















Growth Mixture Analysis

Generalization of conventional random effect growth modeling (multilevel modeling) to include qualitatively different developments (Muthén & Shedden, 1999 in Biometrics). Combination of conventional growth modeling and cluster analysis (finite mixture analysis).

- Setting
 - Longitudinal data
 - A single or multiple items measured repeatedly
 - Hypothesized trajectory classes (categorical latent variable)
 - Individual trajectory variation within classes





Strategies For Finding The Number Of Classes In Growth Mixture Modeling

• Comparing models with different numbers of classes

- BIC low BIC value corresponds to a high loglikelihood value and a parsimonious model
- TECH11 Lo-Mendell-Rubin likelihood ratio test (Biometrika, 2001)
- TECH14 bootstrapped LRT (Version 4)

Residuals and model tests

- TECH7 class-specific comparisons of model-estimated means, variances, and covariances versus posterior probability-weighted sample statistics
- TECH12 class-mixed residuals for univariate skewness and kurtosis
- TECH13 multivariate skew and kurtosis model tests



Strategies For Finding Starting Values In Growth Mixture Modeling (Continued)

- Strategy 1
 - Do a conventional growth analysis (one-class model)
 - Use estimated growth factor means and standard deviations as growth factor mean starting values in a multi-class model mean plus and minus .5 standard deviation
- Strategy 2
 - Estimate a multi-class model with the variances and covariances of the growth factors fixed to zero (LCGA)
 - Use the estimated growth factor means as growth factor mean starting values for a model with growth factor variances and covariances free

Random starts makes it unnecessary to give starting values: starts = 505



Deciding On The Number Of Classes For The LSAY Growth Mixture Model					
n = 935					
Number of Classes	1	2	3		
Loglikelihood	-11,997.653	-11,864.826	-11,856.220		
# parameters	15	29	36		
BIC	24,098	23,928	23,959		
AIC	24,025	23,788	23,784		
Entropy	NA	.468	.474		
TECH11 LRT p-value for k-1 classes	NA	.0000	.4041		
Multivariate skew p-value	.00	.34	.26		
Multivariate kurtosis p-value	.00	.10	.05		





	Input For LSAY Two-Class Growth Mixture Model
TITLE:	2-class varying slopes on mothed and homeres varying Psi varying Theta
DATA:	FILE IS lsay.dat; FORMAT IS 3f8 f8.4 8f8.2 2f8.2;
VARIABLE:	<pre>NAMES ARE cohort id school weight math7 math8 math9 math10 att7 att8 att9 att10 gender mothed homeres; USEOBS = (gender EQ 1 AND cohort EQ 2); MISSING = ALL (999); USEVAR = math7-math10 mothed homeres; CLASSES = c(2);</pre>
ANALYSIS:	TYPE = MIXTURE;
	18

MODEL:	%OVERALL%
	intercpt slope math7@0 math8@1 math9 math10; intercpt slope ON mothed homeres; %c#2%
	intercpt slope ON mothed homeres;
	slope WITH intercept;
OUTPUT:	TECH8 TECH12 TECH13 RESIDUAL;



Output Excerpts LSAY Two-Class Growth Mixture Model (Continued) Classification Information FINAL CLASS COUNTS AND PROPORTIONS OF TOTAL SAMPLE SIZE Class 1 392.19327 0.41946 Class 2 542.80673 0.58054 CLASSIFICATION OF INDIVIDUALS BASED ON THEIR MOST LIKELY CLASS MEMBERSHIP Class Counts and Proportions

Class 1 342 0.36578 Class 2 593 0.63422 Average Class Probabilities by Class 2 1 Class 1 0.853 0.147 Class 2 0.170 0.830 21



Output Excerpts LSAY Two-Class Growth Mixture Model (Continued)

TECHNICAL 13 OUTPUT		
SKEW AND KURTOSIS TESTS OF MODEL FIT		
TWO-SIDED MULTIVARIATE SKEW TEST OF FIT		
Sample Value	1.245	
Mean	0.999	
Standard Deviation	0.275	
P-Value	0.3400	
TWO-SIDED MULTIVARIATE KURTOSIS TEST OF FIT		
Sample Value	29.612	
Mean	27.842	
Standard Deviation	1.015	
P-Value	0.1000	
		23

