Structural Equation Modeling II

Lecture 11

Multivariate statistics

Psychology 613 – Spring 2022

Featuring content by Allison

Tackman

What kinds of models can you test with this method?

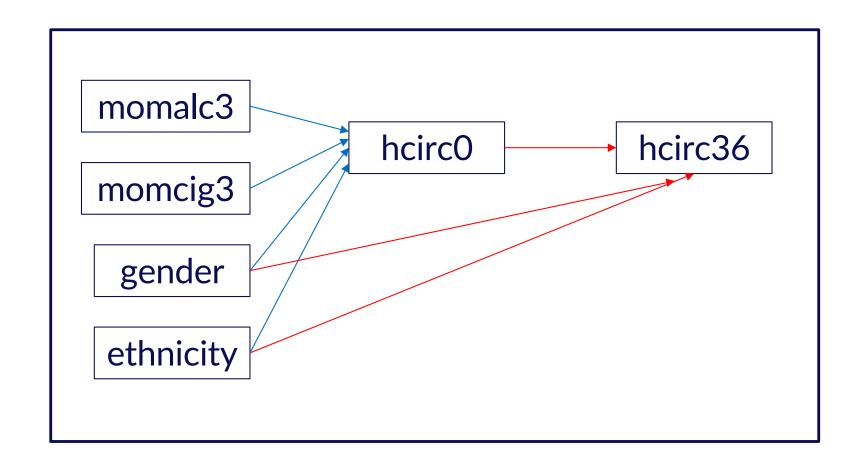
Featuring content by Allison Tackman

What kinds of things can I do with SEM?

- EFA & CFA + measurement invariance
- Path analysis (moderation & mediation)
- Longitudinal data analyses using latent factors
- And so much more...

What kinds of things can I do with SEM?

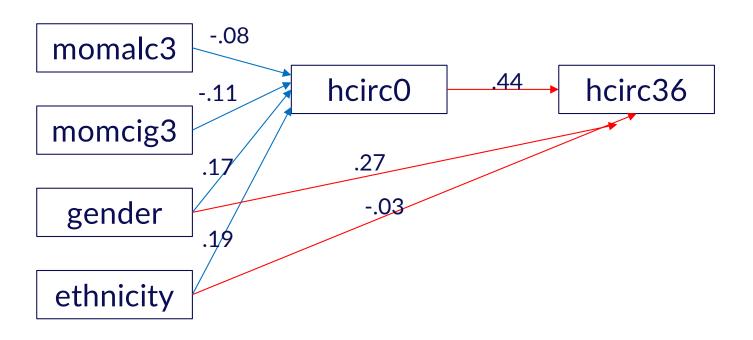
- EFA & CFA + measurement invariance
- Path analysis (moderation & mediation)
- Longitudinal data analyses using latent factors
- And so much more...



Model:

hcirc36 ON hcirc0 gender ethnicity; hcirc0 ON momalc3 momcig3 gender ethnicity;

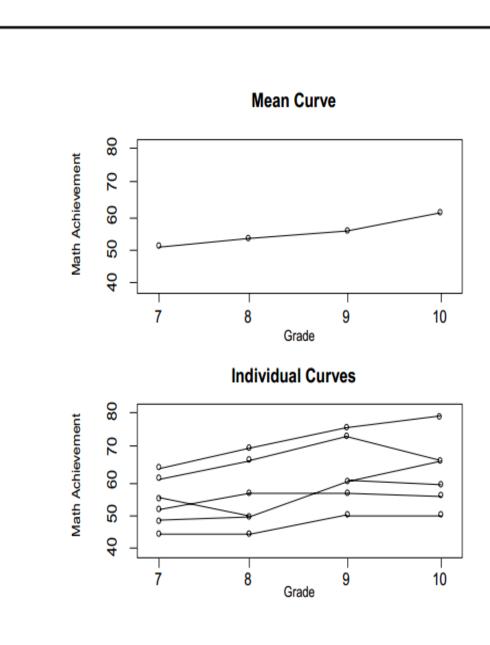
Model Results					
	Estimates	S.E.	Est./S.E.	Std	StdYX
HCIRC36 ON					
HCIRC0	.415	.036	11.382	.415	.439
GENDER	.762	.107	7.146	.762	.270
ETH	094	.107	879	094	033
HCIRCO ON					
MOMALC3	500	.239	-2.090	500	084
MOMCIG3	013	.005	-2.604	013	108
GENDER	.495	.118	4.185	.495	.166
ETH	.578	.125	4.625	.578	.194

