



Latent growth curve analysis in perinatal and pediatric epidemiology

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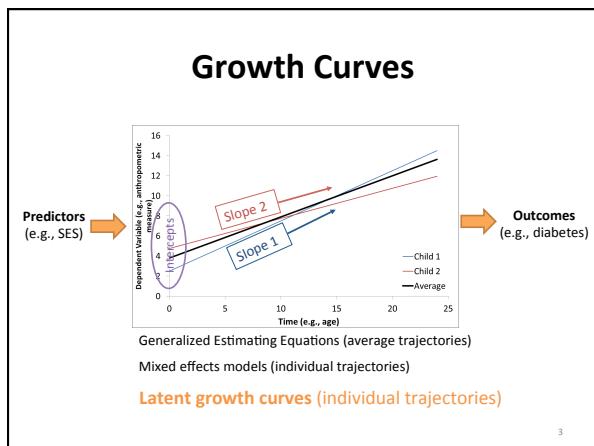
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Objectives

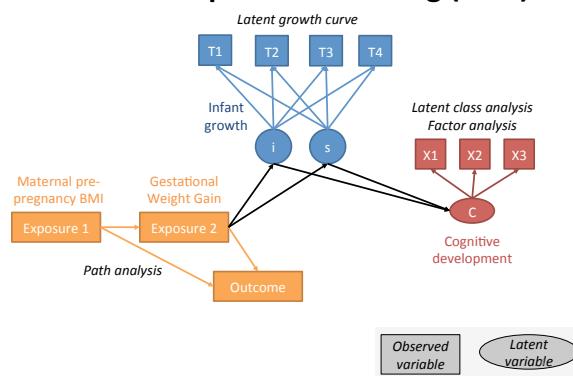
Participants will be able to:

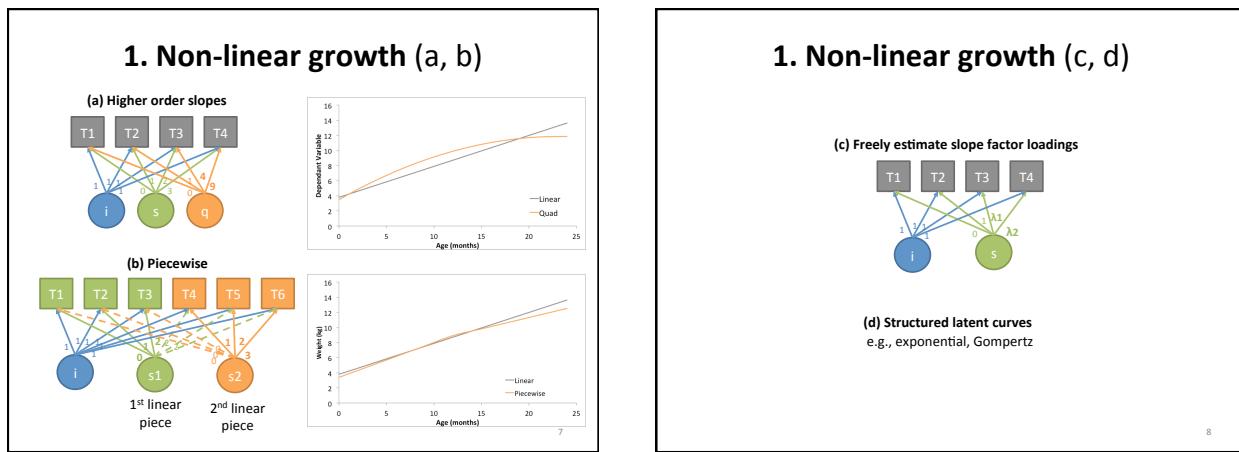
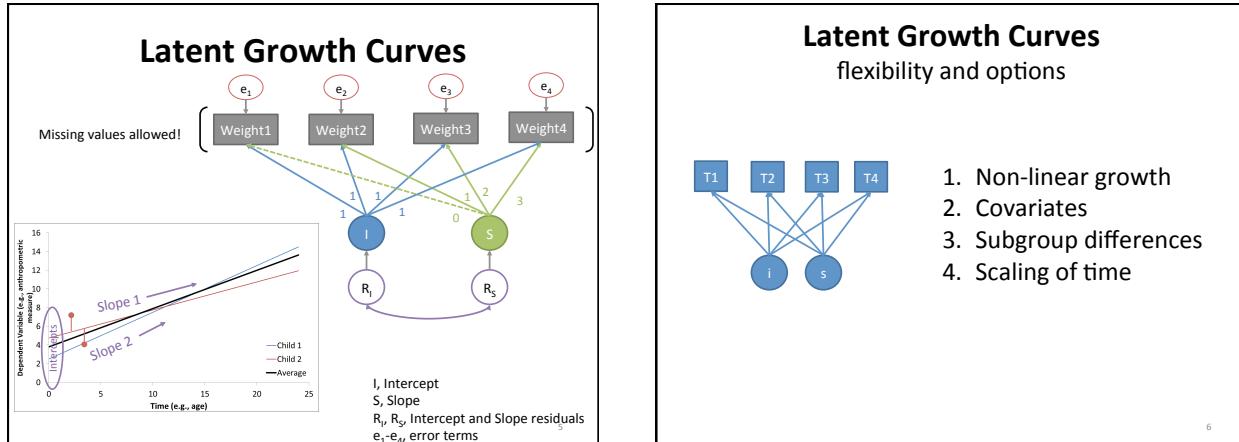
- Describe the strengths and general approach of latent growth curve analysis in existing research
- Evaluate whether latent growth curve analysis is appropriate for their research
- Apply basic latent growth analysis in Mplus (example: infant growth)

