
i1_2 ~~ i4_2

i2 ~~ i4
i2_2 ~~ i4_2

i1 ~~ i5
i1_2 ~~ i5_2

# Fixing latent variance to 1, as we freed first factor loading

```

```

math_t2 ~~ 1 * math_t2

# Fixing latent mean to 0 for identification
math_t2 ~ 0 * 1

# Correlations across time
math_t1 ~~ math_t2
i1 ~~ i1_2
i2 ~~ i2_2
#i3 ~~ i3_2
i4 ~~ i4_2
i5 ~~ i5_2

'

fit_config_comb <- cfa(config_comb, data = dat, group = "sample",
                        estimator = "MLR", missing = "FIML", se = "robust.mlr")

```

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some observed variables are not in the model

24 33 38 44 47 50 56 65 66 67 77 87 99 101 110 119 123 124 130 139 140 164 166 173 177 185

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some observed variables are not in the model

16211 16227 16369 16424 16479 16500 16511 16581 16588 16613 16647 16649 16662 16686 16697

```
fit_config_comb
```

lavaan 0.6-12 ended normally after 98 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	81	
Number of observations per group:	Used	Total
ELS	13926	16197
HSLS	22839	23503
Number of missing patterns per group:		
ELS	89	
HSLS	50	

Model Test User Model:

	Standard	Robust
Test Statistic	274.812	235.207
Degrees of freedom	28	28
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.168
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	218.655	187.143
HSLs	56.158	48.065

```
head(modindices(fit_config_comb, sort. = TRUE, free.remove = FALSE))
```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
97	math_t2	=~	i3	1	1	1	47.242	-0.068	-0.068	-0.071	-0.071
91	math_t1	=~	i2_2	1	1	1	38.012	0.050	0.050	0.056	0.056
90	math_t1	=~	i1_2	1	1	1	35.233	-0.057	-0.057	-0.064	-0.064
109	i2	~~	i3_2	1	1	1	33.516	0.022	0.022	0.078	0.078
101	i1	~~	i5	1	1	1	32.875	-0.018	-0.018	-0.071	-0.071
126	i1_2	~~	i4_2	1	1	1	30.409	0.022	0.022	0.088	0.088

```
s_config_comb <- summary(fit_config_comb, fit.measures = TRUE)
```

## WEAK ELS + HSLs

```
# Only the first latent variance is set to 1, the rest are free.
# Latent means all 0.
```

```
weak_comb <- '
```

```
#####
##                               ELS (Group 1)                               ##
#####
group: ELS

#####
# Time Point 1
#####
```

```

math_t1 =~ NA      * i1 +
           12_1 * i2 +
           13_1 * i3 +
           14_1 * i4 +
           15_1 * i5

# Intercepts
i1 ~ enu1_1 * 1
i2 ~ enu2_1 * 1
i3 ~ enu3_1 * 1
i4 ~ enu4_1 * 1
i5 ~ enu5_1 * 1

# Residual variances
i1 ~~ etheta1_1 * i1
i2 ~~ etheta2_1 * i2
i3 ~~ etheta3_1 * i3
i4 ~~ etheta4_1 * i4
i5 ~~ etheta5_1 * i5

# Fix latent variance to 1 for identification, fix mean to 0
math_t1 ~~ 1 * math_t1
math_t1 ~ 0 * 1

#####
# Time Point 2
#####
math_t2 =~ NA      * i1_2 +
           12_2 * i2_2 +
           13_2 * i3_2 +
           14_2 * i4_2 +
           15_2 * i5_2

# Intercepts
i1_2 ~ enu1_2 * 1
i2_2 ~ enu2_2 * 1
i3_2 ~ enu3_2 * 1
i4_2 ~ enu4_2 * 1
i5_2 ~ enu5_2 * 1

# Residual variances

```

```

i1_2 ~~ etheta1_2 * i1_2
i2_2 ~~ etheta2_2 * i2_2
i3_2 ~~ etheta3_2 * i3_2
i4_2 ~~ etheta4_2 * i4_2
i5_2 ~~ etheta5_2 * i5_2