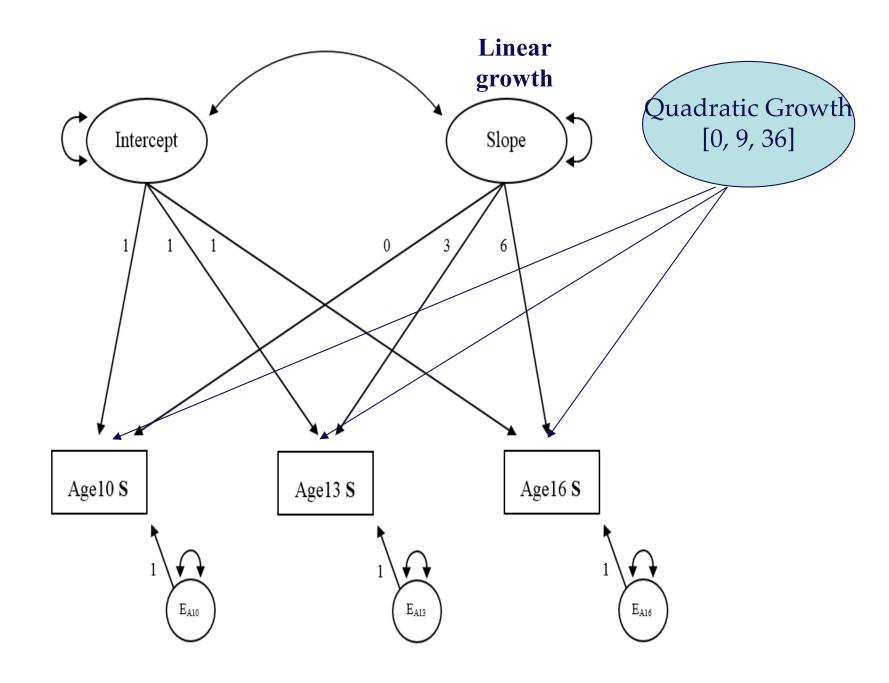


What kinds of things can I do with SEM?

- EFA & CFA + measurement invariance
- Path analysis (moderation & mediation)
- Longitudinal data analyses using latent factors
- And so much more...

Longitudinal data analyses: Univariate latent growth curve modeling





Level 1: $y_{ti} = \eta_{0i} + \eta_{1i} x_t + \varepsilon_{ti}$

Level 2:
$$\eta_{0i} = \alpha_0 + \zeta_{0i}$$

 $\eta_{1i} = \alpha_1 + \zeta_{1i}$

```
M
plu/7
File Edit View Mplus Plot Diagram Window Help
 Title: Linear Latent Growth Curve Model Sleep;
  Data: File is SA ALL REL DATA.csv;
  Variable: Names are gender, tlepsiC, t2epsiC, t3epsiC, T1gradef, T2gradef, T3gradef,
  T1hwf, T2hwf, T3hwf, T1SchEng, T2SchEng, T3SchEng, t1carsch, t2carsch, t3carsch,
  t1parsch, t2parsch, t3parsch, t1sleep, t2sleep, t3sleep, T1PhysA, T2PhysA, T3PhysA,
  t1cesd, t2cesd, t3cesd, t1prsup, t2prsup, t3prsup, t1prdel, t2prdel, t3prdel,
  t1pm, t2pm, t3pm, t1dacc, t2dacc, t3dacc, t1macc, t2macc, t3macc;
  Missing are all (-99999);
  Usevariables are t1sleep t2sleep t3sleep;
  Analysis: Type=general;
  Model: i s | t1sleep@0 t2sleep@3 t3sleep@6;
  Output: sampstat; stdyx;
```

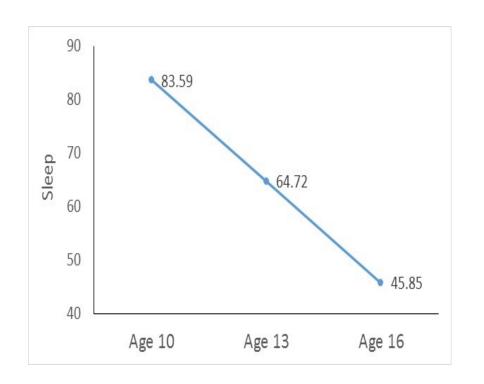
MODEL RESULTS				
				Two-Tailed
	Estimate	S.E.	Est./S.E.	P-Value
I I				
T1SLEEP	1.000	0.000	999.000	999.000
T2SLEEP	1.000	0.000	999.000	999.000
T3SLEEP	1.000	0.000	999.000	999.000
ISSLEEP	1.000	0.000	999.000	999.000
S I				
T1SLEEP	0.000	0.000	999.000	999.000
T2SLEEP	3.000	0.000	999.000	999.000
T3SLEEP	6.000	0.000	999.000	999.000
C MITTH				
S WITH	-22.567	12.864	-1.754	0.079
1	-22.56/	12.004	-1./54	0.079
Means				
I	83.591	1.370	61.010	0.000
S	-6.293	0.493	-12.758	0.000
Intercepts				
T1SLEEP	0.000	0.000	999.000	999.000
T2SLEEP	0.000	0.000	999.000	999.000
T3SLEEP	0.000	0.000	999.000	999.000
Variances				
I	142.292	57.624	2.469	0.014
S	10.035	3.926	2.556	0.011
Residual Variances				
T1SLEEP	19.593	53.717	0.365	0.715
T2SLEEP	158.406	38.025	4.166	0.000
T3SLEEP	82.021	74.761	1.097	0.273