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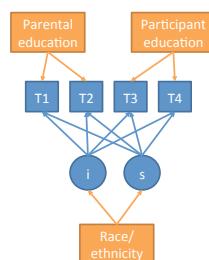


Structural Equation Modeling (SEM)





2. Covariates



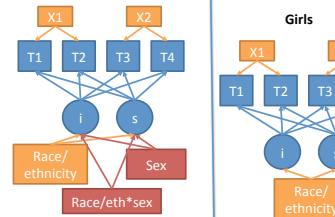
Time varying covariates

Time constant covariates

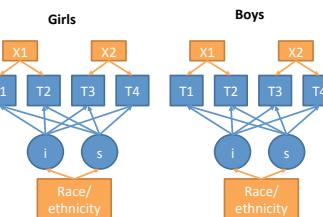
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3. Subgroup differences

Covariate/interaction



Multiple groups



Test: overall model fit in models that (a) constrain parameter(s) to be equal in boys and girls and (b) parameter(s) to differ in boys in girls

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4. Individually varying time points

Scenario 1

	Exam 1	Exam 2	Exam 3	Exam 4	Exam 5	Exam 6
Age (target)	Birth	3 mo	6 mo	9 mo	12 mo	18 mo
Age (actual)	Birth	2-4 mo	5-7 mo	8-10 mo	11-13 mo	14-19 mo

Scenario 2

	Exam 1	Exam 2	Exam 3	Exam 4	Exam 5	Exam 6
Child 1	Birth	2.9 mo	6.2 mo	9.8 mo	12.2 mo	19.7 mo
Child 2	Birth	1.2 mo	1.8 mo	4.4 mo	9.7 mo	12.1 mo

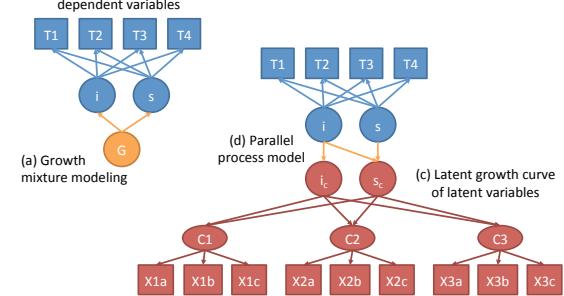
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Other topics

