

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

Turning warnings off (bad idea)

```
options(warn = -1)
```

```
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 0.6-19
lavaan is FREE software! Please report any bugs.

```
library(ltm)
```

Loading required package: MASS

Attaching package: 'MASS'

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

```
select
```

Loading required package: msm

Loading required package: polycor

```
library(sjlabelled)
```

Attaching package: 'sjlabelled'

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

```
as_label
```

```
library(kableExtra)
```

Attaching package: 'kableExtra'

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

```
group_rows
```

```
library(sirt)
```

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

```
library(mirt)
```

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")
```

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)`

```
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")

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 +  
              el2_2 * i2_2 +  
              el3_2 * i3_2 +  
              el4_2 * i4_2 +  
              el5_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")
fit_config_comb

```

lavaan 0.6-19 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	Scaled
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	187.143	187.143
HSLs	48.065	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

```

fitmeasures(fit_config_comb, c("aic", "bic", "cfi", "df"))

```

aic	bic	cfi	df
474900.288	475589.784	0.999	28.000

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

	id	lhs	op	rhs	user	block	group	free	ustart	exo	label	plabel
1	1	math_t1	=~	i1	1	1	1	1	NA	0		.p1.
2	2	math_t1	=~	i2	1	1	1	2	NA	0	el2_1	.p2.
3	3	math_t1	=~	i3	1	1	1	3	NA	0	el3_1	.p3.
4	4	math_t1	=~	i4	1	1	1	4	NA	0	el4_1	.p4.
5	5	math_t1	=~	i5	1	1	1	5	NA	0	el5_1	.p5.
6	6	i1	~1		1	1	1	6	NA	0	enu1_1	.p6.
7	7	i2	~1		1	1	1	7	NA	0	enu2_1	.p7.
8	8	i3	~1		1	1	1	8	NA	0	enu3_1	.p8.
9	9	i4	~1		1	1	1	9	NA	0	enu4_1	.p9.
10	10	i5	~1		1	1	1	10	NA	0	enu5_1	.p10.
11	11	i1	~~	i1	1	1	1	11	NA	0	etheta1_1	.p11.
12	12	i2	~~	i2	1	1	1	12	NA	0	etheta2_1	.p12.
13	13	i3	~~	i3	1	1	1	13	NA	0	etheta3_1	.p13.
14	14	i4	~~	i4	1	1	1	14	NA	0	etheta4_1	.p14.
15	15	i5	~~	i5	1	1	1	15	NA	0	etheta5_1	.p15.
16	16	math_t1	~~	math_t1	1	1	1	0	1	0		.p16.
17	17	math_t1	~1		1	1	1	0	0	0		.p17.
18	18	math_t2	=~	i1_2	1	1	1	16	NA	0		.p18.
19	19	math_t2	=~	i2_2	1	1	1	17	NA	0	el2_2	.p19.
20	20	math_t2	=~	i3_2	1	1	1	18	NA	0	el3_2	.p20.
21	21	math_t2	=~	i4_2	1	1	1	19	NA	0	el4_2	.p21.
22	22	math_t2	=~	i5_2	1	1	1	20	NA	0	el5_2	.p22.
23	23	i1_2	~1		1	1	1	21	NA	0	enu1_2	.p23.
24	24	i2_2	~1		1	1	1	22	NA	0	enu2_2	.p24.
25	25	i3_2	~1		1	1	1	23	NA	0	enu3_2	.p25.
26	26	i4_2	~1		1	1	1	24	NA	0	enu4_2	.p26.
27	27	i5_2	~1		1	1	1	25	NA	0	enu5_2	.p27.
28	28	i1_2	~~	i1_2	1	1	1	26	NA	0	etheta1_2	.p28.
29	29	i2_2	~~	i2_2	1	1	1	27	NA	0	etheta2_2	.p29.
30	30	i3_2	~~	i3_2	1	1	1	28	NA	0	etheta3_2	.p30.
31	31	i4_2	~~	i4_2	1	1	1	29	NA	0	etheta4_2	.p31.
32	32	i5_2	~~	i5_2	1	1	1	30	NA	0	etheta5_2	.p32.
33	33	i1	~~	i2	1	1	1	31	NA	0		.p33.
34	34	i1_2	~~	i2_2	1	1	1	32	NA	0		.p34.