MODEL RESULTS				
	Estimate	S.E.	Est./S.E.	Two-Tailed P-Value
I				
T1SLEEP	1.000	0.000	999.000	999.000
T2SLEEP	1.000	0.000	999.000	999.000
T3SLEEP	1.000	0.000	999.000	999.000
S				
T1SLEEP	0.000	0.000	999.000	999.000
T2SLEEP	3.000	0.000	999.000	999.000
T3SLEEP	6.000	0.000	999.000	999.000
S WITH				
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Intercepts				
T1SLEEP	0.000	0.000	999.000	999.000
T2SLEEP	0.000	0.000	999.000	999.000
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T3SLEEP	82.021	74.761	1.097	0.273

Pred Value for Age 10 (where Int = 1, and Slo = 0): 83.59(1) + (-6.29)(0) = 83.59

Pred Value for Age 10 (where Int = 1, and Slo = 3): 83.59(1) + (-6.29)(3) = 64.72

Pred Value for Age 10 (where Int = 1, and Slo = 6): 83.59(1) + (-6.29)(6) = 45.85



Longitudinal data analyses: Univariate latent growth curve modeling with time-invariant covariates

```
III File Edit View Mplus Plot Diagram Window Help
                  Title: Linear Latent Growth Curve Model Sleep;
 Data: File is SA ALL REL DATA.csv;
 Variable: Names are gender, tlepsiC, t2epsiC, t3epsiC, T1gradef, T2gradef, T3gradef,
 T1hwf, T2hwf, T3hwf, T1SchEng, T2SchEng, T3SchEng, t1carsch, t2carsch, t3carsch,
 t1parsch, t2parsch, t3parsch, t1sleep, t2sleep, t3sleep, T1PhysA, T2PhysA, T3PhysA,
 t1cesd, t2cesd, t3cesd, t1prsup, t2prsup, t3prsup, t1prdel, t2prdel, t3prdel,
 t1pm, t2pm, t3pm, t1dacc, t2dacc, t3dacc, t1macc, t2macc, t3macc;
 Missing are all (-99999);
 Usevariables are t1sleep t2sleep t3sleep gencc;
 Define:
 if (gender==1) THEN gencc = 0;
 if (gender==2) THEN gencc = 1;
 ! Boys are 0
  ! Girls are 1
 Analysis: Type=general;
 Model: i s | t1sleep@0 t2sleep@3 t3sleep@6;
        i s ON gencc;
 Output: sampstat; stdyx;
```

MODEL RESULTS					
					Two-Tailed
		Estimate	S.E.	Est./S.E.	P-Value
		LSCIMACE	3.2.	LSC./S.L.	r-varue
I	1				
_	T1SLEEP	1.000	0.000	999.000	999.000
	T2SLEEP	1.000	0.000	999.000	999.000
	T3SLEEP	1.000	0.000	999.000	999.000
		21000		2221000	222122
S	1				
	T1SLEEP	0.000	0.000	999.000	999.000
	T2SLEEP	3.000	0.000	999.000	999.000
	T3SLEEP	6.000	0.000	999.000	999.000
I	ON				
	GENCC	5.166	2.642	1.955	0.051
s	ON				
	GENCC	-1.655	0.963	-1.719	0.086
,					
S	WITH	22 272	10 746	1 070	0.061
	I	-23.870	12.746	-1.873	0.061
Tn	tercepts				
	T1SLEEP	0.000	0.000	999.000	999.000
	T2SLEEP	0.000	0.000	999.000	999.000
	T3SLEEP	0.000	0.000	999.000	999.000
	I	80.919	1.906	42.448	0.000
	S	-5.377	0.737	-7.297	0.000
		3.377	0.707	,,23,	0.000
Residual Variances					
	T1SLEEP	3.183	53.594	0.059	0.953
	T2SLEEP	159.766	38.415	4.159	0.000
	T3SLEEP	94.112	76.438	1.231	0.218
	I	151.985	58.174	2.613	0.009
	S	9.829	3.757	2.616	0.009

MOD	EL RESULTS				
		Estimate	S.E.	Est./S.E.	Two-Tailed P-Value
I	1				
	T1SLEEP	1.000	0.000	999.000	999.000
	T2SLEEP	1.000	0.000	999.000	999.000
	T3SLEEP	1.000	0.000	999.000	999.000
S	1				
	T1SLEEP	0.000	0.000	999.000	999.000
	T2SLEEP	3.000	0.000	999.000	999.000
	T3SLEEP	6.000	0.000	999.000	999.000
I	ON GENCC	5.166	2.642	1.955	0.051
S	ON				
	GENCC	-1.655	0.963	-1.719	0.086
S	WITH				
	I	-23.870	12.746	-1.873	0.061
In	tercepts				
	T1SLEEP	0.000	0.000	999.000	999.000
	T2SLEEP	0.000	0.000	999.000	999.000
	T3SLEEP	0.000	0.000	999.000	999.000
	I	80.919	1.906	42.448	0.000
	S	-5.377	0.737	-7.297	0.000
Re	sidual Variances				
	T1SLEEP	3.183	53.594	0.059	0.953
	T2SLEEP	159.766	38.415	4.159	0.000
	T3SLEEP	94.112	76.438	1.231	0.218
	I	151.985	58.174	2.613	0.009
000000000000000000000000000000000000000	S	9.829	3.757	2.616	0.009