# Covariances among items
i1 ~~ i2
i1_2 ~~ i2_2

i1 ~~ i3
i1_2 ~~ i3_2

i2 ~~ i3
i2_2 ~~ i3_2

i4 ~~ i5
i4_2 ~~ i5_2

# Fix latent variance to 1 for identification, free mean
math_t2 ~~ 1 * math_t2
math_t2 ~ NA * 1

# Correlations across time
math_t1 ~~ math_t2
i1 ~~ i1_2
i2 ~~ i2_2
i3 ~~ i3_2
i4 ~~ i4_2
i5 ~~ i5_2

#####
##                               HSLs (Group 2)                               ##
#####
group: HSLs

#####
# Time Point 1
#####
math_t1 =~ NA    *i1 +

```

```

        12_1 * i2 +    # Same label as ELS
        # no i3 in HSLS
        14_1 * i4 +    # Same label as ELS
        15_1 * i5      # Same label as ELS

# Intercepts
i1 ~ hnu1_1 * 1
i2 ~ hnu2_1 * 1
# i3 ~ hnu3_1 * 1 (item not in HSLS)
i4 ~ hnu4_1 * 1
i5 ~ hnu5_1 * 1

# Residual variances
i1 ~~ htheta1_1 * i1
i2 ~~ htheta2_1 * i2
# i3 ~~ htheta3_1 * i3 (item not in HSLS)
i4 ~~ htheta4_1 * i4
i5 ~~ htheta5_1 * i5

# Free latent variance and free latent mean
math_t1 ~~ var_hsls * math_t1
math_t1 ~ NA * 1

#####
# Time Point 2
#####
math_t2 =~ NA    * i1_2 +
        12_2 * i2_2 +    # Same label as ELS
        # no i3_2 in HSLS at Time 2
        14_2 * i4_2 +    # Same label as ELS
        15_2 * i5_2      # Same label as ELS

# Intercepts
i1_2 ~ hnu1_2 * 1
i2_2 ~ hnu2_2 * 1
# i3_2 ~ hnu3_2 * 1 (item not in HSLS)
i4_2 ~ hnu4_2 * 1
i5_2 ~ hnu5_2 * 1

# Residual variances

```



```

i1_2 ~~ htheta1_2 * i1_2
i2_2 ~~ htheta2_2 * i2_2
# i3_2 ~~ htheta3_2 * i3_2 (item not in HSLS)
i4_2 ~~ htheta4_2 * i4_2
i5_2 ~~ htheta5_2 * i5_2

# Covariances among items
i1 ~~ i2
i1_2 ~~ i2_2

i1 ~~ i4
i1_2 ~~ i4_2

i2 ~~ i4
i2_2 ~~ i4_2

i1 ~~ i5
i1_2 ~~ i5_2

# Free latent variance and free latent mean
math_t2 ~~ var_hsls_t2 * math_t2
math_t2 ~ NA * 1

# Correlations across time
math_t1 ~~ math_t2
i1 ~~ i1_2
i2 ~~ i2_2
# i3 ~~ i3_2 (item not in HSLS)
i4 ~~ i4_2
i5 ~~ i5_2

'

fit_weak_comb <- cfa(weak_comb, data = dat, group = "sample",
  estimator = "MLR", missing = "FIML", se = "robust.mlr")

```

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some items are not in the model

24 33 38 44 47 50 56 65 66 67 77 87 99 101 110 119 123 124 130 139 140 164 166 173 177 185

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some items are not in the model

16211 16227 16369 16424 16479 16500 16511 16581 16588 16613 16647 16649 16662 16686 16697

Warning in lav\_model\_vcov(lavmodel = lavmodel, lavsamplestats = lavsamplestats, : lavaan WARN  
 The variance-covariance matrix of the estimated parameters (vcov)  
 does not appear to be positive definite! The smallest eigenvalue  
 (= 1.885309e-19) is close to zero. This may be a symptom that the  
 model is not identified.

```
fit_weak_comb
```

lavaan 0.6-12 ended normally after 88 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	86
Number of equality constraints	6
Number of observations per group:	Used Total
ELS	13926 16197
HSLs	22839 23503
Number of missing patterns per group:	
ELS	89
HSLs	50

Model Test User Model:

	Standard	Robust
Test Statistic	364.936	288.592
Degrees of freedom	29	29
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.265
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	259.106	204.901
HSLs	105.830	83.691

```
head(modindices(fit_weak_comb, sort. = TRUE, free.remove = FALSE))
```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
103	math_t2	=~	i3	1	ELS	1	70.861	-0.076	-0.076	-0.079	-0.079
141	math_t1	=~	i5_2	2	HSLs	1	49.758	-0.046	-0.037	-0.048	-0.048
107	i1	~~	i5	1	ELS	1	34.540	-0.019	-0.019	-0.074	-0.074
97	math_t1	=~	i2_2	1	ELS	1	33.184	0.040	0.040	0.045	0.045

