

Scripts & Outputs

“Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

Latent Profile Analysis (LPA) in Mplus

Step 1: Model Parameterization Structure

The parametrization structure for the LPA models was defined based on comparison of different parametrization structures given three class model.

Parametrization Model 1 (Default) - Equal variances and zero covariances

```
Mplus VERSION 8.7
MUTHEN & MUTHEN
07/11/2023 11:26 AM

INPUT INSTRUCTIONS

  TITLE: LPA 3 profile syntax: M1 free means, constrained
  variance, zero covariance
  DATA:
  FILE IS mplus2.csv;
  ! Specifies file location for data file. Make sure data is in
  format appropriate for Mplus
  ! per Mplus manual. This data file is in individual format
  (one row of data per participan
  VARIABLE:
  NAMES ARE ID SCI_02 SCI_02B slpnssN worryN slp
  slpDurC gender genderN yrgrp yrgrpN SCI_02C
  slpnssNC worryNC solC slpDurC;

  ! All variables included in data file should be named here.
  USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
  ! Only variables intended for use in the analysis should be
  listed here
  IDVARIABLE IS ID;
  CLASSES = c (3);
  ! This is where you instruct Mplus on how many
  classes/profiles are being estimated. Initi
  ! model contains only one class/profile, thus it would be
  CLASSES = c (1). Above specifies
  !profiles, and for each further iterative models the number
  in parentheses increases by on
  !three profiles/classes would be c (3), and so on.
  MISSING ARE ALL(9999);
  ! Used to communicate how missing data is coded in data
  file. Here shown with a "." which
  ! all that is included in each cell with missing data in the
  data file
  ANALYSIS:
  TYPE = MIXTURE;
  ! LPA is a version of mixture modeling, and this instructs
  Mplus to analyze in this way
  ESTIMATOR = MLR;
  !FIML robust to non-normal data
  STARTS = 1000 250;
  STITERATIONS = 500;
  ! Default number of starts for each step of the ML
  estimation. First STARTS value specifie
  !number of unique start values to start with, the 250
  represents the 250 best unique start
  !carrying forward to completion. The STITERATIONS
  specifies the number of ML iteration
  !steps for those 250 selected start values to go through to
  be able to converge. This is a

  !maximum number of iteration; if a model converges in less
  than 500 iterations it will sto
  !before reaching 500 iterations.
  !These values can be increased ... see "Four-Profile Final
  Model with Covariate Analysis
  !Syntax" for an example.
  LRTSTARTS = 2 1 50 10;
  LRTBOOTSTRAP = 250;
  !The above start values are for the defaults for the LRT
  statistic being run to compare th
  !model fit with the model fit of a model with one less class
  (k-1). The BOOTSTRAP statemen
  !specifies the number of bootstrap draws to inform Mplus'
  bootstrapped LRT results.
  MODEL:
  !For a default Mplus model the LPA model does not need
  to be specified. However, it can be
  !The model can also be modified from the Mplus default of
  estimating the indicator means
  !(uniquely across profiles) and variances (constrained
  across profiles), as well as the la
  !profile mean.
  %OVERALL%
  [slpnssNC worryNC solC slpDurC]; !estimates the
  indicators means for each profile. Without
  !the means are freely estimated in each profile, not
  constrained.
  slpnssNC worryNC solC slpDurC; !Label Var1-Var5
  constrains the estimates of the variances
  !profiles to be equal.

  OUTPUT:
  TECH11 TECH14;
  ! TECH1 provides parameter specifications and starting
  values for the analysis
  ! TECH8 provides optimization history for this analysis type
  !TECH11 provides LRT results
  !TECH14 provides bootstrapped LRT test
  PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);
  TYPE=PLOT3;
  SAVEDATA:
  FILE IS Step1_LPA3M1.dat;
  ! Tells Mplus where to save the output files from the
  analysis
  SAVE = CPROBABILITIES;
  ! The above command lines are to save the most likely
  profile membership for each particip
  ! and the posterior probabilities for their membership in
  each latent profile.

  *** WARNING in MODEL command
  All variables are uncorrelated with all other variables within
  class.
  Check that this is what is intended.
```

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		mplus2.csv	
LPA 3 profile syntax: M1 free means, constrained variance, zero covariance		Input data format FREE	
SUMMARY OF ANALYSIS		SUMMARY OF DATA	
Number of groups	1	Number of missing data patterns	4
Number of observations	27802	Number of y missing data patterns	4
		Number of u missing data patterns	0
Number of dependent variables	4	COVARIANCE COVERAGE OF DATA	
Number of independent variables	0	Minimum covariance coverage value 0.100	
Number of continuous latent variables	0	PROPORTION OF DATA PRESENT FOR Y	
Number of categorical latent variables	1		
Observed dependent variables			
Continuous		Covariance Coverage	
SLPNSSNC	WORRYNC	SOLC	SLPDURC
Categorical latent variables		SLPNSSNC	
C		WORRYNC	
Variables with special functions		SOLC	
ID variable	ID	SLPDURC	
Estimator	MLR	UNIVARIATE SAMPLE STATISTICS	
Information matrix	OBSERVED	UNIVARIATE HIGHER-ORDER MOMENT	
Optimization Specifications for the Quasi-Newton Algorithm for		DESCRIPTIVE STATISTICS	
Continuous Outcomes		Variable/	Mean/
Maximum number of iterations	100	Percentiles	Skewness/
Convergence criterion	0.100D-05	Sample Size	Minimum/
Optimization Specifications for the EM Algorithm		Min/Max	% with
Maximum number of iterations	500	20%/60%	Maximum
Convergence criteria		40%/80%	Median
Loglikelihood change	0.100D-06	SLPNSSNC	0.933
Relative loglikelihood change	0.100D-06	0.000	1.000
Derivative	0.100D-05	27802.000	0.471
Optimization Specifications for the M step of the EM Algorithm for		-0.928	0.020
Categorical Latent variables		27802.000	1.000
Number of M step iterations	1	0.495	0.969
M step convergence criterion	0.100D-05	SOLC	0.000
Basis for M step termination	ITERATION	1.054	-0.072
Optimization Specifications for the M step of the EM Algorithm for		27123.000	1.000
Censored, Binary or Ordered Categorical (Ordinal), Unordered		-0.072	0.910
Categorical (Nominal) and Count Outcomes		SLPDURC	0.000
Number of M step iterations	1	-1.205	-0.292
M step convergence criterion	0.100D-05	26763.000	1.000
Basis for M step termination	ITERATION	-0.292	0.620
Maximum value for logit thresholds	15	RANDOM STARTS RESULTS RANKED FROM THE BEST TO THE WORST LOGLIKELIHOOD VALUES	
Minimum value for logit thresholds	-15	Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:	
Minimum expected cell size for chi-square	0.100D-01	-128973.336	780698
Maximum number of iterations for H1	2000	-128973.336	569338
Convergence criterion for H1	0.100D-03	-128973.336	699810
Optimization algorithm	EMA	-128973.336	298553
Random Starts Specifications		-128973.336	232559
Number of initial stage random starts	1000	-128973.336	628143
Number of final stage optimizations	250	-128973.336	635245
Number of initial stage iterations	500	-128973.336	27071
Initial stage convergence criterion	0.100D+01	-128973.336	810705
Random starts scale	0.500D+01	-128973.336	545108
Random seed for generating random starts	0	-128973.336	926283
Input data file(s)			269

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-128973.336	268896	124	-128973.336	383902	673
-128973.336	850112	922	-128973.336	887676	22
-128973.336	297518	166	-128973.336	351622	551
-128973.336	738393	619	-128973.336	402699	604
-128973.336	214681	824	-128973.336	130011	587
-128973.336	797594	234	-128973.336	72344	897
-128973.336	425929	508	-128973.336	474357	789
-128973.336	200041	810	-128973.336	318177	748
-128973.336	609089	241	-128973.336	458181	189
-128973.336	853781	716	-128973.336	483369	270
-128973.336	914505	838	-128973.336	92564	583
-128973.336	399848	220	-128973.336	288738	940
-128973.336	444228	860	-128973.336	535063	329
-128973.336	209031	632	-128973.336	475419	987
-128973.336	351807	617	-128973.336	966499	963
-128973.336	937885	426	-128973.336	424223	900
-128973.336	646573	741	-128973.336	848890	95
-128973.336	327140	678	-128973.336	195353	225
-128973.336	996231	310	-128973.336	117391	710
-128973.336	264901	634	-128973.336	326091	759
-128973.336	710445	792	-128973.336	965994	396
-128973.336	568859	49	-128973.336	966603	919
-128973.336	435506	988	-128973.336	89970	223
-128973.336	573096	20	-128973.336	865906	641
-128973.336	85734	411	-128973.336	891347	504
-128973.336	21132	351	-128973.336	484406	421
-128973.336	563002	360	-128973.336	849670	347
-128973.336	538872	949	-128973.336	507218	613
-128973.336	248742	556	-128973.336	900921	984
-128973.336	686482	215	-128973.336	970689	266
-128973.336	705224	953	-128973.336	810594	845
-128973.336	43523	297	-128973.336	192191	629
-128973.336	950604	172	-128973.336	314034	513
-128973.336	499150	216	-128973.336	939021	8
-128973.336	668003	647	-128973.336	438144	271
-128973.336	357866	968	-128973.336	341960	685
-128973.336	284716	713	-128973.336	137888	901
-128973.336	285380	1	-128973.336	281558	184
-128973.336	224950	455	-128973.336	392751	480
-128973.336	67009	564	-128973.336	850545	357
-128973.336	490123	995	-128973.336	928287	197
-128973.336	462228	298	-128973.336	264081	186
-128973.336	344422	296	-128973.336	629720	926
-128973.336	72662	729	-128973.336	572637	989
-128973.336	291112	645	-128973.336	644297	340
-128973.336	805768	879	-128973.336	323588	826
-128973.336	68850	462	-128973.336	210139	991
-128973.336	921023	782	-128973.336	215353	164
-128973.336	710154	831	-128973.336	809240	543
-128973.336	376411	473	-128973.336	432513	803
-128973.336	574942	558	-128973.336	298275	418
-128973.336	804660	260	-128973.336	23012	352
-128973.336	393232	152	-128973.336	979450	913
-128973.336	595759	997	-128973.336	118421	139
-128973.336	364676	27	-128973.336	857122	889
-128973.336	629320	222	-128973.336	669639	699
-128973.336	350608	334	-128973.336	107446	12
-128973.336	700349	401	-128973.336	252949	487
-128973.336	167409	772	-128973.336	417035	149
-128973.336	195763	358	-128973.336	343926	624
-128973.336	147440	514	-128973.336	39136	226
-128973.336	603842	61	-128973.336	696830	668
-128973.336	608849	224	-128973.336	597614	284
-128973.336	415931	10	-128973.336	213532	503
-128973.336	652266	490	-128973.336	551340	766
-128973.336	900631	774	-128973.336	22075	659
-128973.336	231281	542	-128973.336	519357	559
-128973.336	922596	456	-128973.336	404426	344
-128973.336	154575	539	-128973.336	79945	395