Output Excerpts LSAY Two-Class Growth Mixture Model (Continued)				
Model Results				
Class 1	Estimates	S.E.	Est./S.E.	
матн7	1 000	000	000	
МАТНЯ	1 000	000	.000	
МАТН9	1.000	.000	.000	
MATH10	1.000	.000	.000	
SLOPE				
MATH7	.000	.000	.000	
MATH8	1.000	.000	.000	
MATH9	2.422	.133	18.157	
MATH10	3.580	.204	17.570	
INTERCPT ON				
MOTHED	1.656	.626	2.645	
HOMERES	.720	.377	1.911	
SLOPE ON				
MOTHED	.146	.154	.953	
HOMERES	. 228	.087	2.626	24

Growth	Mixture Moo	del (Con	tinued)
	Estimates	S.E.	Est./S.E.
SLOPE WITH INTERCPT	727	1.643	443
	., .,		
Residual Variances			
MATH7	17.198	2.840	6.055
MATH8	15.257	2.077	7.347
MATH9	24.170	3.294	7.337
MATH10	49.112	10.037	4.893
INTERCPT	54.297	6.094	8.910
SLOPE	1.643	.627	2.620
Intercepts			
MATH7	.000	.000	.000
MATH8	.000	.000	.000
MATH9	.000	.000	.000
MATH10	.000	.000	.000
INTERCPT	42.733	1.648	25.922
SLOPE	.816	.366	2.228

Intercept S.E. Est./S.E. Class 2 INTERCPT Intercept Intercept MATH7 1.000 .000 .000 MATH8 1.000 .000 .000 MATH9 1.000 .000 .000 MATH10 1.000 .000 .000 SLOPE Intercept Intercept Intercept MATH7 .000 .000 .000 MATH10 1.000 .000 .000 SLOPE Intercept Intercept Intercept Intercept MATH9 2.422 .133 18.157 INTERCPT ON Intercept Intercept Intercept MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334	Output Excerpts LSAY Two-Class				
Estimates S.E. Est./S.E. Class 2 INTERCPT MATH7 1.000 .000 .000 MATH7 1.000 .000 .000 MATH9 1.000 .000 .000 MATH9 1.000 .000 .000 MATH9 1.000 .000 .000 SLOPE .000 .000 MATH7 .000 .000 .000 MATH7 .000 .000 .000 MATH7 .000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON .028 .079 .358	Growth Mixture Model (Continued)				
INTERCPT MATH7 1.000 .000 .000 MATH8 1.000 .000 .000 MATH9 1.000 .000 .000 MATH10 1.000 .000 .000 SLOPE MATH7 .000 .000 .000 MATH8 1.000 .000 .000 MATH8 1.000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON MOTHED .028 .079 .358	Class 2	Estimates	S.E.	Est./S.E.	
MATH7 1.000 .000 .000 MATH8 1.000 .000 .000 MATH9 1.000 .000 .000 MATH9 1.000 .000 .000 MATH10 1.000 .000 .000 SLOPE .000 .000 MATH7 .000 .000 .000 MATH8 1.000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON .0285 .376 5.545 HOMERES 1.805 .285 6.334	INTERCPT				
MATH8 1.000 .000 .000 MATH9 1.000 .000 .000 MATH10 1.000 .000 .000 SLOPE .000 .000 .000 MATH7 .000 .000 .000 MATH7 .000 .000 .000 MATH8 1.000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON .005 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON .028 .079 .358	MATH7	1.000	.000	.000	
MATH9 1.000 .000 .000 MATH10 1.000 .000 .000 SLOPE	MATH8	1.000	.000	.000	
MATH10 1.000 .000 .000 SLOPE	MATH9	1.000	.000	.000	
SLOPE .000 .000 .000 MATH7 .000 .000 .000 MATH8 1.000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON	MATH10	1.000	.000	.000	
MATH7 .000 .000 .000 MATH8 1.000 .000 .000 MATH8 1.000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON	SLOPE				
MATH8 1.000 .000 .000 MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON MOTHED .028 .079 .358	MATH7	.000	.000	.000	
MATH9 2.422 .133 18.157 MATH10 3.580 .204 17.570 INTERCPT ON MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON MOTHED .028 .079 .358	MATH8	1.000	.000	.000	
MATH10 3.580 .204 17.570 INTERCPT ON MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON MOTHED .028 .079 .358	MATH9	2.422	.133	18.157	
INTERCPT ON MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON MOTHED .028 .079 .358	MATH10	3.580	.204	17.570	
MOTHED 2.085 .376 5.545 HOMERES 1.805 .285 6.334 SLOPE ON .028 .079 .358	INTERCPT ON				
HOMERES 1.805 .285 6.334 SLOPE ON MOTHED .028 .079 .358	MOTHED	2.085	.376	5.545	
SLOPE ON MOTHED .028 .079 .358	HOMERES	1.805	.285	6.334	
MOTHED .028 .079 .358	SLOPE ON				
.020 .079 .000	MOTHED	028	079	358	
HOMERES 054 058 943	HOMERES	054	058	943	
.054 .050 .545	полысыр	:054	.050	.945	26