```

96  math_t1 =~ i1_2      1    ELS      1 31.078 -0.049 -0.049 -0.055 -0.055
106      i1 ~~    i4      1    ELS      1 30.684  0.018  0.018  0.070  0.070

```

```

s_weak_comb <- summary(fit_weak_comb, fit.measures = TRUE, standardized = TRUE)

lavTestLRT(fit_config_comb, fit_weak_comb)

```

Scaled Chi-Squared Difference Test (method = "satorra.bentler.2001")

lavaan NOTE:

The "Chisq" column contains standard test statistics, not the robust test that should be reported per model. A robust difference test is a function of two standard (not robust) statistics.

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit_config_comb	28	474900	475590	274.81			
fit_weak_comb	29	474988	475669	364.94	22.776	1	1.82e-06 ***

---  
Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

## STRONG ELS + HSLS

```

strong_comb <- '

#####
##                               ELS (Group 1)                               ##
#####
group: ELS

#####
# Time Point 1
#####
math_t1 =~ NA      * i1 +
              12_1 * i2 +
              13_1 * i3 +
              14_1 * i4 +
              15_1 * i5

# Intercepts
# Same labels for both groups!

```

```

i1 ~ nu1_1 * 1
i2 ~ nu2_1 * 1
i3 ~ nu3_1 * 1
i4 ~ nu4_1 * 1
i5 ~ nu5_1 * 1

# Residual variances
i1 ~~ etheta1_1 * i1
i2 ~~ etheta2_1 * i2
i3 ~~ etheta3_1 * i3
i4 ~~ etheta4_1 * i4
i5 ~~ etheta5_1 * i5

# Fix latent variance to 1 for identification, fix mean to 0
math_t1 ~~ 1 * math_t1
math_t1 ~ 0 * 1

#####
# Time Point 2
#####
math_t2 =~ 1*i1_2 +
            12_2 * i2_2 +
            13_2 * i3_2 +
            14_2 * i4_2 +
            15_2 * i5_2

# Intercepts
# Same labels for both groups!
i1_2 ~ nu1_2 * 1
i2_2 ~ nu2_2 * 1
i3_2 ~ nu3_2 * 1
i4_2 ~ nu4_2 * 1
i5_2 ~ nu5_2 * 1

# Residual variances
i1_2 ~~ etheta1_2 * i1_2
i2_2 ~~ etheta2_2 * i2_2
i3_2 ~~ etheta3_2 * i3_2
i4_2 ~~ etheta4_2 * i4_2
i5_2 ~~ etheta5_2 * i5_2

```

```

# Covariances among items
i1 ~~ i2
i1_2 ~~ i2_2

i1 ~~ i3
i1_2 ~~ i3_2

i2 ~~ i3
i2_2 ~~ i3_2

i4 ~~ i5
i4_2 ~~ i5_2

# Fix latent variance to 1 for identification, free mean
math_t2 ~~ var_els_t2 * math_t2
math_t2 ~ NA * 1

# Correlations across time
math_t1 ~~ math_t2
i1 ~~ i1_2
i2 ~~ i2_2
i3 ~~ i3_2
i4 ~~ i4_2
i5 ~~ i5_2

#####
##                                HSLs (Group 2)                                ##
#####
group: HSLs

#####
# Time Point 1
#####
math_t1 =~ NA      * i1 +
           12_1 * i2 +   # Same label as ELS
           # no i3 in HSLs
           14_1 * i4 +   # Same label as ELS
           15_1 * i5     # Same label as ELS

# Intercepts

```

```

# Same labels for both groups!
i1 ~ nu1_1 * 1
i2 ~ nu2_1 * 1
# i3 ~ hnu3_1 * 1 (item not in HSLS)
i4 ~ nu4_1 * 1
i5 ~ nu5_1 * 1

# Residual variances
i1 ~~ htheta1_1 * i1
i2 ~~ htheta2_1 * i2
# i3 ~~ htheta3_1 * i3 (item not in HSLS)
i4 ~~ htheta4_1 * i4
i5 ~~ htheta5_1 * i5

# Free latent variance and free latent mean
math_t1 ~~ var_hsls * math_t1
math_t1 ~ NA * 1

#####
# Time Point 2
#####
math_t2 =~ NA * i1_2 +
          l2_2 * i2_2 + # Same label as ELS
          # no i3_2 in HSLS at Time 2
          l4_2 * i4_2 + # Same label as ELS
          l5_2 * i5_2   # Same label as ELS