Pred Value for Age 10 (where Int = 1, and Slo = 0):

Males: 80.92(1) + (-5.38)(0) = 80.92

Females: (80.92 + 5.17)(1) + (-5.38 - 1.66)(0) = 86.09

Pred Value for Age 10 (where Int = 1, and Slo = 3):

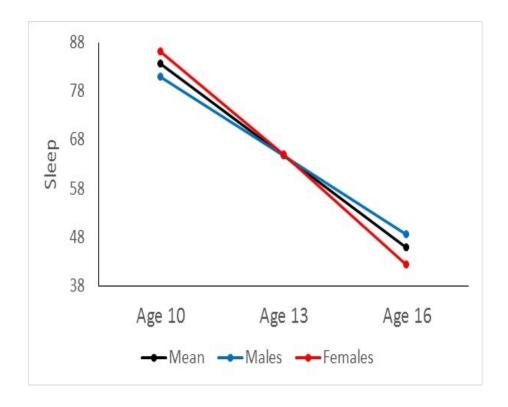
Males: 80.92(1) + (-5.38)(3) = 64.78

Females: (80.92 + 5.17)(1) + (-5.38 - 1.66)(3) = 64.97

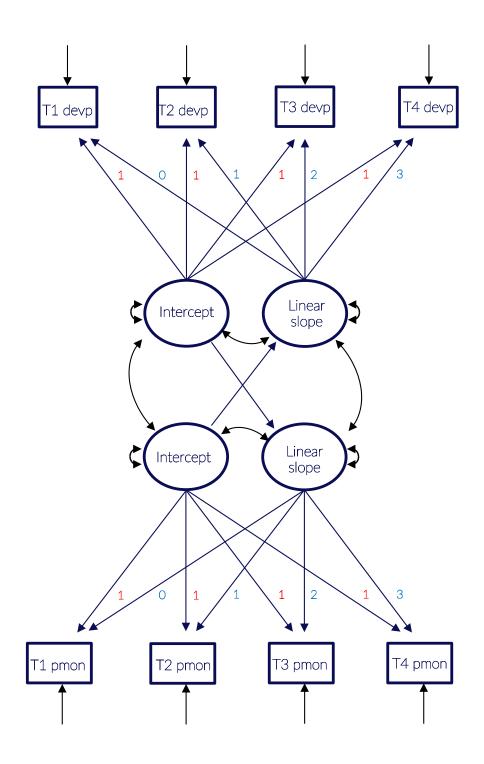
Pred Value for Age 10 (where Int = 1, and Slo = 6):

Males: 80.92(1) + (-5.38)(6) = 48.64

Females: (80.92 + 5.17)(1) + (-5.38 - 1.66)(6) = 42.24



Longitudinal data analyses: Bivariate latent growth curve modeling



MODEL RESULTS Two-Tailed Estimate S.E. Est./S.E. P-Value SMON ON IDEVP 0.259 0.080 3.219 0.001 SDEVP ON IMON 0.124 0.057 2.172 0.030 IMON WITH SMON -0.010 0.047 -0.220 0.826 -0.426 0.078 -5.432 0.000 IDEVP WITH IDEVP SDEVP -0.063 0.062 -1.020 0.308 WITH SMON SDEVP -0.020 0.016 -1.270 0.204

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Questions of interest:

1. Do the slopes of the two constructs correlate?

MODEL RESULTS Two-Tailed Estimate S.E. Est./S.E. P-Value SMON ON 0.259 IDEVP 0.080 3.219 0.001 SDEVP ON IMON 0.124 0.057 2.172 0.030 IMON WITH SMON -0.010 0.047 -0.220 0.826 -0.426 -5.432 0.000 IDEVP 0.078 IDEVP WITH SDEVP -0.063 0.062 -1.020 0.308 SMON WITH 0.204 SDEVP -0.020 0.016 -1.270

- 1. Do the slopes of the two constructs correlate?
- 2. Do the intercepts of the two constructs correlate?