Growth Mixture Model (Continued)				
SIODE WITH	Estimates	S.E.	Est./S.E.	
INTERCPT	.478	.575	.831	
Residual Variances				
MATH7	12.248	1.587	7.720	
MATH8	11.375	1.244	9.141	
MATH9	8.014	1.420	5.642	
MATH10	7.349	1.312	5.600	
INTERCPT	32.431	4.651	6.973	
SLOPE	.191	.264	.724	
Intercepts				
MATH7	.000	.000	.000	
MATH8	.000	.000	.000	
MATH9	.000	.000	.000	
MATH10	.000	.000	.000	
INTERCPT	45.365	1.461	31.054	
SLOPE	2.847	.316	9.015	
LATENT CLASS REGRESS	ION MODEL PART			
M				

(G	Dutput 1 rowth 1	Excerpt Mixture	s LSAY Model	Two-C (Contin	lass ued)	
Residual	ls					
ESTIMATE	D MODEL ANI	RESIDUALS	(OBSERVED	- ESTIMATED) FOR CLASS	
Model Es	timated Mea	ins				
MATH7 48.650	MATH8 50.486	MATH9 53.095	MATH10 55.221	MOTHED 2.255	HOMERES 3.030	
Residual	s for Means	3				
MATH7 -0.113	MATH8 0.072	MATH9 0.236	MATH10 -0.389	MOTHED 0.000	HOMERES 0.000	
Model Es	timated Cov	variances				
MATH7 MATH8 MATH9	MATH7 76.058 60.370	<u>MATH8</u> 78.945	<u>MATH9</u>	MATH10	MOTHED	
MATH10 MOTHED	64.263 1.751	72.253	83.615	141.979 2.507	0.907	
HOMERES	2.309	2.908	3.760	4.454	0.345	

	HOMERES				
HOMERES	2.412				
Devident	f. G.				
Residuals	s Lor Covar	Lances			
-	MATH7	MATH8	MATH9	MATH10	MOTHED
MATH'7	-0.153				
MATH8	-0.109	-0.143			
MATH9	0.413	0.572	0.451		
MATH10	-0.701	-0.614	-0.254	-0.338	
MOTHED	0.210	-0.252	-0.021	0.256	0.000
HOMERES	0.289	-0.254	-0.358	0.720	0.000
	HOMERES				
HOMEDEC	0.000				



Further Readings On Growth Mixture Modeling (Continued)

Muthén, B. & Muthén, L. (2000). Integrating person-centered and variable-centered analysis: growth mixture modeling with latent trajectory classes. <u>Alcoholism: Clinical and Experimental</u> <u>Research</u>, 24, 882-891. (#85)

Muthén, B. & Shedden, K. (1999). Finite mixture modeling with mixture outcomes using the EM algorithm. <u>Biometrics</u>, 55, 463-469. (#78)

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General Growth Mixture Modeling (GGMM)

GGMM goes beyond conventional random effect growth modeling by using latent trajectory classes which

- Allow for heterogeneity with respect to
 - Growth functions different classes correspond to different growth shapes
 - Antecedents different background variables have different importance for different classes
 - Consequences class membership predicts later outcomes
- Allow for prediction of trajectory class membership
- Allow for confirmatory clustering
 - With respect to parameters describing curve shapes
 - With respect to typical individuals known classes
- Allow for classification of individuals
 - Early prediction of problematic development
 - Allow for enhanced preventive intervention analysis
 - Different classes benefit differently and can receive different treatments















Multinomial Logistic Regression Of c ON x

The multinomial logistic regression model expresses the probability that individual *i* falls in class *k* of the latent class variable *c* as a function of the covariate x,

$$P(c_j = k \mid x_i) = \frac{e^{\alpha_k} + \gamma_k x_i}{\sum_{s=1}^{K} e^{\alpha_s} + \gamma_s x_i} , \qquad (90)$$

where $\alpha_{\kappa} = 0$, $\gamma_{\kappa} = 0$ so that $e^{\alpha_{\kappa} + \gamma_{\kappa} x_{i}} = 1$.