35	35	i1	~~	i3	1	1	1	33	NA	0		.p35.
36	36	i1_2	~~	i3_2	1	1	1	34	NA	0		.p36.
37	37	i2	~~	i3	1	1	1	35	NA	0		.p37.
38	38	i2_2	~~	i3_2	1	1	1	36	NA	0		.p38.
39	39	i4	~~	i5	1	1	1	37	NA	0		.p39.
40	40	i4_2	~~	i5_2	1	1	1	38	NA	0		.p40.
41	41	math_t2	~~	math_t2	1	1	1	0	1	0		.p41.
42	42	math_t2	~1		1	1	1	0	0	0		.p42.
43	43	math_t1	~~	math_t2	1	1	1	39	NA	0		.p43.
44	44	i1	~~	i1_2	1	1	1	40	NA	0		.p44.
45	45	i2	~~	i2_2	1	1	1	41	NA	0		.p45.
46	46	i3	~~	i3_2	1	1	1	42	NA	0		.p46.
47	47	i4	~~	i4_2	1	1	1	43	NA	0		.p47.
48	48	i5	~~	i5_2	1	1	1	44	NA	0		.p48.
49	49	math_t1	=~	i1	1	2	2	45	NA	0		.p49.
50	50	math_t1	=~	i2	1	2	2	46	NA	0	hl2_1	.p50.
51	51	math_t1	=~	i4	1	2	2	47	NA	0	hl4_1	.p51.
52	52	math_t1	=~	i5	1	2	2	48	NA	0	hl5_1	.p52.
53	53	i1	~1		1	2	2	49	NA	0	hnu1_1	.p53.
54	54	i2	~1		1	2	2	50	NA	0	hnu2_1	.p54.
55	55	i4	~1		1	2	2	51	NA	0	hnu4_1	.p55.
56	56	i5	~1		1	2	2	52	NA	0	hnu5_1	.p56.
57	57	i1	~~	i1	1	2	2	53	NA	0	htheta1_1	.p57.
58	58	i2	~~	i2	1	2	2	54	NA	0	htheta2_1	.p58.
59	59	i4	~~	i4	1	2	2	55	NA	0	htheta4_1	.p59.
60	60	i5	~~	i5	1	2	2	56	NA	0	htheta5_1	.p60.
61	61	math_t1	~~	math_t1	1	2	2	0	1	0		.p61.
62	62	math_t1	~1		1	2	2	0	0	0		.p62.
63	63	math_t2	=~	i1_2	1	2	2	57	NA	0		.p63.
64	64	math_t2	=~	i2_2	1	2	2	58	NA	0	hl2_2	.p64.
65	65	math_t2	=~	i4_2	1	2	2	59	NA	0	hl4_2	.p65.
66	66	math_t2	=~	i5_2	1	2	2	60	NA	0	hl5_2	.p66.
67	67	i1_2	~1		1	2	2	61	NA	0	hnu1_2	.p67.
68	68	i2_2	~1		1	2	2	62	NA	0	hnu2_2	.p68.
69	69	i4_2	~1		1	2	2	63	NA	0	hnu4_2	.p69.
70	70	i5_2	~1		1	2	2	64	NA	0	hnu5_2	.p70.
71	71	i1_2	~~	i1_2	1	2	2	65	NA	0	htheta1_2	.p71.
72	72	i2_2	~~	i2_2	1	2	2	66	NA	0	htheta2_2	.p72.
73	73	i4_2	~~	i4_2	1	2	2	67	NA	0	htheta4_2	.p73.
74	74	i5_2	~~	i5_2	1	2	2	68	NA	0	htheta5_2	.p74.
75	75	i1	~~	i2	1	2	2	69	NA	0		.p75.
76	76	i1_2	~~	i2_2	1	2	2	70	NA	0		.p76.
77	77	i1	~~	i4	1	2	2	71	NA	0		.p77.

78	78	i1_2	~~	i4_2	1	2	2	72	NA	0	.p78.
79	79	i2	~~	i4	1	2	2	73	NA	0	.p79.
80	80	i2_2	~~	i4_2	1	2	2	74	NA	0	.p80.
81	81	i1	~~	i5	1	2	2	75	NA	0	.p81.
82	82	i1_2	~~	i5_2	1	2	2	76	NA	0	.p82.
83	83	math_t2	~~	math_t2	1	2	2	0	1	0	.p83.
84	84	math_t2	~1		1	2	2	0	0	0	.p84.
85	85	math_t1	~~	math_t2	1	2	2	77	NA	0	.p85.
86	86	i1	~~	i1_2	1	2	2	78	NA	0	.p86.
87	87	i2	~~	i2_2	1	2	2	79	NA	0	.p87.
88	88	i4	~~	i4_2	1	2	2	80	NA	0	.p88.
89	89	i5	~~	i5_2	1	2	2	81	NA	0	.p89.
		start		est							
1		0.791		0.775							
2		0.803		0.759							
3		0.842		0.868							
4		0.798		0.800							
5		0.799		0.792							
6		2.541		2.541							
7		2.357		2.358							
8		2.462		2.462							
9		2.623		2.623							
10		2.650		2.650							
11		0.428		0.261							
12		0.436		0.293							
13		0.464		0.175							
14		0.444		0.248							
15		0.439		0.255							
16		1.000		1.000							
17		0.000		0.000							
18		0.734		0.752							
19		0.745		0.726							
20		0.766		0.752							
21		0.691		0.682							
22		0.760		0.750							
23		2.564		2.565							
24		2.367		2.369							
25		2.444		2.445							
26		2.849		2.850							
27		2.740		2.742							
28		0.394		0.223							
29		0.400		0.271							
30		0.421		0.277							