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-128973.336 164305	128	-128973.336 871851	257
-128973.336 760599	832	-128973.336 76974	16
-128973.336 387701	275	-128973.336 561664	392
-128973.336 792993	859	-128973.336 202790	198
-128973.336 869564	869	-128973.336 150531	154
-128973.336 638577	769	-128973.336 237332	661
-128973.336 407168	44	-128973.336 252346	528
-128973.336 863094	147	-128973.336 62835	642
-128973.336 217744	326	-128973.336 312754	562
-128973.336 782821	272	-128973.336 475420	71
-128973.336 805935	615	-128973.336 704798	530
-128973.336 173191	422	-128973.336 636396	168
-128973.336 81233	825	-128973.336 299977	956
-128973.336 749453	33	-128973.336 456213	160
-128973.336 724087	925	-128973.336 79212	517
-128973.336 752769	253	-128973.336 437181	135
-128973.336 702492	718	-128973.336 640833	434
-128973.336 547702	767	-128973.336 76451	211
-128973.336 988761	475	-128973.336 529496	343
-128973.336 224151	973	-128973.336 441191	361
-128973.336 679448	937	-128973.336 267983	228
-128973.336 724519	333	-128973.336 246794	801
-128973.336 246261	38	-128973.336 225995	847
-128973.336 977800	853	-128973.336 568405	233
-128973.336 220454	288	-128973.336 575700	100
-128973.336 648555	113	-128973.336 345974	622
-128973.336 413564	535	-128973.336 726744	939
-128973.336 476295	969	-128973.336 920593	611
-128973.336 138695	783	-128973.336 379823	905
-128973.336 860102	495	-128973.336 824956	607
-128973.336 781190	219	-128973.336 645052	910
-128973.336 341041	34	-128973.336 742688	594
-128973.336 127362	757		
-128973.336 418686	338		
-128973.336 152496	123		
-128973.336 46437	153		
-128973.336 830292	527		
-128973.336 211281	292		
-128973.336 995913	787		
-128973.336 590834	785		
-128973.336 192071	142		
-128973.336 188640	451		
-128973.336 928624	981		
-128973.336 671390	814		
-128973.336 140849	515		
-128973.336 606576	151		
-128973.336 887580	493		
-128973.336 497522	502		
-128973.336 302046	863		
-128973.336 626208	698		
-128973.336 505244	582		
-128973.336 602032	648		
-128973.336 802256	477		
-128973.336 971853	402		
-128973.336 589483	950		
-128973.336 373702	669		
-128973.336 534864	307		
-128973.336 650371	14		
-128973.336 824126	287		
-128973.336 781489	627		
-128973.336 669634	335		
-128973.336 526324	178		
-128973.336 377584	630		
-128973.336 271809	846		
-128973.336 848331	137		
-128973.336 712702	684		
-128973.336 451258	848		
-128973.336 856612	700		
-128973.336 278661	674		

THE BEST LOGLIKELIHOOD VALUE HAS BEEN REPLICATED. RERUN WITH AT LEAST TWICE THE RANDOM STARTS TO CHECK THAT THE BEST LOGLIKELIHOOD IS STILL OBTAINED AND REPLICATED.

THE MODEL ESTIMATION TERMINATED NORMALLY

MODEL FIT INFORMATION

Number of Free Parameters	18
Loglikelihood	
H0 Value	-128973.336
H0 Scaling Correction Factor for MLR	1.3080

Information Criteria

Akaike (AIC)	257982.672
Bayesian (BIC)	258130.863
Sample-Size Adjusted BIC	258073.660
(n* = (n + 2) / 24)	

FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THE ESTIMATED MODEL

Latent Classes		
1	3094.62154	0.11131
2	6716.69965	0.24159
3	17990.67881	0.64710

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				Estimate	S.E.	Est./S.E.	P-Value
FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON ESTIMATED POSTERIOR PROBABILITIES				Latent Class 1			
Latent Classes				Means			
1	3094.62154	0.11131		SLPNSSNC	1.108	0.014	79.573 0.000
2	6716.69965	0.24159		WORRYNC	-2.040	0.017	-118.142 0.000
3	17990.67881	0.64710		SOLC	1.488	0.034	43.966 0.000
				SLPDURC	1.192	0.024	49.384 0.000
				Variances			
				SLPNSSNC	0.465	0.003	155.639 0.000
				WORRYNC	0.184	0.003	68.676 0.000
				SOLC	0.597	0.008	76.749 0.000
				SLPDURC	0.677	0.007	95.747 0.000
FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THEIR MOST LIKELY LATENT CLASS MEMBERSHIP				Latent Class 2			
Class Counts and Proportions				Means			
Latent Classes				SLPNSSNC	0.982	0.010	98.325 0.000
1	3065	0.11024		WORRYNC	-0.687	0.022	-31.840 0.000
2	6569	0.23628		SOLC	0.444	0.015	30.251 0.000
3	18168	0.65348		SLPDURC	0.517	0.015	34.921 0.000
				Variances			
				SLPNSSNC	0.465	0.003	155.639 0.000
				WORRYNC	0.184	0.003	68.676 0.000
				SOLC	0.597	0.008	76.749 0.000
				SLPDURC	0.677	0.007	95.747 0.000
CLASSIFICATION QUALITY				Latent Class 3			
Entropy	0.863			Means			
Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)				SLPNSSNC	0.885	0.005	167.165 0.000
	1	2	3	WORRYNC	0.607	0.005	127.737 0.000
1	0.921	0.079	0.000	SOLC	-0.416	0.007	-61.368 0.000
2	0.041	0.880	0.078	SLPDURC	-0.395	0.008	-51.870 0.000
3	0.000	0.038	0.962	Variances			
				SLPNSSNC	0.465	0.003	155.639 0.000
				WORRYNC	0.184	0.003	68.676 0.000
				SOLC	0.597	0.008	76.749 0.000
				SLPDURC	0.677	0.007	95.747 0.000
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)				Categorical Latent Variables			
	1	2	3	Means			
1	0.912	0.088	0.000	C#1	-1.760	0.034	-51.148 0.000
2	0.036	0.861	0.103	C#2	-0.985	0.019	-50.760 0.000
3	0.000	0.029	0.971	QUALITY OF NUMERICAL RESULTS			
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)				Condition Number for the Information Matrix			
	1	2	3	0.514E-02			
1	13.724	11.383	0.000	(ratio of smallest to largest eigenvalue)			
2	-1.051	2.123	0.000	TECHNICAL 11 OUTPUT			
3	-13.786	-3.525	0.000	Random Starts Specifications for the k-1 Class Analysis Model			
MODEL RESULTS				Number of initial stage random starts			
				1000			
				Number of final stage optimizations			
				250			
				VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES			
Two-Tailed							

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H0 Loglikelihood Value	-131941.399	Histograms (sample values)
2 Times the Loglikelihood Difference	5936.126	Scatterplots (sample values)
Difference in the Number of Parameters	5	Sample means
Mean	14.458	Estimated means, medians, modes, and percentiles
Standard Deviation	30.006	Sample and estimated means
P-Value	0.0000	Observed individual values
		Estimated means and observed individual values
		Estimated overall and class-specific distributions
LO-MENDELL-RUBIN ADJUSTED LRT TEST		
Value	5822.329	SAVEDATA INFORMATION
P-Value	0.0000	Save file
		Step1_LPA3M1.dat
TECHNICAL 14 OUTPUT		
Order and format of variables		
Random Starts Specifications for the k-1 Class Analysis		
Model		SLPNSSNC F10.3
Number of initial stage random starts	1000	WORRYNC F10.3
Number of final stage optimizations	250	SOLC F10.3
		SLPDURC F10.3
		CPROB1 F10.3
		CPROB2 F10.3
		CPROB3 F10.3
		C F10.3
		ID I6
Random Starts Specification for the k-1 Class Model for Generated Data		
Number of initial stage random starts	2	
Number of final stage optimizations	1	
Random Starts Specification for the k Class Model for Generated Data		
Number of initial stage random starts	50	Save file format
Number of final stage optimizations	10	8F10.3 I6
Number of bootstrap draws requested	250	
		Save file record length 10000
		Save missing symbol *
PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES		
H0 Loglikelihood Value	-131941.399	DIAGRAM INFORMATION
2 Times the Loglikelihood Difference	5936.126	Mplus diagrams are currently not available for Mixture analysis.
Difference in the Number of Parameters	5	No diagram output was produced.
Approximate P-Value	0.0000	
Successful Bootstrap Draws	250	
WARNING: OF THE 250 BOOTSTRAP DRAWS, 152 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 3-CLASS MODEL.		
THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA.		
INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.		
Beginning Time: 11:26:26		
Ending Time: 12:34:43		
Elapsed Time: 01:08:17		
MUTHEN & MUTHEN		
3463 Stoner Ave.		
Los Angeles, CA 90066		
Tel: (310) 391-9971		
Fax: (310) 391-8971		
Web: www.StatModel.com		
Support: Support@StatModel.com		
PLOT INFORMATION		
The following plots are available:		
Copyright (c) 1998-2021 Muthen & Muthen		

Parametrization Model 2 - Varying variances and zero covariance

Mplus VERSION 8.7	FILE IS mplus2.csv;
MUTHEN & MUTHEN	! Specifies file location for data file. Make sure data is in format appropriate for Mplus
07/11/2023 1:23 PM	! per Mplus manual. This data file is in individual format (one row of data per participant)
INPUT INSTRUCTIONS	VARIABLE:
TITLE: LPA 3 profile syntax: M2 Unconstrained variance, 0 covariance	NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol slpDur gender genderN yrgrp yrgrpN SCI_02C slpnssNC worryNC solC slpDurC;
DATA:	

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be
listed here
IDVARIABLE IS ID;
CLASSES = c (3);
! This is where you instruct Mplus on how many
classes/profiles are being estimated. Initi
! model contains only one class/profile, thus it would be
CLASSES = c (1). Above specifies
!profiles, and for each further iterative models the number
in parentheses increases by on
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(9999);
! Used to communicate how missing data is coded in data
file. Here shown with a "." which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 1000 250;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifie
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will sto
!before reaching 500 iterations.
!These values can be increased ... see "Four-Profile Final
Model with Covariate Analysis
!Syntax" for an example.
LRTSTARTS = 2 1 50 10;
LRTBOOTSTRAP = 250;
!The above start values are for the defaults for the LRT
statistic being run to compare th
!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statemen
!specifies the number of bootstrap draws to inform Mplus'
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances
!profiles to be equal.