This implies that the log odds comparing class k to the last class K is

$$log[P(c_i = k \mid x_i)/P(c_i = K \mid x_i)] = \alpha_k + \gamma_k x_i.$$
(91)

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Heavy Drinking And Alcohol Dependence NLSY Cohort64 (n=922)

		HD Classes				
	1 (D	own)	2 (Hi	gh18)	3(ไ	Jp)
	Est.	t	Est.	t	Est.	t
Male	1.21	5.52	1.25	3.48	1.45	4.73
Black	-0.89	-3.43	-3.14	-2.86	-0.06	-0.17
Hisp	-0.65	-2.22	-0.35	-0.86	-0.01	-0.03
ES	1.24	4.79	2.05	5.72	0.71	1.78
FH1	0.03	0.09	-0.21	-0.41	-0.08	-0.16
FH23	0.04	0.15	0.25	0.56	0.08	0.23
FH123	-0.23	-0.58	1.18	2.59	1.00	2.60
HSDRP	0.57	1.98	0.32	0.76	0.91	2.93
Coll	-0.07	-0.31	-1.31	-2.85	-1.08	-2.59

Heavy Drinking And Alcohol Dependence NLSY Cohort64 (n=922) (Continued)

	Probability	Odds Ratio
HD Class 1 (Down)	0.16	3.92
HD Class 2 (High 18)	0.26	7.06
HD Class 3 (Up)	0.60	30.00
HD Class 4 (Norm)	0.05	1.00

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Input Excerpts NLSY Growth Mixture Model With Covariates And A Distal Outcome (Continued)

!	<pre>log age scale: x_t = a*(ln(t-b)</pre>	-	ln(c	-b));	
!	where t is time, a and b are con	ıst	ants	to fit the mean curv	re
!	(chosen as a = 2 and b = 16), and	nd	c is	the centering age,	
!	here set at 25.				
	%c#1%	!	Not	needed	
	[dep94\$1*1 i*2 s1*5 s2*1];	!	Not	needed	
	%c#2%	!	Not	needed	
	[dep94\$1*0 i*1 s1*2 s2*3];	!	Not	needed	
	%c#3%	!	Not	needed	
	[dep94\$1*.6 i*3 s1*1.5 s2*.2];	!	Not	needed	
	%c#4%	!	Not	needed	
	[dep94\$1*2 i*.6 s1*2 s2*1];	!	Not	needed	
				15	
				45	



	Covariate	s And A Dista	l Outcome	(Continu	ed)
C#2	ON				
MALE		1.248	.359	3.481	
BLACH	τ	-3.138	1.097	-2.860	
HISP		346	.401	864	
ES		2.045	.357	5.722	
FH1		211	.514	410	
FH23		.247	.444	.556	
FH123	3	1.178	.456	2.585	
HSDRI)	.323	.428	.756	
COLL		-1.311	.460	-2.851	
C#3	ON				
MALE		1.454	.308	4.727	
BLACH	τ	059	.344	171	
HISP		011	.369	030	
ES		.712	.399	1.784	
FH1		079	.502	157	
FH23		.084	.364	.232	
FH123	3	1.004	.387	2.596	
HSDRI	b	.913	.312	2.926	
COLL		-1.075	.414	-2.594	4

Output Excerpts NLSY Growth Mixture Model With Covariates And A Distal Outcome (Continued)