31	0.375	0.286	0.008
32	0.425	0.290	0.010
33	0.000	0.076	0.009
34	0.000	0.006	0.009
35	0.000	-0.044	0.009
36	0.000	-0.024	0.009
37	0.000	0.017	0.009
38	0.000	0.069	0.009
39	0.000	0.082	0.009
40	0.000	0.049	0.008
41	1.000	1.000	0.000
42	0.000	0.000	0.000
43	0.000	0.532	0.010
44	0.000	0.010	0.004
45	0.000	0.024	0.004
46	0.000	0.003	0.004
47	0.000	0.010	0.004
48	0.000	0.009	0.004
49	0.640	0.654	0.011
50	0.636	0.640	0.007
51	0.613	0.607	0.007
52	0.612	0.624	0.007
53	2.974	2.974	0.005
54	2.720	2.720	0.006
55	3.066	3.066	0.005
56	2.980	2.979	0.005
57	0.287	0.145	0.013
58	0.334	0.256	0.008
59	0.258	0.149	0.007
60	0.266	0.142	0.006
61	1.000	1.000	0.000
62	0.000	0.000	0.000
63	0.730	0.675	0.011
64	0.654	0.686	0.008
65	0.650	0.667	0.007
66	0.650	0.631	0.007
67	2.769	2.769	0.006
68	2.542	2.542	0.006
69	2.912	2.912	0.005
70	2.848	2.848	0.005
71	0.334	0.214	0.014
72	0.394	0.317	0.010
73	0.297	0.150	0.008


```

74 0.303 0.208 0.008
75 0.000 -0.003 0.008
76 0.000 0.010 0.009
77 0.000 -0.001 0.008
78 0.000 0.028 0.008
79 0.000 -0.011 0.006
80 0.000 -0.038 0.008
81 0.000 -0.032 0.007
82 0.000 0.046 0.007
83 1.000 1.000 0.000
84 0.000 0.000 0.000
85 0.000 0.405 0.009
86 0.000 0.007 0.002
87 0.000 0.030 0.003
88 0.000 0.007 0.002
89 0.000 0.013 0.002

```

BETWEEN WEAK ELS + HSLS

```

weak_comb <- '

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

#####
# Time Point 1
#####
math_t1 =~ l1_1 * i1 +
           l2_1 * i2 +
           l3_1 * i3 +
           l4_1 * i4 +
           l5_1 * i5

# Intercepts
i1 ~ 0 * 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

# Free both
math_t1 ~~ var_els_t1 * math_t1
math_t1 ~ mean_els_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1_2 * i1_2 +
           l2_2 * i2_2 +
           l3_2 * i3_2 +
           l4_2 * i4_2 +
           l5_2 * i5_2

# Intercepts
i1_2 ~ 0 * 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 ~~ var_els_t2 * math_t2
math_t2 ~ mean_els_t2 * 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 =~ l1_1 * i1 +
           l2_1 * i2 +   # Same label as ELS
           # no i3 in HSLs
           l4_1 * i4 +   # Same label as ELS
           l5_1 * i5     # Same label as ELS

# Intercepts
i1 ~ 0 * 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_t1 * math_t1
math_t1 ~ mean_hsls_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1_2 * 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
i1_2 ~ 0 * 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 ~ mean_hsls_math_t2 * 1 # free mean

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

```

lavaan 0.6-19 ended normally after 131 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	81	
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	Scaled
Test Statistic	368.014	323.188
Degrees of freedom	34	34
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.139
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	228.778	228.778
HSLs	94.411	94.411

```
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	72.712	-0.102	-0.076	-0.079	-0.079
141	math_t1	=~	i5_2	2	HSLs	1	39.492	-0.049	-0.032	-0.041	-0.041
107	i1	~~	i5	1	ELS	1	35.681	-0.019	-0.019	-0.075	-0.075
97	math_t1	=~	i2_2	1	ELS	1	30.437	0.048	0.038	0.043	0.043
115	i2	~~	i3_2	1	ELS	1	29.842	0.021	0.021	0.079	0.079
139	math_t1	=~	i2_2	2	HSLs	1	28.588	0.048	0.031	0.035	0.035

```
s_weak_comb <- summary(fit_weak_comb, fit.measures = TRUE, standardized = TRUE)
fitMeasures(fit_weak_comb, c("rmsea", "chisq.scaled", "cfi", "tli", "df"))
```

rmsea	chisq.scaled	cfi	tli	df
0.023	323.188	0.998	0.996	34.000

```
#s_weak_comb
#lavTestLRT(fit_config_comb, fit_weak_comb)
```

BETWEEN STRONG ELS + HSLs

```
strong_between <- '
```

```
#####
##                               ELS (Group 1)                               ##
#####
```

group: ELS

```
#####
# Time Point 1
#####
math_t1 =~ l1_1      * i1 +
            l2_1 * i2 +
            l3_1 * i3 +
            l4_1 * i4 +
            l5_1 * i5

# Intercepts
# Same labels for both groups!
i1 ~ 0 * 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

# Free both
math_t1 ~~ var_els_t1 * math_t1
math_t1 ~ mean_els_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1_2      * i1_2 +
            l2_2 * i2_2 +
            l3_2 * i3_2 +
            l4_2 * i4_2 +
            l5_2 * i5_2

# Intercepts
```

```

# Same labels for both groups!
i1_2 ~ 0 * 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 ~ mean_els_t2 * 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 =~ l1_1    * i1 +
           l2_1 * i2 +    # Same label as ELS
           # no i3 in HSLs
           l4_1 * i4 +    # Same label as ELS
           l5_1 * i5      # Same label as ELS