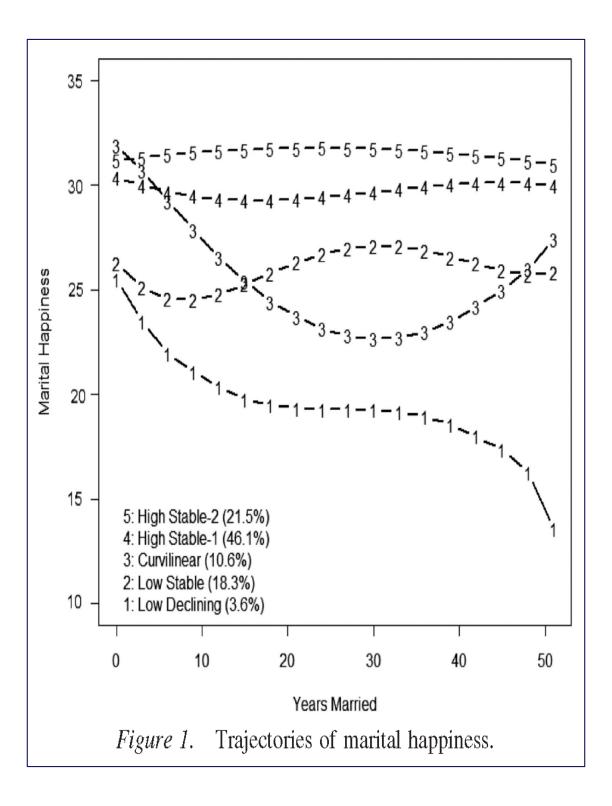
MODEL RESULTS Two-Tailed Estimate S.E. Est./S.E. P-Value SMON ON IDEVP 0.259 0.080 3.219 0.001 SDEVP ON IMON 0.124 0.057 0.030 2.172 IMON WITH SMON -0.010 0.047 -0.220 0.826 IDEVP -0.426 0.078 -5.432 0.000 IDEVP WITH SDEVP -0.063 0.062 -1.020 0.308 SMON WITH 0.204 SDEVP -0.020 0.016 -1.270

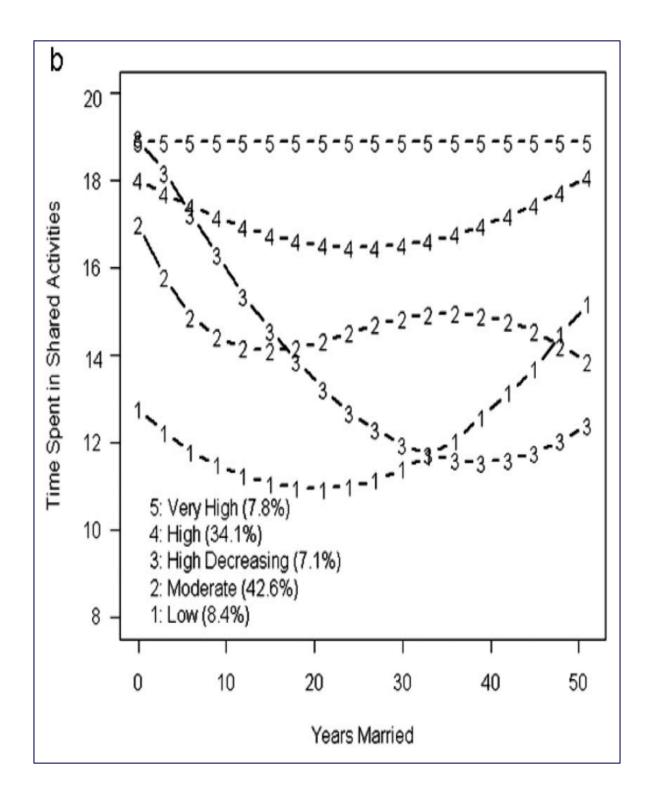
- 1. Do the slopes of the two constructs correlate?
- 2. Do the intercepts of the two constructs correlate?
- 3. Does parental monitoring at T1 predict change in deviant peer affiliation over time?

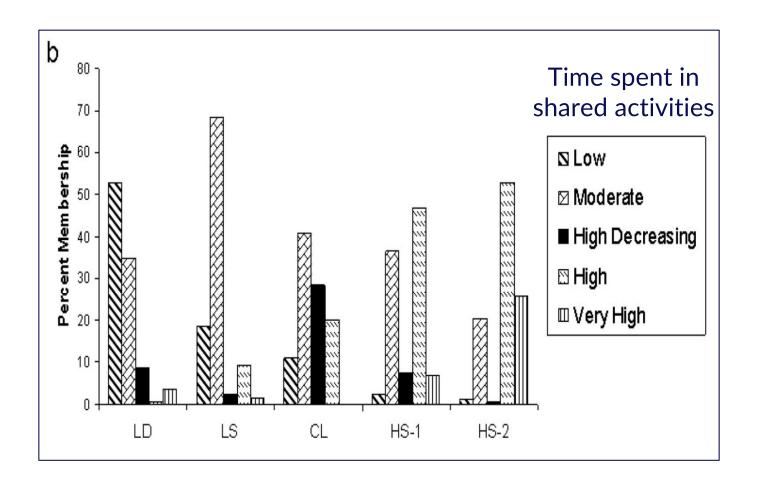
MODEL RESULTS Two-Tailed S.E. Est./S.E. Estimate P-Value SMON ON IDEVP 0.259 0.080 3.219 0.001 SDEVP ON 0.124 0.057 0.030 IMON 2.172 IMON WITH SMON -0.010 0.047 -0.220 0.826 IDEVP -0.4260.078 -5.4320.000 IDEVP WITH SDEVP -0.063 0.062 -1.0200.308 SMON WITH 0.204 SDEVP -0.020 0.016 -1.270