# Intercepts
# Same labels for both groups!

i1_2 ~ nu1_2 * 1
i2_2 ~ nu2_2 * 1
# i3_2 ~ hnu3_2 * 1 (item not in HSLS)
i4_2 ~ nu4_2 * 1
i5_2 ~ nu5_2 * 1

# Residual variances
i1_2 ~~ htheta1_2 * i1_2
i2_2 ~~ htheta2_2 * i2_2
# i3_2 ~~ htheta3_2 * i3_2 (item not in HSLS)

```

```

i4_2 ~~ htheta4_2 * i4_2
i5_2 ~~ htheta5_2 * i5_2

# Covariances among items
i1 ~~ i2
i1_2 ~~ i2_2

i1 ~~ i4
i1_2 ~~ i4_2

i2 ~~ i4
i2_2 ~~ i4_2

i1 ~~ i5
i1_2 ~~ i5_2

# Free latent variance and free latent mean
math_t2 ~~ var_hsls_t2 * math_t2
math_t2 ~ NA * 1

# Correlations across time
math_t1 ~~ math_t2
i1 ~~ i1_2
i2 ~~ i2_2
# i3 ~~ i3_2 (item not in HSLs)
i4 ~~ i4_2
i5 ~~ i5_2

'

fit_strong_comb <- cfa(strong_comb, data = dat, group = "sample",
                      estimator = "MLR", missing = "FIML", se = "robust.mlr")

```

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some items are not in the data matrix: 24 33 38 44 47 50 56 65 66 67 77 87 99 101 110 119 123 124 130 139 140 164 166 173 177 185

Warning in lav\_data\_full(data = data, group = group, cluster = cluster, : lavaan WARNING: some items are not in the data matrix: 16211 16227 16369 16424 16479 16500 16511 16581 16588 16613 16647 16649 16662 16686 16697

```
fit_strong_comb
```

lavaan 0.6-12 ended normally after 172 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	86	
Number of equality constraints	14	
Number of observations per group:	Used	Total
ELS	13926	16197
HSLs	22839	23503
Number of missing patterns per group:		
ELS	89	
HSLs	50	

Model Test User Model:

	Standard	Robust
Test Statistic	794.745	705.843
Degrees of freedom	37	37
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.126
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	521.566	463.223
HSLs	273.178	242.620

```
head(modindices(fit_strong_comb, sort. = TRUE, free.remove = FALSE))
```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
112	math_t2	=~	i4	1	ELS	1	237.142	0.027	0.021	0.022	0.022
152	math_t2	=~	i4	2	HSLs	1	222.944	-0.025	-0.017	-0.023	-0.023
153	math_t2	=~	i5	2	HSLs	1	168.010	0.022	0.015	0.021	0.021
113	math_t2	=~	i5	1	ELS	1	166.953	-0.024	-0.018	-0.019	-0.019
9	i4	~1		1	ELS	1	122.263	-0.053	-0.053	-0.056	-0.056
10	i5	~1		1	ELS	1	92.968	0.047	0.047	0.051	0.051

```
s_strong_comb <- summary(fit_strong_comb, fit.measures = TRUE, standardized = TRUE)
lavTestLRT(fit_weak_comb, fit_strong_comb)
```

Scaled Chi-Squared Difference Test (method = "satorra.bentler.2001")



lavaan NOTE:

The "Chisq" column contains standard test statistics, not the robust test that should be reported per model. A robust difference test is a function of two standard (not robust) statistics.

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
fit_weak_comb	29	474988	475669	364.94			
fit_strong_comb	37	475402	476015	794.74	689.28	8	< 2.2e-16 ***

---

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

**Strong invariance holds! What if across groups AND time?**

## HSLs ONLY

```
# hsls_time <- '  
#   math_T1 =~ NA * i1 +  
#               12_1 * i2 +  
#               14_1 * i4 +  
#               15_1 * i5  
#  
#  
#   # Fixing latent variance to 1, as we freed first factor loading  
#   math_T1 ~~ 1 * math_T1  
#  
#   # Fixing latent mean to 0 for identification?  
#   math_T1 ~ 0 * 1  
#  
#   # Time Point 2  
#   math_T2 =~ NA * i1_2 +  
#               12_2 * i2_2 +  
#               14_2 * i4_2 +  
#               15_2 * i5_2  
#  
#  
#  
#   # Adding the covariances  
#   i1_2 ~~ i2_2  
#   i2_2 ~~ i4_2
```