# Intercepts
# Same labels for both groups!
i1 ~ 0 * 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 both
math_t1 ~~ var_hsls_t1 * math_t1
math_t1 ~ mean_hsls_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1_2    * 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 ~ 0 * 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 ~ mean_hsls_t2 * 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_between <- sem(strong_between, data = dat, group = "sample",
                           estimator = "MLR", missing = "FIML", se = "robust.mlr")
fit_strong_between

```

lavaan 0.6-19 ended normally after 112 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	81
Number of equality constraints	12

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	Scaled
Test Statistic	979.397	878.847
Degrees of freedom	40	40
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.114
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	612.296	612.296
HSLs	266.550	266.550

```

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

```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
102	math_t1	=~	i1_2	1	ELS	1	176.902	-0.033	-0.027	-0.030	-0.030
111	math_t2	=~	i5	1	ELS	1	170.928	0.030	0.023	0.025	0.025
67	i1_2	~1		2	HSLs	1	156.308	0.085	0.085	0.103	0.103
23	i1_2	~1		1	ELS	1	156.308	-0.085	-0.085	-0.095	-0.095
110	math_t2	=~	i4	1	ELS	1	149.746	-0.028	-0.021	-0.022	-0.022
63	math_t2	=~	i1_2	2	HSLs	1	149.526	0.030	0.021	0.026	0.026

```
s_strong_between <- summary(fit_strong_between, fit.measures = TRUE, standardized = TRUE)
#s_strong_between
fitMeasures(fit_strong_between, c("rmsea", "chisq.scaled", "cfi", "tli", "df"))
```

rmsea	chisq.scaled	cfi	tli	df
0.036	878.847	0.995	0.990	40.000

```
fit_weak_comb
```

lavaan 0.6-19 ended normally after 131 iterations

Estimator	ML
Optimization method	NLMINB
Number of model parameters	81
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	Scaled
Test Statistic	368.014	323.188
Degrees of freedom	34	34
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.139
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	228.778	228.778
HSLs	94.411	94.411

```
#lavTestLRT(fit_weak_comb, fit_strong_between)
```

WITHIN STRONG ELS+HSLS

```
strong_within <- '

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

#####
# Time Point 1
#####
math_t1 =~ e11      * i1 +
            e12 * i2 +
            e13 * i3 +
            e14 * i4 +
            e15 * i5

# Intercepts
# Same labels for both times!
i1 ~ 0 * 1
i2 ~ nu2_2 * 1
i3 ~ nu3_3 * 1
i4 ~ nu4_4 * 1
i5 ~ nu5_5 * 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 ~ mean_els_t1 * 1

#####
# Time Point 2
#####
```

```

math_t2 =~ e11 * i1_2 +
           e12 * i2_2 +
           e13 * i3_2 +
           e14 * i4_2 +
           e15 * i5_2

# Intercepts
# Same labels for both groups!
i1_2 ~ 0 * 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 ~ mean_els_t2 * 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 =~ h11    * i1 +
            h12 * i2 + # Same label as ELS
            # no i3 in HSLs
            h14 * i4 + # Same label as ELS
            h15 * i5   # Same label as ELS

# Intercepts
# Same labels for both times!
i1 ~ 0 * 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 ~ mean_hsls_t1 * 1

#####
# Time Point 2

```

```
#####
math_t2 =~ h11    * i1_2 +
           h12 * i2_2 +   # Same label as ELS
           # no i3_2 in HSLS at Time 2
           h14 * i4_2 +   # Same label as ELS
           h15 * i5_2     # Same label as ELS

# Intercepts
# Same labels for both times!

i1_2 ~ 0 * 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 ~ mean_hsls_t2 * 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_within <- cfa(strong_within, data = dat, group = "sample",
  estimator = "MLR", missing = "FIML", se = "robust.mlr",
  std.lv = TRUE)

fit_strong_within

```

lavaan 0.6-19 ended normally after 123 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	81	
Number of equality constraints	10	
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	Scaled
Test Statistic	571.360	515.496
Degrees of freedom	38	38
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.108
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	416.217	416.217
HSLs	99.278	99.278

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

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all
16	math_t1	~~	math_t1	1	ELS	1	90.352	0.183	1.000	1.000
41	math_t2	~~	math_t2	1	ELS	1	90.352	-0.183	-1.000	-1.000
103	math_t1	=~	i4_2	1	ELS	1	52.783	-0.053	-0.053	-0.058
111	i1	~~	i5	1	ELS	1	49.761	-0.022	-0.022	-0.088
102	math_t1	=~	i3_2	1	ELS	1	38.385	-0.045	-0.045	-0.047
61	math_t1	~~	math_t1	2	HSLs	1	38.010	-0.097	-1.000	-1.000
	sepc.nox									
16			1.000							
41			-1.000							
103			-0.058							
111			-0.088							
102			-0.047							
61			-1.000							