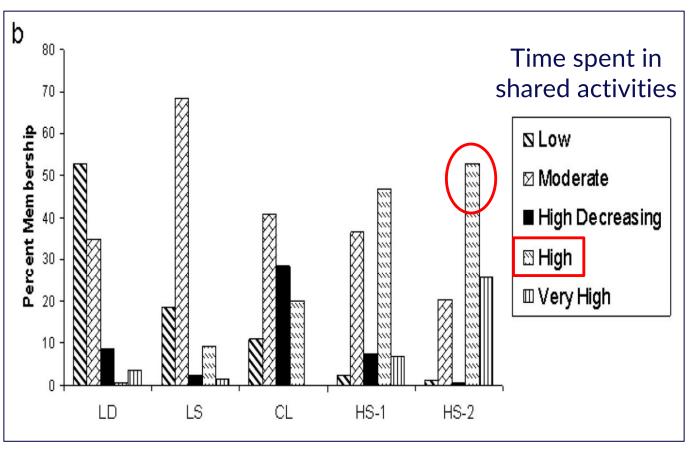
- 1. Do the slopes of the two constructs correlate?
- 2. Do the intercepts of the two constructs correlate?
- 3. Does parental monitoring at T1 predict change in deviant peer affiliation over time?
- 4. Does deviant peer affiliation at T1 predict change in parental monitoring over time?

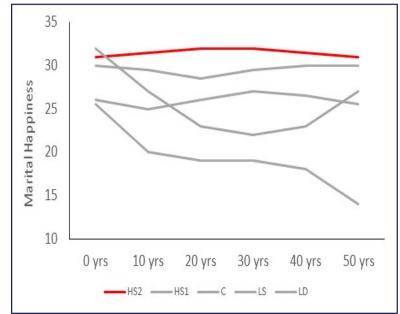
Longitudinal data analyses: Latent class growth analysis / growth mixture modeling