(b) Binary, ordinal dependent variables

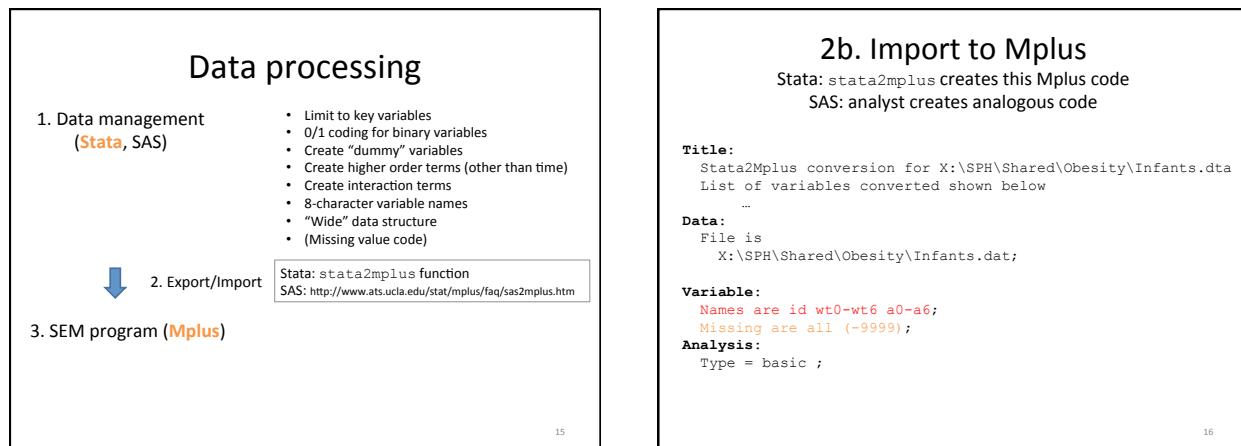
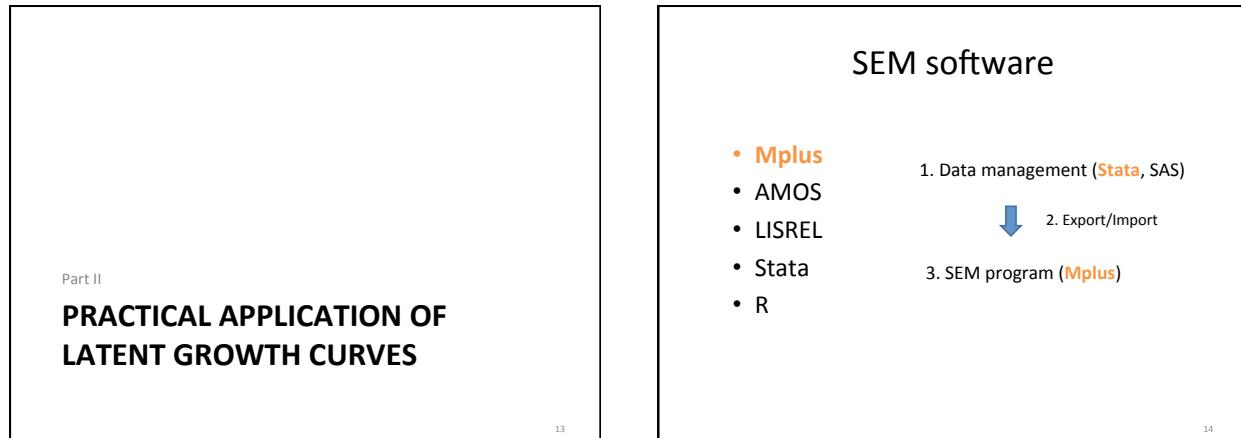


(a) Growth mixture modeling



Cognitive development factors

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3. Mplus analysis

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Example: Infant growth

Kaiser Permanente NW Region (KPNW) Electronic Medical Record Data MAT MORB (CDC-Funded, KPNW-based study)

Kaiser-based collaborators

Stephen P. Fortmann, MD
Kimberly Vesco, MD

A horizontal orange arrow pointing right, representing a timeline from 'Live Birth' to '2 years'.