```
s_strong_within <- summary(fit_strong_within, fit.measures = TRUE, standardized = TRUE)
#s_strong_within
fitMeasures(fit_strong_within, c("rmsea", "chisq.scaled", "cfi", "tli", "df"))
```

rmsea	chisq.scaled	cfi	tli	df
0.028	515.496	0.997	0.994	38.000

WITHIN+BETWEEN WEAK

```
# Latent variance and mean set to 1 and 0 in ELS time 1,
# free variance, mean set to 0 everywhere else

both_weak_comb <- '

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

#####
# Time Point 1
#####
math_t1 =~ l1 * i1 +
           l2 * i2 +
           l3 * i3 +
```

```

14 * i4 +
15 * i5

# Intercepts
i1 ~ 0 * 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

# Free both
math_t1 ~~ var_els_t1 * math_t1
math_t1 ~ mean_els_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1 * i1_2 +
           12 * i2_2 +
           13 * i3_2 +
           14 * i4_2 +
           15 * i5_2

# Intercepts
i1_2 ~ 0 * 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 ~~ i5
i1_2 ~~ i5_2

i1 ~~ i3
i1_2 ~~ i3_2

i1 ~~ i4
i1_2 ~~ i4_2

i2 ~~ i3
i2_2 ~~ i3_2

i4 ~~ i5
i4_2 ~~ i5_2

# Free both
math_t2 ~~ var_els_t2 * math_t2
math_t2 ~ mean_els_t2 * 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 =~ l1 * i1 +
           12 * i2 +   # Same label as ELS
           # no i3 in HSLS
           14 * i4 +   # Same label as ELS
           15 * i5     # Same label as ELS

# Intercepts
i1 ~ 0 * 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 both
math_t1 ~~ var_hsls_t1 * math_t1
math_t1 ~ mean_hsls_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1 * i1_2 +
           12 * i2_2 +   # Same label as ELS
           # no i3_2 in HSLS at Time 2
           14 * i4_2 +   # Same label as ELS
           15 * i5_2     # Same label as ELS

# Intercepts
i1_2 ~ 0 * 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 ~~ i5
i1_2 ~~ i5_2

# i1 ~~ i3
# i1_2 ~~ i3_2

i1 ~~ i4
i1_2 ~~ i4_2

i4 ~~ i5
i4_2 ~~ i5_2

# Free latent variance and free latent mean
math_t2 ~~ var_hsls_t2 * math_t2
math_t2 ~ mean_hsls_t2 * 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_both_weak_comb <- sem(both_weak_comb, data = dat, group = "sample",
                          estimator = "MLR", missing = "FIML", se = "robust.mlr")
fit_both_weak_comb

```

lavaan 0.6-19 ended normally after 112 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	81	
Number of equality constraints	10	
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	Scaled
Test Statistic	402.536	368.810
Degrees of freedom	38	38
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.091
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	295.386	295.386
HSLs	73.424	73.424

```
fitMeasures(fit_both_weak_comb, c("rmsea", "chisq.scaled", "cfi", "tli", "df", "aic", "bic"))
```

rmsea	chisq.scaled	cfi	tli	df	aic
0.023	368.810	0.998	0.996	38.000	475008.011
bic					
475612.385					

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

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
108	math_t2	=~	i4	1	ELS	1	50.106	0.059	0.043	0.047	0.047
104	math_t1	=~	i5_2	1	ELS	1	40.568	0.050	0.042	0.046	0.046
101	math_t1	=~	i2_2	1	ELS	1	35.772	0.046	0.038	0.044	0.044
112	i1	~~	i3_2	1	ELS	1	34.023	-0.021	-0.021	-0.103	-0.103
102	math_t1	=~	i3_2	1	ELS	1	33.877	-0.051	-0.042	-0.045	-0.045
121	i3	~~	i4	1	ELS	1	24.866	0.017	0.017	0.088	0.088

```
s_both_weak_comb <- summary(fit_both_weak_comb, fit.measures = TRUE, standardized = TRUE)
#s_both_weak_comb
```

WITHIN+BETWEEN STRONG

```
both_strong_comb <- '

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

#####
# Time Point 1
#####
math_t1 =~ l1 * i1 +
           12 * i2 +
           13 * i3 +
           14 * i4 +
           15 * i5

#####
# Equal Intercepts #
#####
i1 ~ 0 * 1
i2 ~ nu2 * 1
i3 ~ nu3 * 1
i4 ~ nu4 * 1
i5 ~ nu5 * 1

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

# Free latent mean and variance
math_t1 ~~ var_els_t1 * math_t1
```



```
math_t1 ~ mean_els_t1 * 1
```

```
#####
```

```
# Time Point 2
```

```
#####
```

```
math_t2 =~ l1 * i1_2 +  
           l2 * i2_2 +  
           l3 * i3_2 +  
           l4 * i4_2 +  
           l5 * i5_2
```

```
#####
```

```
# Equal Intercepts #
```

```
#####
```

```
# Intercepts
```

```
i1_2 ~ 0 * 1  
i2_2 ~ nu2 * 1  
i3_2 ~ nu3 * 1  
i4_2 ~ nu4 * 1  
i5_2 ~ nu5 * 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 ~~ i5  
i1_2 ~~ i5_2
```