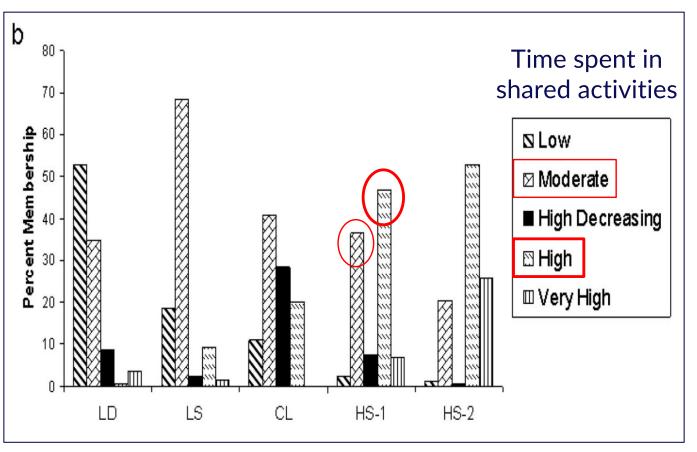


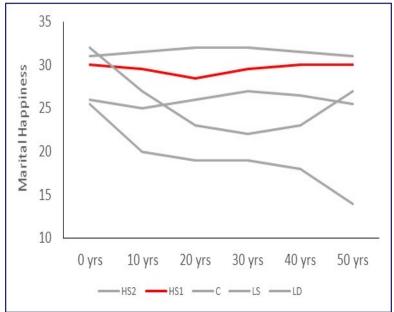


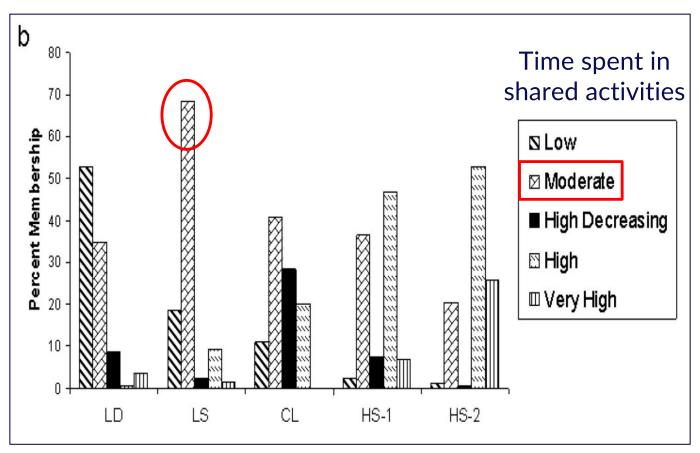


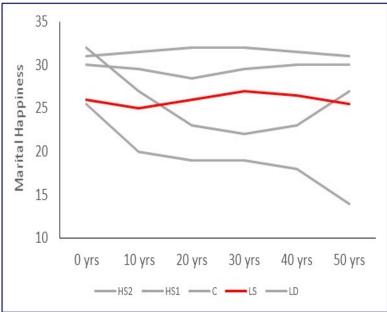


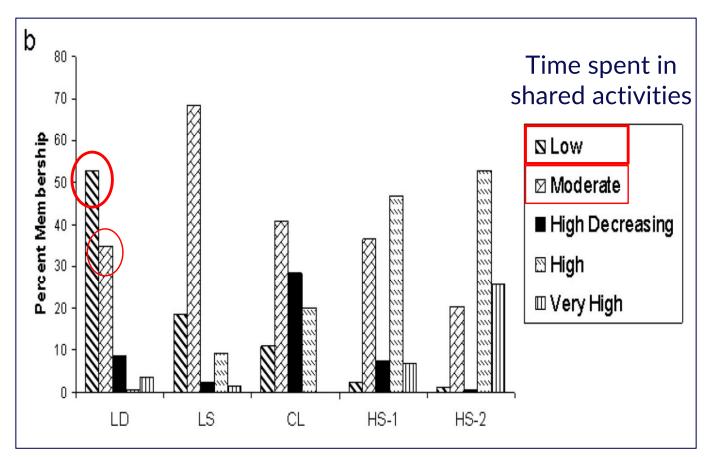


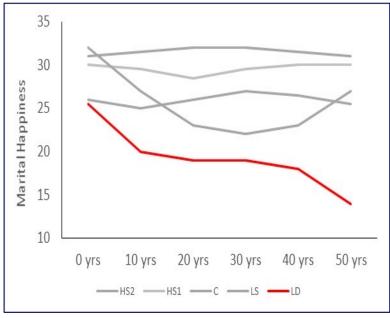


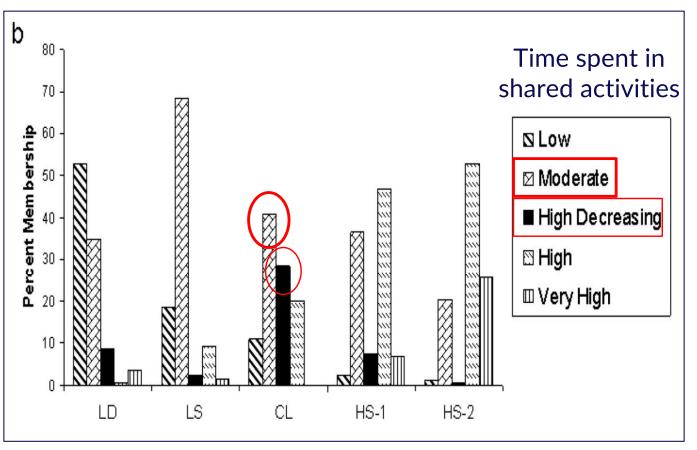


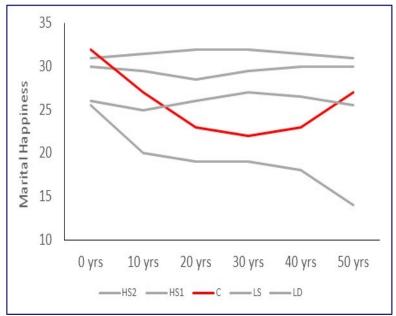












Violations of normality

Lecture 11-b

Violations of normality

Four assumptions of the GLM:

Independent observations

Randomly drawn from population

Error variances are equal across range of IV

(i.e., continuous predictor or groups)

Errors are normally distributed

The Box-Cox transformation

Identifies the exponent that will make a set of residuals closest to normal

Compute using R...

- 1) Put your data into tab-delimited or SPSS form
- 2) Import using: read.table or read.spss
- 3) Parse the data:

$$Y = data C1$$
 $X = data C2$ (where C1 and C2 are the first and second cols)

The Box-Cox transformation

Compute using R...

4) Then compute the actual transformation: boxcox(DV~IV, lambda=seq(from,to,by), plotit=T)

here,

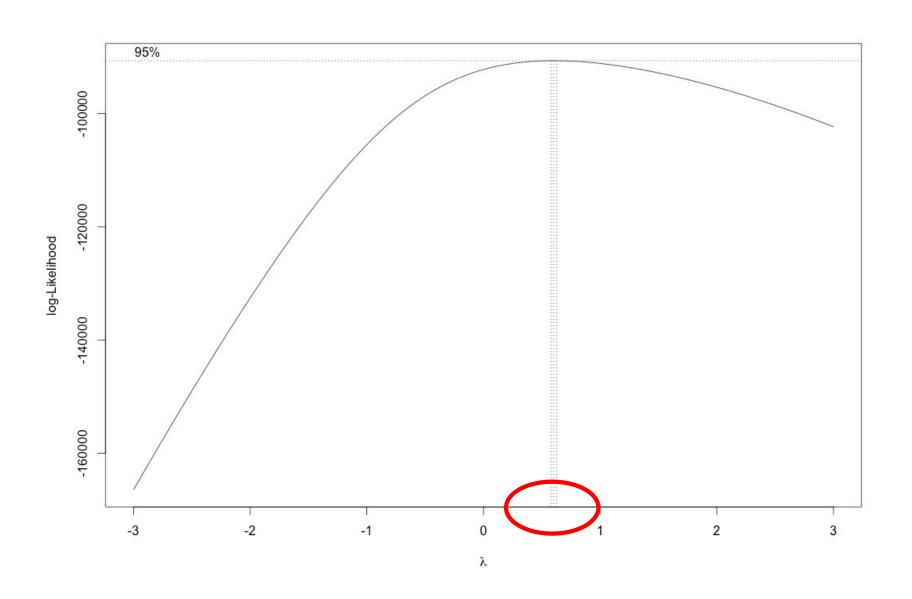
boxcox(Y~X,lambda=seq(-3,3,.01),plotit=T)