%c#1%

[slpnssNC worryNC solC slpDurC];
slpnssNC worryNC solC slpDurC;
%c#2%
[slpnssNC worryNC solC slpDurC];
slpnssNC worryNC solC slpDurC;
%c#3%
[slpnssNC worryNC solC slpDurC];
slpnssNC worryNC solC slpDurC;

OUTPUT:
TECH11 TECH14;
! TECH1 provides parameter specifications and starting
values for the analysis
! TECH8 provides optimization history for this analysis type
!TECH11 provides LRT results
!TECH14 provides bootstrapped LRT test
PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);
TYPE=PLOT3;
SAVEDATA:
FILE IS Step1_LPA3M2.dat;
! Tells Mplus where to save the output files from the
analysis
SAVE = CPROBABILITIES;
! The above command lines are to save the most likely
profile membership for each particip
! and the posterior probabilities for their membership in
each latent profile.

*** WARNING in MODEL command
All variables are uncorrelated with all other variables within
class.
Check that this is what is intended.

LPA 3 profile syntax: m2 Unconstrained variance, 0
covariance

SUMMARY OF ANALYSIS

Number of groups 1
Number of observations 27802

Number of dependent variables 4
Number of independent variables 0
Number of continuous latent variables 0
Number of categorical latent variables 1

Observed dependent variables

Continuous
SLPNSSNC WORRYNC SOLC SLPDURC

Categorical latent variables
C

Variables with special functions

ID variable ID

Estimator MLR
Information matrix OBSERVED
Optimization Specifications for the Quasi-Newton Algorithm
for
Continuous Outcomes
Maximum number of iterations 100

```

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Convergence criterion	0.100D-05	UNIVARIATE HIGHER-ORDER MOMENT	
Optimization Specifications for the EM Algorithm		DESCRIPTIVE STATISTICS	
Maximum number of iterations	500		
Convergence criteria		Variable/ Mean/ Skewness/ Minimum/ % with Percentiles	
Loglikelihood change	0.100D-06	Sample Size Variance Kurtosis Maximum	
Relative loglikelihood change	0.100D-06	Min/Max 20%/60% 40%/80% Median	
Derivative	0.100D-05		
Optimization Specifications for the M step of the EM Algorithm for		SLPNSSNC 0.933 0.086 0.000 27.10%	
Categorical Latent variables		0.000 1.000 1.000	
Number of M step iterations	1	27802.000 0.471 -0.878 2.000 20.40%	
M step convergence criterion	0.100D-05	1.000 2.000	
Basis for M step termination	ITERATION	WORRYNC 0.000 -1.036 -2.825 1.71%	
Optimization Specifications for the M step of the EM Algorithm for		-0.928 0.020 0.495	
Censored, Binary or Ordered Categorical (Ordinal), Unordered		27802.000 1.000 0.165 0.969 28.55%	
Categorical (Nominal) and Count Outcomes		0.495 0.969	
Number of M step iterations	1	SOLC 0.000 0.967 -1.054 31.63%	
M step convergence criterion	0.100D-05	1.054 -0.072 -0.072	
Basis for M step termination	ITERATION	27123.000 1.000 0.467 2.875 2.77%	
Maximum value for logit thresholds	15	-0.072 0.910	
Minimum value for logit thresholds	-15	SLPDURC 0.000 0.568 -1.205 25.70%	
Minimum expected cell size for chi-square	0.100D-01	-1.205 -0.292 -0.292	
Maximum number of iterations for H1	2000	26763.000 1.000 -0.470 2.445 3.44%	
Convergence criterion for H1	0.100D-03	-0.292 0.620	
Optimization algorithm	EMA	RANDOM STARTS RESULTS RANKED FROM THE BEST TO THE WORST LOGLIKELIHOOD VALUES	
Random Starts Specifications		Unperturbed starting value run did not converge in the initial stage	
Number of initial stage random starts	1000	optimizations.	
Number of final stage optimizations	250		
Number of initial stage iterations	500		
Initial stage convergence criterion	0.100D+01	999 perturbed starting value run(s) did not converge in the initial stage	
Random starts scale	0.500D+01	optimizations.	
Random seed for generating random starts	0		
Input data file(s)		Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:	
mplus2.csv			
Input data format	FREE		
SUMMARY OF DATA		250 perturbed starting value run(s) did not converge or were rejected in the third stage.	
Number of missing data patterns	4		
Number of y missing data patterns	4		
Number of u missing data patterns	0		
COVARIANCE COVERAGE OF DATA		THE ESTIMATED COVARIANCE MATRIX FOR THE Y VARIABLES IN CLASS 1 COULD NOT BE INVERTED. PROBLEM INVOLVING VARIABLE SLPDURC. COMPUTATION COULD NOT BE COMPLETED IN ITERATION 17. CHANGE YOUR MODEL AND/OR STARTING VALUES. THIS MAY BE DUE TO A ZERO ESTIMATED VARIANCE, THAT IS, NO WITHIN-CLASS VARIATION FOR THE VARIABLE. THE MODEL ESTIMATION DID NOT TERMINATE NORMALLY. ESTIMATES CANNOT BE TRUSTED.	
Minimum covariance coverage value	0.100		
PROPORTION OF DATA PRESENT FOR Y			
Covariance Coverage			
SLPNSSNC			
WORRYNC			
SOLC			
SLPDURC			
SLPNSSNC	1.000		
WORRYNC	1.000	1.000	
SOLC	0.976	0.976	0.976
SLPDURC	0.963	0.963	0.940 0.963
UNIVARIATE SAMPLE STATISTICS		THE MODEL ESTIMATION DID NOT TERMINATE NORMALLY DUE TO AN ERROR IN THE COMPUTATION. CHANGE YOUR MODEL AND/OR STARTING VALUES.	
		MODEL RESULTS	
		Estimate	
		Latent Class 1	

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

```

Means
SLPNSSNC      -0.197
WORRYNC       55.382
SOLC          -1907.460
SLPDURC      -266.045

Variances
SLPNSSNC       6.868
WORRYNC       37.105
SOLC          3564.902
SLPDURC        0.000

Latent Class 2

Means
SLPNSSNC       0.508
WORRYNC       -0.016
SOLC          -0.134
SLPDURC       1.084

Variances
SLPNSSNC       0.302
WORRYNC       0.399
SOLC           0.078
SLPDURC       0.598

Latent Class 3

Means
SLPNSSNC       0.941
WORRYNC        0.000
SOLC           0.003
SLPDURC      -0.020

Variances
SLPNSSNC       0.470
WORRYNC       1.011
SOLC           1.015
SLPDURC       0.986

Categorical Latent Variables

Means
C#1          -233.972
C#2          -3.905

MODEL COMMAND WITH FINAL ESTIMATES USED AS
STARTING VALUES

%OVERALL%

[ c#1*-233.97177 ];
[ c#2*-3.90489 ];

%C#1%

[ slpnssnc*-0.19736 ];
[ worrync*55.38240 ];
[ solc*-1907.45996 ];
[ slpdurc*-266.04523 ];

slpnssnc*6.86776;
worrync*37.10458;
solc*3564.90186;
slpdurc*0.00000;

%C#2%

[ slpnssnc*0.50800 ];
[ worrync*-0.01614 ];
[ solc*-0.13390 ];
[ slpdurc*1.08362 ];

slpnssnc*0.30176;
worrync*0.39880;
solc*0.07827;
slpdurc*0.59777;

%C#3%

[ slpnssnc*0.94078 ];
[ worrync*0.00030 ];
[ solc*0.00289 ];
[ slpdurc*-0.01953 ];

slpnssnc*0.47027;
worrync*1.01098;
solc*1.01502;
slpdurc*0.98561;

TECHNICAL 11 OUTPUT

Random Starts Specifications for the k-1 Class Analysis
Model
Number of initial stage random starts      1000
Number of final stage optimizations        250

TECHNICAL 14 OUTPUT

Random Starts Specifications for the k-1 Class Analysis
Model
Number of initial stage random starts      1000
Number of final stage optimizations        250

Random Starts Specification for the k-1 Class Model for
Generated Data
Number of initial stage random starts      2
Number of final stage optimizations        1
Random Starts Specification for the k Class Model for
Generated Data
Number of initial stage random starts      50
Number of final stage optimizations        10
Number of bootstrap draws requested        250

SAVEDATA INFORMATION

Class probabilities were not computed.
No data were saved.

DIAGRAM INFORMATION

Mplus diagrams are currently not available for Mixture
analysis.
No diagram output was produced.