```

#
#   i1 ~~ i5
#   i1_2 ~~ i5_2
#
#   # Fixing latent variance to 1, as we freed first factor loading
#   math_T2 ~~ 1 * math_T2
#
#   # Fixing latent mean to 0 for identification?
#   math_T2 ~ 0 * 1
#
#   # Correlations across time
#   math_T1 ~~ math_T2
#   i1 ~~ i1_2
#   i2 ~~ i2_2
#   i4 ~~ i4_2
#   i5 ~~ i5_2
# '
#
# fit_hsls_time <- cfa(hsls_time, data = hsls,
#                       estimator = "MLR", missing = "FIML", se = "robust.mlr")
# fit_hsls_time
# head(modindices(fit_hsls_time, sort. = TRUE, free.remove = FALSE))
#
#
# s_hsls_time <- summary(fit_hsls_time, fit.measures = TRUE)
#
#

```

## ELS ONLY

```

#
# els_time <- '
#   math_T1 =~ NA * i1 +
#               12_1 * i2 +
#               13_1 * i3 +
#               14_1 * i4 +
#               15_1 * i5
#
#
#   # Fixing latent variance to 1, as we freed first factor loading

```

```

# math_T1 ~~ 1 * math_T1
#
# # Fixing latent mean to 0 for identification?
# math_T1 ~ 0 * 1
#
# # Time Point 2
# math_T2 =~ NA * i1_2 +
#           12_2 * i2_2 +
#           13_2 * i3_2 +
#           14_2 * i4_2 +
#           15_2 * i5_2
#
# # Adding the covariances
# i1 ~~ i2
# i1_2 ~~ i2_2
#
# i2 ~~ i3
# i2_2 ~~ i3_2
#
# i4 ~~ i5
# i4_2 ~~ i5_2
#
#
#
# # Fixing latent variance to 1, as we freed first factor loading
# math_T2 ~~ 1 * math_T2
#
# # Fixing latent mean to 0 for identification?
# math_T2 ~ 0 * 1
#
# # Correlations across time
# math_T1 ~~ math_T2
# i1 ~~ i1_2
# i2 ~~ i2_2
# i3 ~~ i3_2
# i4 ~~ i4_2
# i5 ~~ i5_2
# '
#
# fit_els_time <- cfa(els_time, data = els,

```

```
#               estimator = "MLR", missing = "FIML", se = "robust.mlr")
# fit_els_time
# head(modindices(fit_els_time, sort. = TRUE, free.remove = FALSE))
#
# s_els_time <- summary(fit_hsls_time, fit.measures = TRUE)
```

## HSLs ONLY WEAK

```
#
# hsls_weak <- '
#   math_T1 =~ 1 * i1 +
#             12 * i2 +
#             14 * i4 +
#             15 * i5
#
#
#   # Freeing latent variance, as we restrict first factor loading instead
#   math_T1 ~~ NA * math_T1
#
#   # Fixing latent mean to 0 for identification?
#   math_T1 ~ 0 * 1
#
#   # Time Point 2
#   math_T2 =~ 1 * i1_2 +
#             12 * i2_2 +
#             14 * i4_2 +
#             15 * i5_2
#
#
#   # Adding the covariances
#   i1 ~~ i2
#   i1_2 ~~ i2_2
#
#   i2 ~~ i4
#   i2_2 ~~ i4_2
#
#   i1 ~~ i5
#   i1_2 ~~ i5_2
#
```

```

# # Freeing latent variance, as we restrict first factor loading
# math_T2 ~~ NA * math_T2
#
# # Fixing latent mean to 0 for identification?
# math_T2 ~ 0 * 1
#
# # Correlations across time
# math_T1 ~~ math_T2
# i1 ~~ i1_2
# i2 ~~ i2_2
# i4 ~~ i4_2
# i5 ~~ i5_2
# '
#
# fit_hsls_weak <- cfa(hsls_weak, data = hsls,
#                       estimator = "MLR", missing = "FIML", se = "robust.mlr")
# fit_hsls_weak
# head(modindices(fit_hsls_weak, sort. = TRUE, free.remove = FALSE))
#
#
# s_hsls_weak <- summary(fit_hsls_weak, fit.measures = TRUE)
#

```