```
i1 ~~ i3  
i1_2 ~~ i3_2
```

```
i1 ~~ i4  
i1_2 ~~ i4_2
```

```
i2 ~~ i3
```

```

i2_2 ~~ i3_2

i4 ~~ i5
i4_2 ~~ i5_2

# Free latent variance, free mean
math_t2 ~~ var_els_t2 * math_t2
math_t2 ~ mean_els_t2 * 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 =~ l1 * i1 +
           12 * i2 +   # Same label as ELS
           # no i3 in HSLs
           14 * i4 +   # Same label as ELS
           15 * i5     # Same label as ELS

#####
# Equal Intercepts #
#####

i1 ~ 0 * 1
i2 ~ nu2 * 1
#i3 ~ nu3 * 1 (item not in HSLs)
i4 ~ nu4 * 1
i5 ~ nu5 * 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 mean
math_t1 ~~ var_hsls_t1 * math_t1
math_t1 ~ mean_hsls_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1 * i1_2 +
           12 * i2_2 +   # Same label as ELS
           # no i3_2 in HSLS at Time 2
           14 * i4_2 +   # Same label as ELS
           15 * i5_2     # Same label as ELS

#####
# Equal Intercepts #
#####

i1_2 ~ 0 * 1
i2_2 ~ nu2 * 1
#i3_2 ~ nu3 * 1 (item not in HSLS)
i4_2 ~ nu4 * 1
i5_2 ~ nu5 * 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 ~~ i5
i1_2 ~~ i5_2

# i1 ~~ i3
# i1_2 ~~ i3_2

i1 ~~ i4
i1_2 ~~ i4_2

i4 ~~ i5
i4_2 ~~ i5_2

# Free latent variance
math_t2 ~~ var_hsls_t2 * math_t2
math_t2 ~ mean_hsls_t2 * 1 #free mean

# 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_both_strong_comb <-
  sem(both_strong_comb, data = dat, group = "sample",
      estimator = "MLR", missing = "FIML", se = "robust.mlr")

fit_both_strong_comb

```

lavaan 0.6-19 ended normally after 97 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	81	
Number of equality constraints	20	
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	Scaled
Test Statistic	1542.603	1436.560
Degrees of freedom	48	48
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.074
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	1191.896	1191.896
HSLs	244.665	244.665

```
fitMeasures(fit_both_strong_comb, c("rmsea", "chisq.scaled", "cfi", "tli", "df", "aic", "b
```

rmsea	chisq.scaled	cfi	tli	df	aic
0.041	1436.560	0.992	0.987	48.000	476128.078
bic					
476647.328					

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

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
9	i4	~1		1	ELS	1	378.675	-0.095	-0.095	-0.102	-0.102
118	math_t2	=~	i4	1	ELS	1	375.201	-0.039	-0.029	-0.031	-0.031
4	math_t1	=~	i4	1	ELS	1	306.319	-0.032	-0.027	-0.029	-0.029
26	i4_2	~1		1	ELS	1	299.280	0.097	0.097	0.109	0.109
113	math_t1	=~	i4_2	1	ELS	1	297.412	0.039	0.032	0.037	0.037
21	math_t2	=~	i4_2	1	ELS	1	239.919	0.032	0.024	0.027	0.027

```
s_both_strong_comb <- summary(fit_both_strong_comb, fit.measures = TRUE, standardized = TR
#s_both_strong_comb
```

W/B Partial Strong

```
both_strong_partial_ei4 <- '

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

#####
# Time Point 1
#####
math_t1 =~ l1 * i1 +
           l2 * i2 +
           l3 * i3 +
           l4 * i4 +
           l5 * i5

#####
# Equal Intercepts #
#####
i1 ~ 0 * 1
i2 ~ nu2 * 1
i3 ~ nu3 * 1

# Unique i4 intercept
i4 ~ nu4_1 * 1

i5 ~ nu5 * 1

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

# Free latent mean and variance
math_t1 ~~ var_els_t1 * math_t1
math_t1 ~ mean_els_t1 * 1
```

```
#####
# Time Point 2
#####
math_t2 =~ l1 * i1_2 +
            l2 * i2_2 +
            l3 * i3_2 +
            l4 * i4_2 +
            l5 * i5_2

#####
# Equal Intercepts #
#####
# Intercepts
i1_2 ~ 0 * 1
i2_2 ~ nu2 * 1
i3_2 ~ nu3 * 1
i4_2 ~ nu4 * 1
i5_2 ~ nu5 * 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 ~~ i5
i1_2 ~~ i5_2

i1 ~~ i3
i1_2 ~~ i3_2

i1 ~~ i4
i1_2 ~~ i4_2

i2 ~~ i3
i2_2 ~~ i3_2

i4 ~~ i5
```

```

i4_2 ~~ i5_2

# Free latent variance, free mean
math_t2 ~~ var_els_t2 * math_t2
math_t2 ~ mean_els_t2 * 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 =~ l1 * i1 +
           l2 * i2 +   # Same label as ELS
           # no i3 in HSLs
           l4 * i4 +   # Same label as ELS
           l5 * i5     # Same label as ELS