- Dependent variable: **weight** (kg), 2 weeks to 24 months
 - In this example, selected measure within age intervals, closest to the following time points: birth, 3 months, 6 months, 9 months, 12 months, 18 months, 24 months
 - Latent Growth Curve analysis (MPlus 7.4)

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Mplus: basic syntax

Title: Linear LGM; Any (preferably informative) title

DATA: File is X:\SPH\Shared\Obesity\Infants.dat; File location

VARIABLE: Names are id wt0-wt6 a0-a6; ALL variables in dataset

Usevariables are wt0-wt6; Variables in the current analysis

Missing are all (-9999); Missing data code assigned in export step

ANALYSIS: ...

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Latent growth curve: linear [input]

Title: Linear LGM;

DATA: File is x:\SPH\Shared\Obesity\Infants.dat;

VARIABLE: Names are id wt0-wt6 a0-a6; Usevariables are wt0-wt6; Missing are all (-9999);

MODEL: i 1 wt@0 wt@.3 wt@.6 wt@.9 wt@1.2 wt@1.8 wt@2.4;

OUTPUT: TECH1;

User-specified names:
i = intercept term
l = linear slope term

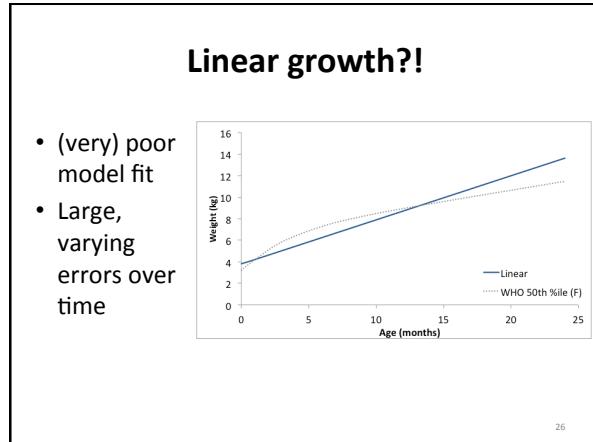
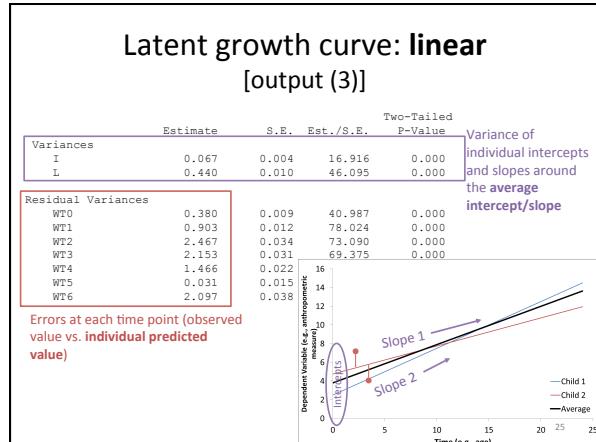
Note: FIML (for missing data) is the default