Beginning Time: 13:23:15
Ending Time: 13:26:44
Elapsed Time: 00:03:29

MUTHEN & MUTHEN
3463 Stoner Ave.
Los Angeles, CA 90066

```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

Tel: (310) 391-9971
 Fax: (310) 391-8971
 Web: www.StatModel.com
 Support: Support@StatModel.com

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Parametrization Model 3 - Equal variances and equal covariances

Mplus VERSION 8.7
 MUTHEN & MUTHEN
 07/11/2023 1:29 PM

INPUT INSTRUCTIONS

TITLE: LPA 3 profile syntax: M3 constrained variance and equal covariance

DATA:

FILE IS mplus2.csv;

! Specifies file location for data file. Make sure data is in format appropriate for Mplus

! per Mplus manual. This data file is in individual format (one row of data per participant)

VARIABLE:

NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol slpDur genderN yrgrp yrgrpN SCI_02C slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.

USEVARIABLES ARE slpnssNC worryNC solC slpDurC;

! Only variables intended for use in the analysis should be listed here

IDVARIABLE IS ID;

CLASSES = c (3);

! This is where you instruct Mplus on how many classes/profiles are being estimated. Initi

! model contains only one class/profile, thus it would be CLASSES = c (1). Above specifies

!profiles, and for each further iterative models the number in parentheses increases by on

!three profiles/classes would be c (3), and so on.

MISSING ARE ALL(9999);

! Used to communicate how missing data is coded in data file. Here shown with a “.” which

! all that is included in each cell with missing data in the data file

ANALYSIS:

TYPE = MIXTURE;

! LPA is a version of mixture modeling, and this instructs Mplus to analyze in this way

ESTIMATOR = MLR;

!FIML robust to non-normal data

STARTS = 1000 250;

STITERATIONS = 500;

! Default number of starts for each step of the ML estimation. First STARTS value specifies

!number of unique start values to start with, the 250 represents the 250 best unique start

!carrying forward to completion. The STITERATIONS specifies the number of ML iteration

!steps for those 250 selected start values to go through to be able to converge. This is a

!maximum number of iteration; if a model converges in less than 500 iterations it will stop

!before reaching 500 iterations.

!These values can be increased ... see “Four-Profile Final Model with Covariate Analysis

!Syntax” for an example.

LRTSTARTS = 2 1 50 10;

LRTBOOTSTRAP = 250;

!The above start values are for the defaults for the LRT statistic being run to compare th

!model fit with the model fit of a model with one less class (k-1). The BOOTSTRAP statemen

!specifies the number of bootstrap draws to inform Mplus' bootstrapped LRT results.

MODEL:

!For a default Mplus model the LPA model does not need to be specified. However, it can be

!The model can also be modified from the Mplus default of estimating the indicator means

!(uniquely across profiles) and variances (constrained across profiles), as well as the la

!profile mean.

%OVERALL%

[slpnssNC worryNC solC slpDurC]; !estimates the indicators means for each profile. Without

!the means are freely estimated in each profile, not constrained.

slpnssNC worryNC solC slpDurC; !Label Var1-Var5 constrains the estimates of the variances

!profiles to be equal.

!covariances

slpnssNC WITH worryNC solC slpDurC;

worryNC WITH solC slpDurC;

solC WITH slpDurC;

OUTPUT:

TECH11 TECH14;

! TECH1 provides parameter specifications and starting values for the analysis

! TECH8 provides optimization history for this analysis type

!TECH11 provides LRT results

!TECH14 provides bootstrapped LRT test

PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);

TYPE=PLOT3;

SAVEDATA:

FILE IS Step1_LPA3M3.dat;

! Tells Mplus where to save the output files from the analysis

SAVE = CPROBABILITIES;

! The above command lines are to save the most likely profile membership for each participant

! and the posterior probabilities for their membership in each latent profile.

LPA 3 profile syntax: M3 constrained variance and equal covariance

SUMMARY OF ANALYSIS

Number of groups	1
Number of observations	27802
Number of dependent variables	4
Number of independent variables	0
Number of continuous latent variables	0
Number of categorical latent variables	1
Observed dependent variables	

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Continuous	SLPNSSNC	WORRYNC	SOLC	SLPDURC	SLPNSSNC	WORRYNC	SOLC
Categorical latent variables					SLPNSSNC	1.000	
C					WORRYNC	1.000	1.000
					SOLC	0.976	0.976 0.976
Variables with special functions					SLPDURC	0.963	0.963 0.940 0.963
ID variable	ID				UNIVARIATE SAMPLE STATISTICS		
Estimator			MLR		UNIVARIATE HIGHER-ORDER MOMENT		
Information matrix			OBSERVED		DESCRIPTIVE STATISTICS		
Optimization Specifications for the Quasi-Newton Algorithm for					Variable/	Mean/	Skewness/
Continuous Outcomes					Sample Size	Variance	Kurtosis
Maximum number of iterations			100		Minimum/	% with	
Convergence criterion			0.100D-05		Percentiles	Maximum	
Optimization Specifications for the EM Algorithm					Min/Max	20%/60%	40%/80% Median
Maximum number of iterations			500		SLPNSSNC	0.933	0.086 0.000 27.10%
Convergence criteria					0.000 1.000 1.000		
Loglikelihood change			0.100D-06		27802.000	0.471	-0.878 2.000 20.40%
Relative loglikelihood change			0.100D-06		1.000 2.000		
Derivative			0.100D-05		WORRYNC	0.000	-1.036 -2.825 1.71%
Optimization Specifications for the M step of the EM					-0.928 0.020 0.495		
Algorithm for					27802.000 1.000 0.165	0.969	28.55%
Categorical Latent variables					0.495 0.969		
Number of M step iterations			1		SOLC	0.000	0.967 -1.054 31.63%
M step convergence criterion			0.100D-05		1.054 -0.072 -0.072		
Basis for M step termination			ITERATION		27123.000 1.000 0.467	2.875	2.77%
Optimization Specifications for the M step of the EM					-0.072 0.910		
Algorithm for					SLPDURC	0.000	0.568 -1.205 25.70%
Censored, Binary or Ordered Categorical (Ordinal),					-1.205 -0.292 -0.292		
Unordered					26763.000 1.000 -0.470	2.445	3.44%
Categorical (Nominal) and Count Outcomes					-0.292 0.620		
Number of M step iterations			1		RANDOM STARTS RESULTS RANKED FROM THE BEST		
M step convergence criterion			0.100D-05		TO THE WORST LOGLIKELIHOOD VALUES		
Basis for M step termination			ITERATION		8 perturbed starting value run(s) did not converge in the		
Maximum value for logit thresholds			15		initial stage		
Minimum value for logit thresholds			-15		optimizations.		
Minimum expected cell size for chi-square			0.100D-01		Final stage loglikelihood values at local maxima, seeds, and		
Maximum number of iterations for H1			2000		initial stage start numbers:		
Convergence criterion for H1			0.100D-03		-126772.383	349562	359
Optimization algorithm			EMA		-126772.383	869564	869
Random Starts Specifications					-126772.383	347515	24
Number of initial stage random starts			1000		-126772.383	848331	137
Number of final stage optimizations			250		-126772.383	588923	606
Number of initial stage iterations			500		-126772.383	781489	627
Initial stage convergence criterion			0.100D+01		-126772.383	440841	118
Random starts scale			0.500D+01		-126772.383	77571	980
Random seed for generating random starts			0		-126772.383	407168	44
Input data file(s)					-126772.383	351807	617
mplus2.csv					-126772.383	63231	935
Input data format			FREE		-126772.383	631413	439
SUMMARY OF DATA					-126772.383	415931	10
Number of missing data patterns			4		-126772.383	294811	637
Number of y missing data patterns			4		-126772.383	354208	196
Number of u missing data patterns			0		-126772.383	319144	176
COVARIANCE COVERAGE OF DATA					-126772.383	535804	111
Minimum covariance coverage value			0.100		-126772.383	57226	208
PROPORTION OF DATA PRESENT FOR Y					-126772.383	782821	272
Covariance Coverage					-126772.383	967902	52
					-126772.383	173191	422
					-126772.383	444228	860
					-126772.383	251641	784
					-126772.383	669639	699
					-126772.383	370957	554

Scripts & Outputs *“Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”*

-126772.383	965994	396	-126772.383	424223	900
-126772.383	995648	936	-126772.383	66276	217
-126772.383	592219	119	-126772.383	939021	8
-126772.383	617658	660	-126772.383	939709	112
-126772.383	634782	979	-126772.383	244349	736
-126772.383	140442	500	-126772.383	804561	59
-126772.383	213189	605	-126772.383	501995	791
-126772.383	505244	582	-126772.383	569131	26
-126772.383	471398	74	-126772.383	391179	78
-126772.383	860102	495	-126772.383	529455	268
-126772.383	567165	319	-126772.383	366533	484
-126772.383	226322	478	-126772.383	404426	344
-126772.383	260601	36	-126772.383	520865	763
-126772.383	147440	514	-126772.383	266340	683
-126772.383	85114	385	-126772.383	836515	341
-126772.383	965639	463	-126772.383	622860	259
-126772.383	887676	22	-126772.383	748692	204
-126772.383	164305	128	-126772.383	462821	745
-126772.383	50887	389	-126772.383	202790	198
-126772.383	422103	62	-126772.383	638577	769
-126772.383	425149	878	-126772.383	344422	296
-126772.383	777492	972	-126772.383	118438	601
-126772.383	98068	998	-126772.383	79212	517
-126772.383	636396	168	-126772.383	539389	544
-126772.383	414828	322	-126772.383	328661	813
-126772.383	417035	149	-126772.383	488125	368
-126772.383	609089	241	-126772.383	714997	399
-126772.383	859432	770	-126772.383	227563	63
-126772.383	406935	958	-126772.383	686482	215
-126772.383	94610	884	-126772.383	213532	503
-126772.383	387701	275	-126772.383	441191	361
-126772.383	473343	844	-126772.383	973369	202
-126772.383	579995	183	-126772.383	105435	265
-126772.383	726744	939	-126772.383	434915	552
-126772.383	251680	794	-126772.383	712702	684
-126772.383	68850	462	-126772.383	431271	943
-126772.383	760850	739	-126772.383	284716	713
-126772.383	677062	680	-126772.383	136842	58
-126772.383	642386	662	-126772.383	65651	214
-126772.383	425982	485	-126772.383	850112	922
-126772.383	370466	41	-126772.383	130011	587
-126772.383	484687	306	-126772.383	40327	971
-126772.383	508482	446	-126772.383	12477	155
-126772.383	648555	113	-126772.383	926797	406
-126772.383	502157	799	-126772.383	783165	170
-126772.383	423661	875	-126772.383	455617	242
-126772.383	195353	225	-126772.383	610181	711
-126772.383	93468	3	-126772.383	271809	846
-126772.383	650371	14	-126772.383	193569	440
-126772.383	972873	157	-126772.383	858804	534
-126772.383	742688	594	-126772.383	402699	604
-126772.383	89970	223	-126772.383	203508	806
-126772.383	194143	510	-126772.383	4091	548
-126772.383	884242	957	-126772.383	942358	644
-126772.383	343926	624	-127270.473	638977	643
-126772.383	903633	553	-127270.473	210870	383
-126772.383	891347	504	-127270.473	476498	179
-126772.383	131856	90	-127270.473	584397	428
-126772.383	40340	188	-127270.473	789528	837
-126772.383	22089	143	-127270.473	945065	255
-126772.383	326091	759	-127270.473	224950	455
-126772.383	210139	991	-127270.473	476295	969
-126772.383	917702	693	-127270.473	608496	4
-126772.383	322790	636	-127270.473	739214	807
-126772.383	549244	756	-127270.473	534483	290
-126772.383	838615	677	-127270.473	354624	448
-126772.383	966499	963	-127270.473	902278	21
-126772.383	617243	237	-127270.473	137377	397
-126772.383	628143	854	-127270.473	691041	590
-126772.383	36714	201	-127270.473	625191	350

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-127270.473	303834	798	-127270.473	845580	805
-127270.473	963967	941	-127270.473	484406	421
-127270.473	152496	123	-127270.473	643311	888
-127270.473	392751	480	-127270.473	259507	53
-127270.473	666720	671	-127270.473	688839	273
-127270.473	967237	48	-127270.473	915642	40
-127270.473	341519	653	-127270.473	207896	25
-127270.473	188498	258	-127270.473	207917	976
-127270.473	848590	694	-127270.473	964570	701
-127270.473	985387	381	-127270.473	153053	378
-127270.473	599729	658	-127270.473	311214	64
-127270.473	582296	452	-127270.473	531546	702
-127270.473	396614	918	-127270.473	220454	288
-127270.473	618000	190	-127270.473	161421	519
-127270.473	741888	138	-127270.473	248742	556
-127270.473	59674	240			
-127270.473	67009	564			
-127270.473	879211	453	THE BEST LOGLIKELIHOOD VALUE HAS BEEN		
-127270.473	995249	525	REPLICATED. RERUN WITH AT LEAST TWICE THE		
-127270.473	70118	104	RANDOM STARTS TO CHECK THAT THE BEST		
-127270.473	690596	858	LOGLIKELIHOOD IS STILL OBTAINED AND		