#####
# Equal Intercepts #
#####
i1 ~ 0 * 1
i2 ~ nu2 * 1
#i3 ~ nu3 * 1 (item not in HSLs)
i4 ~ nu4 * 1
i5 ~ nu5 * 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 mean
math_t1 ~~ var_hsls_t1 * math_t1
math_t1 ~ mean_hsls_t1 * 1

#####
# Time Point 2
#####
math_t2 =~ l1 * i1_2 +
           12 * i2_2 +   # Same label as ELS
           # no i3_2 in HSLS at Time 2
           14 * i4_2 +   # Same label as ELS
           15 * i5_2     # Same label as ELS

#####
# Equal Intercepts #
#####

i1_2 ~ 0 * 1
i2_2 ~ nu2 * 1
#i3_2 ~ nu3 * 1 (item not in HSLS)
i4_2 ~ nu4 * 1
i5_2 ~ nu5 * 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 ~~ i5
i1_2 ~~ i5_2

# i1 ~~ i3

```

```

# i1_2 ~~ i3_2

i1 ~~ i4
i1_2 ~~ i4_2

i4 ~~ i5
i4_2 ~~ i5_2

# Free latent variance
math_t2 ~~ var_hsls_t2 * math_t2
math_t2 ~ mean_hsls_t2 * 1 #free mean

# 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_both_strong_partial_ei4 <- sem(both_strong_partial_ei4, data = dat, group = "sample",
                                estimator = "MLR", missing = "FIML", se = "robust.mlr")
fit_both_strong_partial_ei4

```

lavaan 0.6-19 ended normally after 99 iterations

Estimator	ML	
Optimization method	NLMINB	
Number of model parameters	81	
Number of equality constraints	19	
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	Scaled
----------	--------

Test Statistic	1049.413	977.026
Degrees of freedom	47	47
P-value (Chi-square)	0.000	0.000
Scaling correction factor		1.074
Yuan-Bentler correction (Mplus variant)		
Test statistic for each group:		
ELS	731.286	731.286
HSLs	245.739	245.739

```
fitMeasures(fit_both_strong_partial_ei4, c("rmsea", "chisq.scaled", "cfi", "tli", "df", "a
```

rmsea	chisq.scaled	cfi	tli	df	aic
0.034	977.026	0.994	0.991	47.000	475636.889
bic					
476164.652					

```
head(modindices(fit_both_strong_partial_ei4, sort. = TRUE, free.remove = FALSE))
```

	lhs	op	rhs	block	group	level	mi	epc	sepc.lv	sepc.all	sepc.nox
55	i1	~1		2	HSLs	1	203.047	0.063	0.063	0.083	0.083
153	math_t2	=~	i1	2	HSLs	1	181.936	0.021	0.015	0.019	0.019
51	math_t1	=~	i1	2	HSLs	1	180.790	0.020	0.013	0.018	0.018
112	math_t1	=~	i4_2	1	ELS	1	175.604	0.030	0.025	0.029	0.029
109	math_t1	=~	i1_2	1	ELS	1	170.958	-0.029	-0.025	-0.027	-0.027
26	i4_2	~1		1	ELS	1	169.082	0.072	0.072	0.082	0.082

```
s_both_strong_comb <- summary(fit_both_strong_partial_ei4, fit.measures = TRUE, standardiz
```

HSLs ONLY

```
# hs1s_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 =~ l1_1 * NA * i1 +
#               l2_1 * i2 +
#               l3_1 * i3 +
#               l4_1 * i4 +
#               l5_1 * i5
#
#
#   # Fixing latent variance to 1, as we freed first factor loading
#   math_T1 ~~ NA * math_T1
#
#   # Fixing latent mean to 0 for identification?
#   math_T1 ~ 0 * 1
#
#   # Time Point 2
#   math_T2 =~ NA * i1_2 +
#               l2_2 * i2_2 +
#               l3_2 * i3_2 +
#               l4_2 * i4_2 +
#               l5_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 ~~ 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
# 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_els_time, fit.measures = TRUE, standardized = TRUE)

```

K-fold validation

```
library(caret)
```

Loading required package: ggplot2

Attaching package: 'ggplot2'

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

```
as_label
```

```

k <- 10 # Number of folds
set.seed(42)
folds <- createFolds(dat$sample, k = k, list = TRUE)

```

```

cfi_list <- numeric(k)
rmsea_list <- numeric(k)
srmr_list <- numeric(k)
chisq_list <- numeric(k)

for (i in seq_along(folds)) {
  # Split into training and validation
  validation_indices <- folds[[i]]
  train_data <- dat[-validation_indices, ]
  validation_data <- dat[validation_indices, ]

  # Fit model on training set
  fit_train <- fit_train <- sem(both_strong_partial_ei4, data = train_data,
                                group = "sample",
                                estimator = "MLR",
                                missing = "FIML",
                                se = "robust.mlr")

  # Fit model on validation set using training constraints
  fit_validation <- fit_validation <- sem(both_strong_partial_ei4,
                                           data = validation_data,
                                           group = "sample",
                                           estimator = "MLR",
                                           missing = "FIML", se = "robust.mlr")