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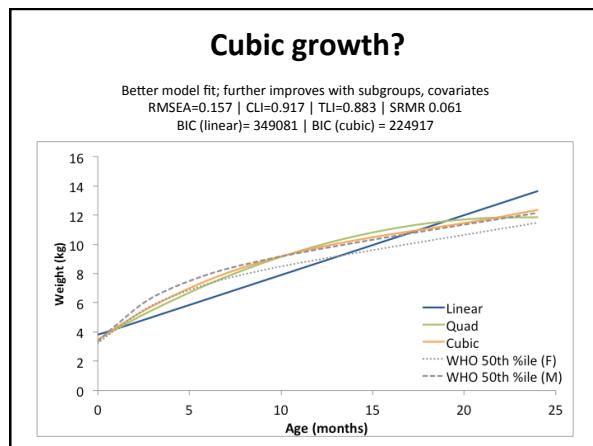
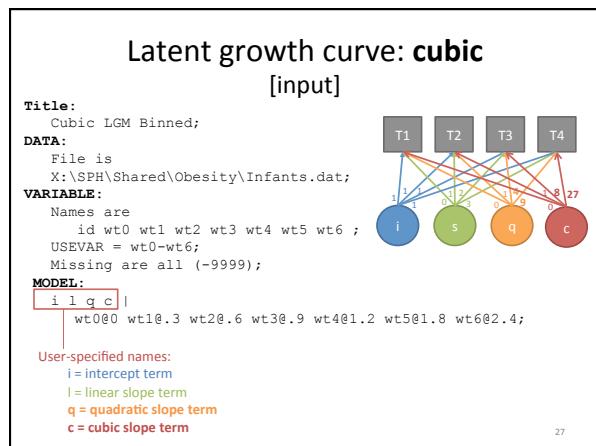
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Latent growth curve: linear [output (1)]					
THE MODEL ESTIMATION TERMINATED NORMALLY					
MODEL FIT INFORMATION					
Loglikelihood					
H0 Value		-174480.963			
H1 Value		-108311.313			
Information Criteria Useful for comparing models ($BIC_{diff} > 10$)					
Akaike (AIC)		348985.927			
Bayesian (BIC)		349081.857			
Sample-Size Adjusted BIC		349043.721			
$(n^* = (n + 2) / 24)$					
Chi-Square Test of Model Fit					
Value		132339.300			
Degrees of Freedom		23			
P-Value		0.0000			
...					
21					
Latent growth curve: linear [output (2)]					
...					
RMSEA (Root Mean Square Error Of Approximation)					
Estimate		0.513			
90 Percent C.I.		0.510 0.515			
Probability RMSEA $\leq .05$		0.000			
CFI/TLI					
CFI		0.000			
TLI		-0.247			
Chi-Square Test of Model Fit for the Baseline Model					
Value		96903.981			
Degrees of Freedom		21			
P-Value		0.0000			
SRMR (Standardized Root Mean Square Residual)					
Value		1.208			
≤ 0.05 is good					
This model has terrible fit					
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Latent growth curve: linear [output (2)]			
MODEL RESULTS			
I		Estimate	S.E. Est./S.E. Two-Tailed P-Value
WT0		1.000	0.000 999.000 999.000
WT1		1.000	0.000 999.000 999.000
WT2	Fixed at 1	1.000	0.000 999.000 999.000
WT3		1.000	0.000 999.000 999.000
WT4		1.000	0.000 999.000 999.000
WT5		1.000	0.000 999.000 999.000
WT6		1.000	0.000 999.000 999.000
L		Estimate	S.E. Est./S.E. Two-Tailed P-Value
WT0		0.000	0.000 999.000 999.000
WT1		0.300	0.000 999.000 999.000
WT2	Fixed path loadings for each time point	0.600	0.000 999.000 999.000
WT3		0.900	0.000 999.000 999.000
WT4		1.200	0.000 999.000 999.000
WT5		1.800	0.000 999.000 999.000
WT6		2.400	0.000 999.000 999.000
Latent growth curve: linear [output (2)]			
MODEL RESULTS			
L	WITH	Estimate	S.E. Est./S.E. Two-Tailed P-Value
WT0		0.139	0.004 33.084 0.000
WT1		3.847	0.009 411.701 0.000
WT2		4.097	0.009 456.723 0.000
WT3		0.000	0.000 999.000 999.000
WT4		0.000	0.000 999.000 999.000
WT5		0.000	0.000 999.000 999.000
WT6		0.000	0.000 999.000 999.000
Covariance between intercept and slope			
Mean intercept and slope			
Errors			
24			