-127270.473	539751	459	REPLICATED.		
-127270.473	286735	175			
-127270.473	49910	829	THE MODEL ESTIMATION TERMINATED NORMALLY		
-127270.473	977800	853			
-127270.473	216565	474	MODEL FIT INFORMATION		
-127270.473	561664	392			
-127270.473	214681	824	Number of Free Parameters	24	
-127270.473	851945	18			
-127270.473	872743	374	Loglikelihood		
-127270.473	563584	657			
-127270.473	807534	850	H0 Value	-126772.383	
-127270.473	392418	28	H0 Scaling Correction Factor	1.2282	
-127270.473	238906	430	for MLR		
-127270.473	939870	655			
-127270.473	126371	526	Information Criteria		
-127270.473	302046	863			
-127270.473	999211	628	Akaike (AIC)	253592.765	
-127270.473	364676	27	Bayesian (BIC)	253790.354	
-127270.473	137888	901	Sample-Size Adjusted BIC	253714.083	
-127270.473	158612	581	(n* = (n + 2) / 24)		
-127270.473	345726	461			
-127270.473	736574	414	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-127270.473	544048	87	LATENT CLASSES		
-127270.473	978061	488	BASED ON THE ESTIMATED MODEL		
-127270.473	849670	347			
-127270.473	648035	836	Latent		
-127270.473	783102	433	Classes		
-127270.473	58507	793	1	3218.02371	0.11575
-127270.473	73576	213	2	18879.70470	0.67908
-127270.473	377504	294	3	5704.27159	0.20517
-127270.473	436892	565			
-127270.473	758647	951	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-127270.473	347222	533	LATENT CLASSES		
-127270.473	489927	520	BASED ON ESTIMATED POSTERIOR PROBABILITIES		
-127270.473	unperturbed	0			
-127270.473	529496	343	Latent		
-127270.473	154575	539	Classes		
-127270.473	732596	320	1	3218.02371	0.11575
-127270.473	871438	561	2	18879.70470	0.67908
-127270.473	685657	69	3	5704.27159	0.20517
-127270.473	11984	934			
-127270.473	3607	873			
-127270.473	77210	712	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-127270.473	863094	147	LATENT CLASSES		
-127270.473	791678	974	BASED ON THEIR MOST LIKELY LATENT CLASS		
-127270.473	283492	435	MEMBERSHIP		
-127270.473	496710	386			
-127270.473	265218	924	Class Counts and Proportions		
-127270.473	402224	91			

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Latent Classes					SLPNSSNC	0.469	0.003	156.716	0.000
					WORRYNC	0.168	0.002	89.473	0.000
1	3167	0.11391			SOLC	0.754	0.009	88.524	0.000
2	18928	0.68081			SLPDURC	0.790	0.007	107.265	0.000
3	5707	0.20527							
CLASSIFICATION QUALITY					Latent Class 2				
Entropy	0.870				SLPNSSNC WITH				
Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)					WORRYNC	-0.018	0.003	-7.135	0.000
	1	2	3		SOLC	0.118	0.004	30.181	0.000
1	0.908	0.000	0.092		SLPDURC	0.030	0.004	7.611	0.000
2	0.000	0.975	0.025		WORRYNC WITH				
3	0.060	0.074	0.866		SOLC	-0.155	0.005	-30.483	0.000
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					SLPDURC	-0.129	0.004	-29.670	0.000
	1	2	3		SOLC WITH				
1	0.894	0.000	0.106		SLPDURC	0.225	0.006	35.821	0.000
2	0.000	0.978	0.022		Means				
3	0.051	0.082	0.867		SLPNSSNC	0.907	0.005	175.707	0.000
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					WORRYNC	0.581	0.004	138.481	0.000
	1	2	3		SOLC	-0.314	0.006	-48.571	0.000
1	2.131	-11.006	0.000		SLPDURC	-0.301	0.007	-43.319	0.000
2	-10.015	3.778	0.000		Variances				
3	-2.834	-2.352	0.000		SLPNSSNC	0.469	0.003	156.716	0.000
MODEL RESULTS					WORRYNC	0.168	0.002	89.473	0.000
					SOLC	0.754	0.009	88.524	0.000
					SLPDURC	0.790	0.007	107.265	0.000
					Latent Class 3				
					SLPNSSNC WITH				
					WORRYNC	-0.018	0.003	-7.135	0.000
					SOLC	0.118	0.004	30.181	0.000
					SLPDURC	0.030	0.004	7.611	0.000
					WORRYNC WITH				
					SOLC	-0.155	0.005	-30.483	0.000
					SLPDURC	-0.129	0.004	-29.670	0.000
					SOLC WITH				
					SLPDURC	0.225	0.006	35.821	0.000
					Means				
					SLPNSSNC	0.959	0.011	88.715	0.000
					WORRYNC	-0.777	0.017	-45.568	0.000
					SOLC	0.444	0.017	26.141	0.000
					SLPDURC	0.491	0.017	29.459	0.000
					Variances				
					SLPNSSNC	0.469	0.003	156.716	0.000
					WORRYNC	0.168	0.002	89.473	0.000
					SOLC	0.754	0.009	88.524	0.000
					SLPDURC	0.790	0.007	107.265	0.000
					Categorical Latent Variables				
					Means				
					C#1	-0.572	0.038	-15.206	0.000
					C#2	1.197	0.019	62.898	0.000
					QUALITY OF NUMERICAL RESULTS				
					Condition Number for the Information Matrix				
					0.123E-02				
					(ratio of smallest to largest eigenvalue)				
					TECHNICAL 11 OUTPUT				
					Means				
					SLPNSSNC	1.037	0.014	72.590	0.000
					WORRYNC	-2.034	0.019	-106.696	0.000
					SOLC	1.096	0.033	33.163	0.000
					SLPDURC	0.922	0.026	35.171	0.000
					Variances				

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<p>Random Starts Specifications for the k-1 Class Analysis Model</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>1000</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>250</td> </tr> </table> <p>VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES</p> <table border="0"> <tr> <td>H0 Loglikelihood Value</td> <td>-128233.817</td> </tr> <tr> <td>2 Times the Loglikelihood Difference</td> <td>2922.868</td> </tr> <tr> <td>Difference in the Number of Parameters</td> <td>5</td> </tr> <tr> <td>Mean</td> <td>12.618</td> </tr> <tr> <td>Standard Deviation</td> <td>15.491</td> </tr> <tr> <td>P-Value</td> <td>0.0000</td> </tr> </table> <p>LO-MENDELL-RUBIN ADJUSTED LRT TEST</p> <table border="0"> <tr> <td>Value</td> <td>2866.836</td> </tr> <tr> <td>P-Value</td> <td>0.0000</td> </tr> </table> <p>TECHNICAL 14 OUTPUT</p> <p>Random Starts Specifications for the k-1 Class Analysis Model</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>1000</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>250</td> </tr> </table> <p>Random Starts Specification for the k-1 Class Model for Generated Data</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>2</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>1</td> </tr> </table> <p>Random Starts Specification for the k Class Model for Generated Data</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>50</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>10</td> </tr> <tr> <td>Number of bootstrap draws requested</td> <td>250</td> </tr> </table> <p>PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES</p> <table border="0"> <tr> <td>H0 Loglikelihood Value</td> <td>-128233.817</td> </tr> <tr> <td>2 Times the Loglikelihood Difference</td> <td>2922.868</td> </tr> <tr> <td>Difference in the Number of Parameters</td> <td>5</td> </tr> <tr> <td>Approximate P-Value</td> <td>0.0000</td> </tr> <tr> <td>Successful Bootstrap Draws</td> <td>250</td> </tr> </table> <p>WARNING: OF THE 250 BOOTSTRAP DRAWS, 174 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 3-CLASS MODEL. THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA. INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.</p> <p>PLOT INFORMATION</p>	Number of initial stage random starts	1000	Number of final stage optimizations	250	H0 Loglikelihood Value	-128233.817	2 Times the Loglikelihood Difference	2922.868	Difference in the Number of Parameters	5	Mean	12.618	Standard Deviation	15.491	P-Value	0.0000	Value	2866.836	P-Value	0.0000	Number of initial stage random starts	1000	Number of final stage optimizations	250	Number of initial stage random starts	2	Number of final stage optimizations	1	Number of initial stage random starts	50	Number of final stage optimizations	10	Number of bootstrap draws requested	250	H0 Loglikelihood Value	-128233.817	2 Times the Loglikelihood Difference	2922.868	Difference in the Number of Parameters	5	Approximate P-Value	0.0000	Successful Bootstrap Draws	250	<p>The following plots are available:</p> <ul style="list-style-type: none"> Histograms (sample values) Scatterplots (sample values) Sample means Estimated means, medians, modes, and percentiles Sample and estimated means Observed individual values Estimated means and observed individual values Estimated overall and class-specific distributions <p>SAVEDATA INFORMATION</p> <p>Save file Step1_LPA3M3.dat</p> <p>Order and format of variables</p> <table border="0"> <tr> <td>SLPNSSNC</td> <td>F10.3</td> </tr> <tr> <td>WORRYNC</td> <td>F10.3</td> </tr> <tr> <td>SOLC</td> <td>F10.3</td> </tr> <tr> <td>SLPDURC</td> <td>F10.3</td> </tr> <tr> <td>CPROB1</td> <td>F10.3</td> </tr> <tr> <td>CPROB2</td> <td>F10.3</td> </tr> <tr> <td>CPROB3</td> <td>F10.3</td> </tr> <tr> <td>C</td> <td>F10.3</td> </tr> <tr> <td>ID</td> <td>I6</td> </tr> </table> <p>Save file format 8F10.3 I6</p> <p>Save file record length 10000</p> <p>Save missing symbol *</p> <p>DIAGRAM INFORMATION</p> <p>Mplus diagrams are currently not available for Mixture analysis. No diagram output was produced.</p> <p>Beginning Time: 13:29:05 Ending Time: 14:43:06 Elapsed Time: 01:14:01</p> <p>MUTHEN & MUTHEN 3463 Stoner Ave. Los Angeles, CA 90066</p> <p>Tel: (310) 391-9971 Fax: (310) 391-8971 Web: www.StatModel.com Support: Support@StatModel.com</p> <p>Copyright (c) 1998-2021 Muthen & Muthen</p>	SLPNSSNC	F10.3	WORRYNC	F10.3	SOLC	F10.3	SLPDURC	F10.3	CPROB1	F10.3	CPROB2	F10.3	CPROB3	F10.3	C	F10.3	ID	I6
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Parametrization Model 4 - Varying variances and equal covariances

<p>Mplus VERSION 8.7 MUTHEN & MUTHEN 07/11/2023 3:04 PM</p> <p>INPUT INSTRUCTIONS</p>	<p>TITLE: LPA 3 profile syntax: M4: Varying means, varying variances, and equal covariances</p> <p>DATA: FILE IS mplus2.csv; ! Specifies file location for data file. Make sure data is in format appropriate for Mplus</p>
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Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```

! per Mplus manual. This data file is in individual format
(one row of data per participant)
VARIABLE:
  NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol
slpDur gender genderN yrgrp yrgrpN SCI_02C
slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be
listed here
IDVARIABLE IS ID;
CLASSES = c (3);
! This is where you instruct Mplus on how many
classes/profiles are being estimated. Initi
! model contains only one class/profile, thus it would be
CLASSES = c (1). Above specifies
!profiles, and for each further iterative models the number
in parentheses increases by one
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(9999);
! Used to communicate how missing data is coded in data
file. Here shown with a "." which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
  TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 1000 250;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifies
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will stop
!before reaching 500 iterations.
!These values can be increased ... see "Four-Profile Final
Model with Covariate Analysis
!Syntax" for an example.
LRTSTARTS = 2 1 50 10;
LRTBOOTSTRAP = 250;
!The above start values are for the defaults for the LRT
statistic being run to compare the
!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statement
!specifies the number of bootstrap draws to inform Mplus'
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances

!profiles to be equal.

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#1%
[slpnssNC worryNC solC slpDurC];
slpnssNC worryNC solC slpDurC;

%c#2%
[slpnssNC worryNC solC slpDurC];
slpnssNC worryNC solC slpDurC;

%c#3%
[slpnssNC worryNC solC slpDurC];
slpnssNC worryNC solC slpDurC;

OUTPUT:
TECH11 TECH14;
! TECH1 provides parameter specifications and starting
values for the analysis
! TECH8 provides optimization history for this analysis type
!TECH11 provides LRT results
!TECH14 provides bootstrapped LRT test
PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);
TYPE=PLOT3;
SAVEDATA:
FILE IS Step1_LPA3M4.dat;
! Tells Mplus where to save the output files from the
analysis
SAVE = CPROBABILITIES;
! The above command lines are to save the most likely
profile membership for each participant
! and the posterior probabilities for their membership in
each latent profile.

LPA 3 profile syntax: M4: Varying means, varying variances,
and equal covariances

SUMMARY OF ANALYSIS

Number of groups 1
Number of observations 27802

Number of dependent variables 4
Number of independent variables 0
Number of continuous latent variables 0
Number of categorical latent variables 1

Observed dependent variables

Continuous
SLPNSSNC WORRYNC SOLC SLPDURC

Categorical latent variables
C

Variables with special functions

ID variable ID

Estimator MLR