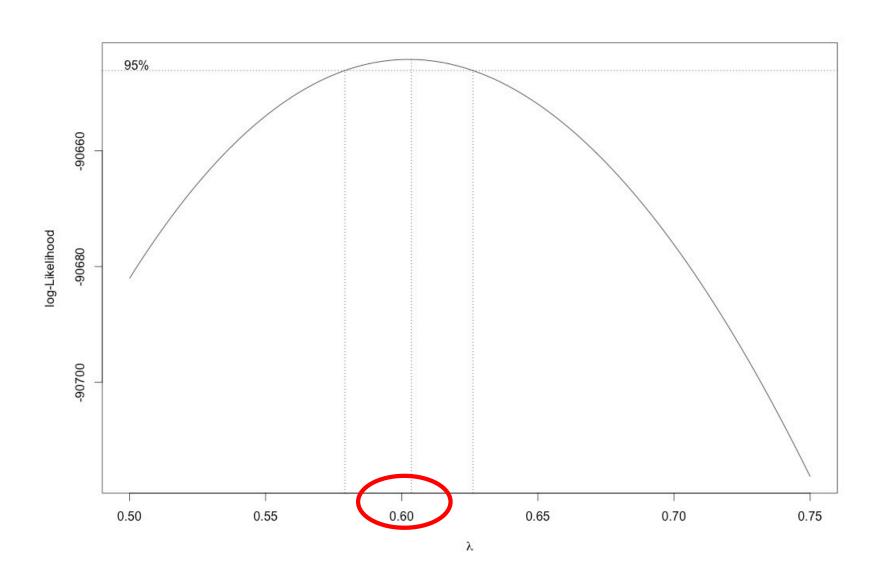
- 5) Zoom in to figure out the exact value
- 6) Transform your data (ynew = y^lambda)
- 7) Re-check: boxcox(ynew~x)

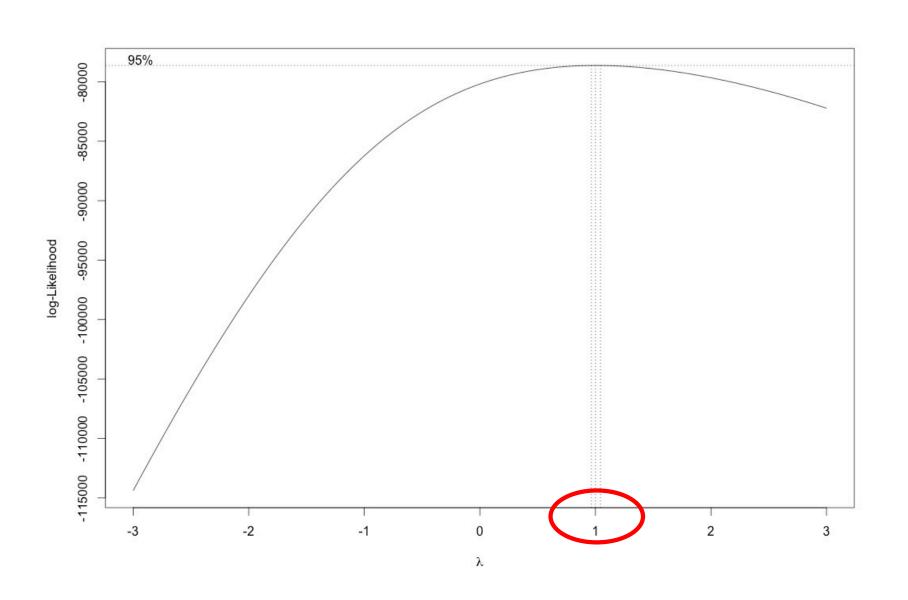
> summary(data)

	Length	Class	Mode
subj	23524	-none-	numeric
rt	23524	-none-	numeric
type	23524	factor	numeric

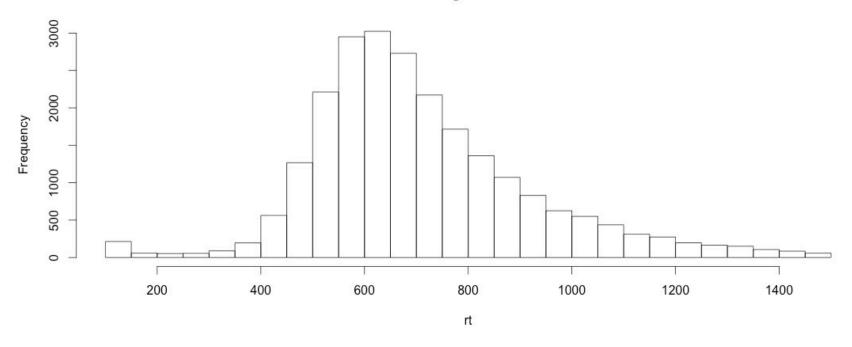
- > rt <- data\$rt
- > type <- data\$type
- > boxcox(rt~type, lambda = seq(-3,3,.01), plotit=1)







Histogram of rt



Histogram of rt_trans