  # Extract fit measures for the validation set
  cfi_list[i] <- fitMeasures(fit_validation, "cfi")
  rmsea_list[i] <- fitMeasures(fit_validation, "rmsea")
  srmr_list[i] <- fitMeasures(fit_validation, "srmr")
  chisq_list[i] <- fitMeasures(fit_validation, "chisq.scaled")
}

fit_measures_summary <- data.frame(
  Fold = seq_len(k),
  CFI = cfi_list,
  RMSEA = rmsea_list,
  SRMR = srmr_list,
  chisquare = chisq_list
)

fit_measures_summary <- rbind(

```

```

fit_measures_summary,
data.frame(
  Fold = "Mean",
  CFI = mean(cfi_list),
  RMSEA = mean(rmsea_list),
  SRMR = mean(srmr_list),
  chisquare = mean(chisq_list)
),
data.frame(
  Fold = "SD",
  CFI = sd(cfi_list),
  RMSEA = sd(rmsea_list),
  SRMR = sd(srmr_list),
  chisquare = sd(chisq_list)
)
)

# Display fit measures summary
print(fit_measures_summary)

```

	Fold	CFI	RMSEA	SRMR	chisquare
1	1	0.9934423414	0.036906512	0.023831975	154.28390
2	2	0.9940840973	0.035169185	0.022386103	142.96924
3	3	0.9947781295	0.032514042	0.022549146	126.08007
4	4	0.9927581038	0.038744826	0.023552413	161.83597
5	5	0.9947909686	0.033259938	0.021451555	135.07740
6	6	0.9958110729	0.029014786	0.019990899	113.89821
7	7	0.9949440057	0.032159835	0.021995136	130.04654
8	8	0.9930883801	0.037331820	0.024971764	158.72218
9	9	0.9941189973	0.034335904	0.023643486	139.13529
10	10	0.9943338072	0.034375132	0.022724958	141.25962
11	Mean	0.9942149904	0.034381198	0.022709744	140.33084
12	SD	0.0009283821	0.002852367	0.001397406	15.05467

```

k <- 10 # Number of folds
set.seed(42)
folds_2 <- createFolds(dat$sample, k = k, list = TRUE)

cfi_list_2 <- numeric(k)
rmsea_list_2 <- numeric(k)
srmr_list_2 <- numeric(k)

```



```

chisq_list_2 <- numeric(k)

for (i in seq_along(folds_2)) {
  # Split into training and validation
  validation_indices_2 <- folds_2[[i]]
  train_data_2 <- dat[-validation_indices_2, ]
  validation_data_2 <- dat[validation_indices_2
                           , ]

  # Fit model on training set
  fit_train_2 <- sem(both_strong_comb, data = train_data,
                    group = "sample",
                    estimator = "MLR",
                    missing = "FIML",
                    se = "robust.mlr")

  # Fit model on validation set using training constraints
  fit_validation_2 <- sem(both_strong_comb,
                        data = validation_data_2,
                        group = "sample",
                        estimator = "MLR",
                        missing = "FIML", se = "robust.mlr")

  # Extract fit measures for the validation set
  cfi_list_2[i] <- fitMeasures(fit_validation_2, "cfi")
  rmsea_list_2[i] <- fitMeasures(fit_validation_2, "rmsea")
  srmr_list_2[i] <- fitMeasures(fit_validation_2, "srmr")
  chisq_list_2[i] <- fitMeasures(fit_validation_2, 'chisq.scaled')
}

fit_measures_summary_2 <- data.frame(
  Fold = seq_len(k),
  CFI = cfi_list_2,
  RMSEA = rmsea_list_2,
  SRMR = srmr_list_2,
  chisquare = chisq_list_2
)

# Calculate mean and standard deviation of fit measures across folds
fit_measures_summary_2 <- rbind(
  fit_measures_summary_2,

```

```

data.frame(
  Fold = "Mean",
  CFI = mean(cfi_list_2),
  RMSEA = mean(rmse_list_2),
  SRMR = mean(srmr_list_2),
  chisquare = mean(chisq_list_2)
),
data.frame(
  Fold = "SD",
  CFI = sd(cfi_list_2),
  RMSEA = sd(rmse_list_2),
  SRMR = sd(srmr_list_2),
  chisquare = sd(chisq_list_2)
)
)

# Display fit measures summary
print(fit_measures_summary_2)

```

	Fold	CFI	RMSEA	SRMR	chisquare
1	1	0.990991231	0.042804521	0.024319173	196.07829
2	2	0.991997652	0.040475182	0.023497891	179.29671
3	3	0.992362988	0.038908811	0.023790774	165.49312
4	4	0.989604868	0.045933643	0.025101776	214.26489
5	5	0.991708560	0.041522764	0.023012711	189.75043
6	6	0.993863180	0.034751052	0.021054695	147.09210
7	7	0.991884270	0.040318338	0.022852693	182.78645
8	8	0.990979591	0.042201770	0.026034085	194.37595
9	9	0.990069483	0.044150645	0.025314445	205.96086
10	10	0.991194644	0.042403381	0.024350190	195.62530
11	Mean	0.991465647	0.041347011	0.023932843	187.07241
12	SD	0.001202646	0.003058123	0.001434509	19.58996