```

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Information matrix	OBSERVED	UNIVARIATE HIGHER-ORDER MOMENT DESCRIPTIVE STATISTICS				
Optimization Specifications for the Quasi-Newton Algorithm for		Variable/ Percentiles	Mean/ Sample Size	Skewness/ Variance	Minimum/ Kurtosis	% with Maximum
Continuous Outcomes		Min/Max	20%/60%	40%/80%	Median	
Maximum number of iterations	100	SLPNSSNC	0.933	0.086	0.000	27.10%
Convergence criterion	0.100D-05	0.000	1.000	1.000		
Optimization Specifications for the EM Algorithm		27802.000	0.471	-0.878	2.000	20.40%
Maximum number of iterations	500	1.000	2.000			
Convergence criteria		WORRYNC	0.000	-1.036	-2.825	1.71%
Loglikelihood change	0.100D-06	-0.928	0.020	0.495		
Relative loglikelihood change	0.100D-06	27802.000	1.000	0.165	0.969	28.55%
Derivative	0.100D-05	0.495	0.969			
Optimization Specifications for the M step of the EM Algorithm for		SOLC	0.000	0.967	-1.054	31.63%
Categorical Latent variables		1.054	-0.072	-0.072		
Number of M step iterations	1	27123.000	1.000	0.467	2.875	2.77%
M step convergence criterion	0.100D-05	-0.072	0.910			
Basis for M step termination	ITERATION	SLPDURC	0.000	0.568	-1.205	25.70%
Optimization Specifications for the M step of the EM Algorithm for		-1.205	-0.292	-0.292		
Censored, Binary or Ordered Categorical (Ordinal), Unordered		26763.000	1.000	-0.470	2.445	3.44%
Categorical (Nominal) and Count Outcomes		-0.292	0.620			
Number of M step iterations	1	RANDOM STARTS RESULTS RANKED FROM THE BEST TO THE WORST LOGLIKELIHOOD VALUES				
M step convergence criterion	0.100D-05	Unperturbed starting value run did not converge in the initial stage				
Basis for M step termination	ITERATION	optimizations.				
Maximum value for logit thresholds	15	657 perturbed starting value run(s) did not converge in the initial stage				
Minimum value for logit thresholds	-15	optimizations.				
Minimum expected cell size for chi-square	0.100D-01	Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:				
Maximum number of iterations for H1	2000	-122273.536	464179	106		
Convergence criterion for H1	0.100D-03	-122273.536	791396	243		
Optimization algorithm	EMA	-122273.536	208797	931		
Random Starts Specifications		-122273.536	985387	381		
Number of initial stage random starts	1000	-122273.536	507218	613		
Number of final stage optimizations	250	-122273.536	408713	450		
Number of initial stage iterations	500	-122273.536	961454	665		
Initial stage convergence criterion	0.100D+01	-122273.536	856536	144		
Random starts scale	0.500D+01	-122273.536	520865	763		
Random seed for generating random starts	0	-122273.536	640833	434		
Input data file(s)		-122273.536	210139	991		
mplus2.csv		-122273.536	996231	310		
Input data format	FREE	-122273.536	622173	992		
SUMMARY OF DATA		-122273.536	970689	266		
Number of missing data patterns	4	-122273.536	838615	677		
Number of y missing data patterns	4	-122273.536	783165	170		
Number of u missing data patterns	0	-122273.536	49293	707		
COVARIANCE COVERAGE OF DATA		-122273.536	86698	893		
Minimum covariance coverage value	0.100	-122273.536	436892	565		
PROPORTION OF DATA PRESENT FOR Y		-122273.536	724519	333		
Covariance Coverage		-122273.536	512836	289		
SLPNSSNC		-122273.536	152496	123		
WORRYNC		-122273.536	987560	985		
SOLC		-122273.536	669639	699		
SLPDURC		-122273.536	538872	949		
SLPNSSNC	1.000	-122273.536	46437	153		
WORRYNC	1.000	-122273.536	995913	787		
SOLC	0.976	-122273.536	662718	460		
SLPDURC	0.963	-122273.536	903369	134		
UNIVARIATE SAMPLE STATISTICS		-122273.536	153053	378		
		-122273.536	535303	923		