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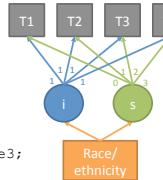


Latent growth curve: predictor(s) [input]

```
Title: Cube LGM, race as covariate;
DATA: File is X:\SPH\Shared\Obesity\Infants.dat;
VARIABLE: Names are id wt0-wt6 race1 race2 race3;
Usevariables are wt0-wt6 race1 race2 race3;
Missing are all (-9999);
MODEL: i l q c |
wt0@0 wt1@.3 wt2@.6 wt3@.9 wt4@1.2 wt5@1.8 wt6@2.4;
i on race1 race2 race3;
l on race1 race2 race3;
q on race1 race2 race3;
c on race1 race2 race3;
```

i on race1 race2 race3;
l on race1 race2 race3;
q on race1 race2 race3;
c on race1 race2 race3;

Rereg each growth term on
race indicator variables

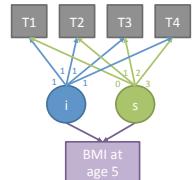


Latent growth curve: outcomes [input]

```
Title: Cube LGM, BMI as outcome;
DATA: File is X:\SPH\Shared\Obesity\Infants_BMI.dat;
VARIABLE: Names are id wt0-wt6 bmi;
Usevariables are wt0-wt6 bmi;
Missing are all (-9999);
MODEL: i l q c |
wt0@0 wt1@.3 wt2@.6 wt3@.9 wt4@1.2 wt5@1.8 wt6@2.4;
bmi on i;
bmi on l;
bmi on q;
bmi on c;
```

bmi on i;
bmi on l;
bmi on q;
bmi on c;

BMI at
age 5



Latent growth curve: subgroup differences [input]

```
Title: Cube LGM, by gender;
DATA: File is X:\SPH\Shared\Obesity\Infants_Gender.dat;
VARIABLE: Names are id wt0-wt6 bmi;
Usevariables are wt0-wt6 gender;
Missing are all (-9999);
GROUPING = gender(0=Male 1=Female);
MODEL: i l q c |
wt0@0 wt1@.3 wt2@.6 wt3@.9 wt4@1.2 wt5@1.8 wt6@2.4;
```

Test: overall model fit in models that (a) constrain parameter(s) to be equal in boys and girls and (b) parameter(s) to differ in boys in girls

Latent growth curve: subgroup differences [input]

```
Title: Cube LGM, by gender w/ equal parameters;
DATA: File is X:\SPH\Shared\Obesity\Infants_Gender.dat;
VARIABLE: ...
GROUPING = gender(0=Male 1=Female);
MODEL: i l q c |
wt0@0 wt1@.3 wt2@.6 wt3@.9 wt4@1.2 wt5@1.8 wt6@2.4;
Model: [i] (1);
[1] (2);
[q] (3);
[c] (4);
Model Female: [i] (1);
[1] (2);
[q] (3);
[c] (4);
```

Constraints:
Male intercepts = Female intercept
Male linear slope = Female linear slope
Male quadratic slope = Female quadratic slope
Male cubic slope = Female cubic slope

Intercept & slopes vary by gender: BIC 237906
Intercept & slopes equal: BIC 239834

Issues and challenges

- Conceptual challenges
 - Flexibility → more decisions, relies on strong theoretical framework
- Operational challenges:
 - Specialized software required (often Mplus)
 - Mplus learning curve
 - Difficulties in model convergence
 - Specify starting values
 - Increase number of iterations
 - Fixing variances, means, covariances

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Questions?

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