Class 1 Thresholds DEP94\$1	1.631	0.248	6.574	
Class 2 Thresholds DEP94\$1	1.041	0.338	3.077	
Class 3 Thresholds DEP94\$1	-0.406	0.272	-1.493	
Class 4 Thresholds DEP94\$1	2.987	0.208	14.392	
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	Input For Growth Mixture Model For Reading Skills Development	
TITLE:	Growth mixture model for reading skills development	
DATA:	FILE IS newran.dat;	
VARIABLE:	<pre>NAMES ARE gender eth wc pal-pa4 wrl-wr8 ll-l4 sl rl rnamingl rnaming2 rnaming3 rnaming4; USEVAR = pal-wr8 rnaming4; MISSING ARE ALL (999); CLASSES = c(5);</pre>	s2 r
ANALYSIS:	TYPE = MIXTURE MISSING;	
MODEL:	%OVERALL% il sl pal@-3 pa2@-2 pa3@-1 pa4@0; i2 s2 wr1@-7 wr2@-6 wr3@-5 wr4@-4 wr5@-3 wr6@-2 wr7@-1 wr8@0; c#1-c#4 ON rnaming4;	
OUTPUT:	TECH8;	55





				Full Model		I	
		1.00	2.00	3.00	4.00	5.00	Total
к	1.00	28	7	3			38
Only	2.00	10	29	16			55
	3.00	8	33	100	25		166
	4.00		1	24	63	17	105
	5.00		1	1	12	32	46
Total		46	71	144	100	49	410
				Full Model			
		1.00	2.00	3.00	4.00	5.00	Total
K + 1	1.00	28	7	3			38
Only	2.00	15	44	24			83
	3.00	3	20	112	20		155
	4.00			5	79	4	88
	5.00				1	45	46
Total		46	71	144	100	49	410
				Full Model			
		1.00	2.00	3.00	4.00	5.00	Total
K + 2	1.00	28	8				36
Only	2.00	16	54	22			92
-	3.00	2	9	119	7		137
	4.00			4	91	4	99
	5.00				2	45	47
Total		46	71	145	100	49	411

				Full Model			
	-	1.00	2.00	3.00	4 00	5.00	Total
K + 3	1.00	37	12	0.00		0.00	49
Only	2.00	9	53	8			70
,	3.00	-	6	136	4		146
	4.00		-	1	95	1	97
	5.00				1	48	49
Total		46	71	145	100	49	411
	I		1	1			
				Full Model			
		1.00	2.00	3.00	4.00	5.00	Total
K + 4	1.00	45	11				56
Only	2.00	1	57	3			61
	3.00		3	141	2		146
	4.00			1	97		98
	5.00				1	49	50
Total		46	71	145	100	49	411
				Full Model			
		1.00	2.00	3.00	4.00	5.00	Total
K + 5	1.00	45	3				48
Only	2.00	1	66				67
	3.00		2	145	1		148
	4.00				98		98
	5.00				1	49	50
Total		46	71	145	100	49	411

		Full Model					
		1.00	2.00	3.00	4.00	5.00	Total
K + 6	1.00	46					46
Only	2.00		69				69
	3.00		1	145	1		147
	4.00				98		98
	5.00				1	49	50
Total		46					
		40	70	145	100	49	410
		40		145	100	49	410











TITLE: growt DATA: FILE VARIABLE: NAMES	h mixtures in randomized trials
DATA: FILE VARIABLE: NAMES	TO 4444 Jak.
VARIABLE: NAMES	IS toca.dat;
sctaa MISSI USEVA CLASS	ARE sctaallf sctaalls sctaal2f sctaal2s sctaal3s .14s sctaal5s sctaal6s sctaal7s intngrp; NG ARE ALL (999); RIABLES ARE sctaal1f-sctaal7s tx; SES = c(3);
DEFINE: tx =	(intngrp==4);
ANALYSIS: TYPE	= MIXTURE MISSING;
MODEL: %OVER ac bo sctaa sctaa qc@0; bc qc sctaa	ALL% gc sctaallf@0 sctaalls@0.5 sctaal2f@1 .12s@1.5 sctaal3s@2.5 sctaal4s@3.5 sctaal5s@4.5 .16s@5.5 sctaal7s@6.5; 2 ON tx; .11f WITH sctaalls; sctaal2f WITH sctaal2s;

Input For Growth Mixtures In Randomized Trials (Continued)

%c#1%
[ac*3 bc qc]; bc qc ON tx;
%c#2%
[ac*2 bc qc]; bc qc ON tx;
%c#3%
[ac*1 bc qc]; bc qc ON tx;
ac sctaallf-sctaal7s;