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-122273.536	972873	157	-122273.536	499347	881
-122273.536	307843	664	-122273.536	81951	907
-122273.536	609185	181	-122273.536	168762	200
-122273.536	85114	385	-122273.536	741484	441
-122273.536	85734	411	-122273.536	130541	896
-122273.536	876943	650	-122273.536	432513	803
-122273.536	610181	711	-122273.536	954914	911
-122273.536	395754	388	-122273.536	212934	568
-122273.536	634782	979	-122273.536	421731	886
-122273.536	416463	467	-122273.536	605161	409
-122273.536	59963	670	-122273.536	505244	582
-122273.536	751153	110	-122273.536	579138	706
-122273.536	534193	689	-122273.536	966603	919
-122273.536	865906	641	-122273.536	751054	117
-122273.536	726744	939	-122273.536	364676	27
-122273.536	920593	611	-122273.536	597614	284
-122273.536	263268	165	-122273.536	484501	163
-122273.536	443917	60	-122273.536	121425	511
-122273.536	529455	268	-122273.536	323588	826
-122273.536	645664	39	-122273.536	637345	19
-122273.536	749453	33	-122273.536	978781	497
-122273.536	344422	296	-122273.536	942358	644
-122273.536	871722	355	-122273.536	327475	518
-122273.536	278692	342	-122273.536	643311	888
-122273.536	939870	655	-122273.536	425929	508
-122273.536	813779	92	-122273.536	762461	425
-122273.536	840031	276	-122273.536	366706	29
-122273.536	514326	325	-122273.536	177175	851
-122273.536	383979	603	-122273.536	3607	873
-122273.536	206099	363	-122273.536	475419	987
-122273.536	347515	24	-122273.536	782821	272
-122273.536	485635	876	-122273.536	210870	383
-122273.536	215353	164	-122273.536	561664	392
-122273.536	791285	416	-122273.536	384199	882
-122273.536	81233	825	-122273.536	319575	499
-122273.536	964570	701	-122273.536	805768	879
-122273.536	301717	823	-122273.536	50887	389
-122273.536	549244	756	-122273.536	327140	678
-122273.536	777492	972	-122273.536	137377	397
-122273.536	376411	473	-122273.536	580181	691
-122273.536	863691	481	-122273.536	569833	85
-122273.536	436806	883	-122273.536	349263	263
-122273.536	22362	365	-122273.536	636396	168
-122273.536	268217	83	-122273.536	292884	103
-122273.536	930872	277	-122273.536	789985	67
-122273.536	211281	292	-122273.536	227563	63
-122273.536	772131	407	-122273.536	489927	520
-122273.536	765586	709	-122273.536	965994	396
-122273.536	134830	780	-122273.536	241197	747
-122273.536	241299	912	-122273.536	629320	222
-122273.536	313306	612	-122273.536	473942	574
-122273.536	178181	753	-122273.536	710154	831
-122273.536	917702	693	-122273.536	214681	824
-122273.536	752769	253	-122273.536	326091	759
-122273.536	436460	89	-122273.536	741888	138
-122273.536	243346	930	-122273.536	580539	786
-122273.536	670281	721	-122273.536	387701	275
-122273.536	507154	387	-122273.536	648555	113
-122273.536	148918	682	-122273.536	569131	26
-122273.536	118421	139	-122273.536	508482	446
-122273.536	999211	628	-122273.536	833196	715
-122273.536	462953	7	-122273.536	699810	571
-122273.536	700270	855	-122273.536	804104	566
-122273.536	857799	315	-122273.536	765392	382
-122273.536	784664	75	-122273.536	140442	500
-122273.536	679832	302	-122273.536	465160	862
-122273.536	891531	206	-122273.536	238317	812
-122273.536	471398	74	-122273.536	937588	293
-122273.536	167409	772	-122273.536	22874	588
-122273.536	787985	482	-122273.536	458181	189

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-122273.536	953843	651	-122273.536	318177	748
-122273.536	650371	14	-122273.536	213189	605
-122273.536	930323	267	-122273.536	42523	730
-122273.536	495490	990	-122273.536	848969	173
-122273.536	665121	417	-122273.536	991329	182
-122273.536	638977	643	-122273.536	123985	248
-122273.536	314034	513	-122273.536	738393	619
-122273.536	793035	187			
-122273.536	544077	809			
-122273.536	618000	190	2 perturbed starting value run(s) did not converge or were rejected in the third stage.		
-122273.536	939021	8			
-122273.536	529496	343	THE BEST LOGLIKELIHOOD VALUE HAS BEEN REPLICATED. RERUN WITH AT LEAST TWICE THE RANDOM STARTS TO CHECK THAT THE BEST LOGLIKELIHOOD IS STILL OBTAINED AND REPLICATED.		
-122273.536	836515	341			
-122273.536	829540	324			
-122273.536	77571	980			
-122273.536	804561	59			
-122273.536	178475	231			
-122273.536	118438	601	THE MODEL ESTIMATION TERMINATED NORMALLY		
-122273.536	193042	316			
-122273.536	682718	733	MODEL FIT INFORMATION		
-122273.536	715255	523			
-122273.536	604861	800	Number of Free Parameters		32
-122273.536	588923	606			
-122273.536	673496	218	Loglikelihood		
-122273.536	231400	738			
-122273.536	197223	332			
-122273.536	264521	929	H0 Value	-122273.536	
-122273.536	798839	312	H0 Scaling Correction Factor	1.0161	
-122273.536	440395	917			
-122273.536	66276	217			
-122273.536	459221	652	Information Criteria		
-122273.536	312754	562			
-122273.536	995249	525	Akaike (AIC)	244611.072	
-122273.536	855760	593	Bayesian (BIC)	244874.523	
-122273.536	29591	633	Sample-Size Adjusted BIC	244772.828	
-122273.536	92564	583		(n* = (n + 2) / 24)	
-122273.536	194143	510	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THE ESTIMATED MODEL		
-122273.536	791678	974			
-122273.536	92091	649			
-122273.536	303834	798			
-122273.536	67009	564	Latent		
-122273.536	268896	124	Classes		
-122273.536	291149	536			
-122273.536	872743	374	1	9597.87522	0.34522
-122273.536	843555	952	2	10278.69205	0.36971
-122273.536	875667	367	3	7925.43273	0.28507
-122273.536	783110	72			
-122273.536	359578	458	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON ESTIMATED POSTERIOR PROBABILITIES		
-122273.536	391179	78			
-122273.536	888905	444			
-122273.536	695155	150			
-122273.536	794236	127	Latent		
-122273.536	679615	942	Classes		
-122273.536	17896	592			
-122273.536	573367	986	1	9597.87522	0.34522
-122273.536	694303	282	2	10278.69205	0.36971
-122273.536	281462	285	3	7925.43273	0.28507
-122273.536	474357	789			
-122273.536	551639	55	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THEIR MOST LIKELY LATENT CLASS MEMBERSHIP		
-122273.536	645052	910			
-122273.536	352277	42			
-122273.536	685268	596			
-122273.536	677720	681			
-122273.536	153394	429	Class Counts and Proportions		
-122273.536	761302	744			
-122273.536	848331	137			
-122273.536	471040	403	Latent		
-122273.536	746978	410	Classes		
-122273.536	267983	228			
-122273.536	486622	522	1	11537	0.41497
			2	8897	0.32001

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

3	7368	0.26502								SLPNSSNC WITH					
										WORRYNC	-0.010	0.002	-4.859	0.000	
										SOLC	0.077	0.003	23.898	0.000	
										SLPDURC	0.008	0.004	2.133	0.033	
CLASSIFICATION QUALITY															
										WORRYNC WITH					
	Entropy									SOLC	-0.048	0.002	-22.837	0.000	
										SLPDURC	-0.021	0.002	-11.280	0.000	
Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)															
										SOLC WITH					
										SLPDURC	0.045	0.003	13.331	0.000	
										Means					
										SLPNSSNC	0.899	0.009	99.109	0.000	
										WORRYNC	0.178	0.030	5.939	0.000	
										SOLC	-0.160	0.014	-11.305	0.000	
										SLPDURC	0.059	0.021	2.866	0.004	
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)															
										Variances					
										SLPNSSNC	0.469	0.005	94.190	0.000	
										WORRYNC	0.273	0.017	15.609	0.000	
										SOLC	0.373	0.008	44.113	0.000	
										SLPDURC	0.684	0.017	39.807	0.000	
										Latent Class 3					
										SLPNSSNC WITH					
										WORRYNC	-0.010	0.002	-4.859	0.000	
										SOLC	0.077	0.003	23.898	0.000	
										SLPDURC	0.008	0.004	2.133	0.033	
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)															
										WORRYNC WITH					
										SOLC	-0.048	0.002	-22.837	0.000	
										SLPDURC	-0.021	0.002	-11.280	0.000	
										SOLC WITH					
										SLPDURC	0.045	0.003	13.331	0.000	
										Means					
										SLPNSSNC	1.042	0.010	99.351	0.000	
										WORRYNC	-1.206	0.023	-53.488	0.000	
										SOLC	1.012	0.031	33.071	0.000	
										SLPDURC	0.812	0.019	42.391	0.000	
										Variances					
										SLPNSSNC	0.479	0.006	86.226	0.000	
										WORRYNC	0.803	0.017	48.499	0.000	
										SOLC	1.125	0.015	73.837	0.000	
										SLPDURC	1.016	0.015	68.881	0.000	
MODEL RESULTS															
										Categorical Latent Variables					
										Means					
										C#1	0.191	0.057	3.332	0.001	
										C#2	0.260	0.034	7.668	0.000	
										QUALITY OF NUMERICAL RESULTS					
										Condition Number for the Information Matrix					
										0.933E-04					
										(ratio of smallest to largest eigenvalue)					
										TECHNICAL 11 OUTPUT					
										Random Starts Specifications for the k-1 Class Analysis Model					
										Number of initial stage random starts				1000	
										Number of final stage optimizations				250	
										Latent Class 2					

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<p>VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES</p> <table border="0"> <tr> <td>H0 Loglikelihood Value</td> <td>-124198.073</td> </tr> <tr> <td>2 Times the Loglikelihood Difference</td> <td>3849.074</td> </tr> <tr> <td>Difference in the Number of Parameters</td> <td>9</td> </tr> <tr> <td>Mean</td> <td>27.434</td> </tr> <tr> <td>Standard Deviation</td> <td>33.435</td> </tr> <tr> <td>P-Value</td> <td>0.0000</td> </tr> </table>	H0 Loglikelihood Value	-124198.073	2 Times the Loglikelihood Difference	3849.074	Difference in the Number of Parameters	9	Mean	27.434	Standard Deviation	33.435	P-Value	0.0000	<p>Histograms (sample values) Scatterplots (sample values) Sample means Estimated means, medians, modes, and percentiles Sample and estimated means Observed individual values Estimated means and observed individual values Estimated overall and class-specific distributions</p>																				
H0 Loglikelihood Value	-124198.073																																
2 Times the Loglikelihood Difference	3849.074																																
Difference in the Number of Parameters	9																																
Mean	27.434																																
Standard Deviation	33.435																																
P-Value	0.0000																																
<p>LO-MENDELL-RUBIN ADJUSTED LRT TEST</p> <table border="0"> <tr> <td>Value</td> <td>3807.728</td> </tr> <tr> <td>P-Value</td> <td>0.0000</td> </tr> </table>	Value	3807.728	P-Value	0.0000	<p>SAVE DATA INFORMATION</p> <p>Save file Step1_LPA3M4.dat</p>																												
Value	3807.728																																
P-Value	0.0000																																
<p>TECHNICAL 14 OUTPUT</p> <p>Random Starts Specifications for the k-1 Class Analysis Model</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>1000</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>250</td> </tr> </table> <p>Random Starts Specification for the k-1 Class Model for Generated Data</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>2</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>1</td> </tr> </table> <p>Random Starts Specification for the k Class Model for Generated Data</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>50</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>10</td> </tr> <tr> <td>Number of bootstrap draws requested</td> <td>250</td> </tr> </table>	Number of initial stage random starts	1000	Number of final stage optimizations	250	Number of initial stage random starts	2	Number of final stage optimizations	1	Number of initial stage random starts	50	Number of final stage optimizations	10	Number of bootstrap draws requested	250	<p>Order and format of variables</p> <table border="0"> <tr> <td>SLPNSSNC</td> <td>F10.3</td> </tr> <tr> <td>WORRYNC</td> <td>F10.3</td> </tr> <tr> <td>SOLC</td> <td>F10.3</td> </tr> <tr> <td>SLPDURC</td> <td>F10.3</td> </tr> <tr> <td>CPROB1</td> <td>F10.3</td> </tr> <tr> <td>CPROB2</td> <td>F10.3</td> </tr> <tr> <td>CPROB3</td> <td>F10.3</td> </tr> <tr> <td>C</td> <td>F10.3</td> </tr> <tr> <td>ID</td> <td>I6</td> </tr> </table> <p>Save file format 8F10.3 I6</p> <p>Save file record length 10000</p> <p>Save missing symbol *</p>	SLPNSSNC	F10.3	WORRYNC	F10.3	SOLC	F10.3	SLPDURC	F10.3	CPROB1	F10.3	CPROB2	F10.3	CPROB3	F10.3	C	F10.3	ID	I6
Number of initial stage random starts	1000																																
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<p>PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES</p> <table border="0"> <tr> <td>H0 Loglikelihood Value</td> <td>-124198.073</td> </tr> <tr> <td>2 Times the Loglikelihood Difference</td> <td>3849.074</td> </tr> <tr> <td>Difference in the Number of Parameters</td> <td>9</td> </tr> <tr> <td>Approximate P-Value</td> <td>0.0000</td> </tr> <tr> <td>Successful Bootstrap Draws</td> <td>250</td> </tr> </table> <p>WARNING: OF THE 250 BOOTSTRAP DRAWS, 199 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 3-CLASS MODEL. THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA. INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.</p>	H0 Loglikelihood Value	-124198.073	2 Times the Loglikelihood Difference	3849.074	Difference in the Number of Parameters	9	Approximate P-Value	0.0000	Successful Bootstrap Draws	250	<p>DIAGRAM INFORMATION</p> <p>Mplus diagrams are currently not available for Mixture analysis. No diagram output was produced.</p> <p>Beginning Time: 15:04:04 Ending Time: 16:37:00 Elapsed Time: 01:32:56</p>																						
H0 Loglikelihood Value	-124198.073																																
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<p>PLOT INFORMATION</p> <p>The following plots are available:</p>	<p>MUTHEN & MUTHEN 3463 Stoner Ave. Los Angeles, CA 90066</p> <p>Tel: (310) 391-9971 Fax: (310) 391-8971 Web: www.StatModel.com Support: Support@StatModel.com</p> <p>Copyright (c) 1998-2021 Muthen & Muthen</p>																																

Parametrization Model 5 - Equal variances and varying covariances

<p>Mplus VERSION 8.7 MUTHEN & MUTHEN 07/12/2023 10:37 AM</p> <p>INPUT INSTRUCTIONS</p> <p>TITLE: LPA 3 profile syntax: M5: Varying means, equal variances, and varying covariances</p> <p>DATA: FILE IS mplus2.csv;</p>	<p>! Specifies file location for data file. Make sure data is in format appropriate for Mplus ! per Mplus manual. This data file is in individual format (one row of data per participant) VARIABLE: NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol slpDur gender genderN yrgrp yrgrpN SCI_02C slpnssNC worryNC solC slpDurC;</p> <p>! All variables included in data file should be named here. USEVARIABLES ARE slpnssNC worryNC solC slpDurC;</p>
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Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

<p>! Only variables intended for use in the analysis should be listed here</p> <p>IDVARIABLE IS ID; CLASSES = c (3); ! This is where you instruct Mplus on how many classes/profiles are being estimated. Initi</p> <p>! model contains only one class/profile, thus it would be CLASSES = c (1). Above specifies</p> <p>!profiles, and for each further iterative models the number in parentheses increases by on</p> <p>!three profiles/classes would be c (3), and so on.</p> <p>MISSING ARE ALL(9999); ! Used to communicate how missing data is coded in data file. Here shown with a “.” which</p> <p>! all that is included in each cell with missing data in the data file</p> <p>ANALYSIS: TYPE = MIXTURE; ! LPA is a version of mixture modeling, and this instructs Mplus to analyze in this way</p> <p>ESTIMATOR = MLR; !FIML robust to non-normal data STARTS = 1000 250; STITERATIONS = 500; ! Default number of starts for each step of the ML estimation. First STARTS value specifie</p> <p>!number of unique start values to start with, the 250 represents the 250 best unique start</p> <p>!carrying forward to completion. The STITERATIONS specifies the number of ML iteration</p> <p>!steps for those 250 selected start values to go through to be able to converge. This is a</p> <p>!maximum number of iteration; if a model converges in less than 500 iterations it will sto</p> <p>!before reaching 500 iterations.</p> <p>!These values can be increased ... see “Four-Profile Final Model with Covariate Analysis</p> <p>!Syntax” for an example.</p> <p>LRTSTARTS = 2 1 50 10; LRTBOOTSTRAP = 250; !The above start values are for the defaults for the LRT statistic being run to compare th</p> <p>!model fit with the model fit of a model with one less class (k-1). The BOOTSTRAP statemen</p> <p>!specifies the number of bootstrap draws to inform Mplus’ bootstrapped LRT results.</p> <p>MODEL: !For a default Mplus model the LPA model does not need to be specified. However, it can be</p> <p>!The model can also be modified from the Mplus default of estimating the indicator means</p> <p>!(uniquely across profiles) and variances (constrained across profiles), as well as the la</p> <p>!profile mean.</p> <p>%OVERALL% [slpnssNC worryNC solC slpDurC]; !estimates the indicators means for each profile. Without</p> <p>!the means are freely estimated in each profile, not constrained.</p> <p>slpnssNC worryNC solC slpDurC; !Label Var1-Var5 constrains the estimates of the variances</p> <p>!profiles to be equal.</p> <p>%c#1%</p> <p>[slpnssNC worryNC solC slpDurC];</p> <p>!covariances slpnssNC WITH worryNC solC slpDurC; worryNC WITH solC slpDurC;</p>	<p>solC WITH slpDurC;</p> <p>%c#2%</p> <p>[slpnssNC worryNC solC slpDurC];</p> <p>!covariances slpnssNC WITH worryNC solC slpDurC; worryNC WITH solC slpDurC; solC WITH slpDurC;</p> <p>%c#3%</p> <p>[slpnssNC worryNC solC slpDurC];</p> <p>!covariances slpnssNC WITH worryNC solC slpDurC; worryNC WITH solC slpDurC; solC WITH slpDurC;</p> <p>OUTPUT: TECH11 TECH14; ! TECH1 provides parameter specifications and starting values for the analysis</p> <p>! TECH8 provides optimization history for this analysis type</p> <p>!TECH11 provides LRT results</p> <p>!TECH14 provides bootstrapped LRT test</p> <p>PLOT: SERIES=slpnssNC worryNC solC slpDurC(*); TYPE=PLOT3; SAVEDATA: FILE IS Step1_LPA3M5.dat; ! Tells Mplus where to save the output files from the analysis</p> <p>SAVE = CPROBABILITIES; ! The above command lines are to save the most likely profile membership for each particip</p> <p>! and the posterior probabilities for their membership in each latent profile.</p> <p>*** WARNING in MODEL command</p> <p>All variables are uncorrelated with all other variables within class.</p> <p>Check that this is what is intended.</p> <p>LPA 3 profile syntax: M5: Varying means, equal variances, and varying covariances</p> <p>SUMMARY OF ANALYSIS</p> <table border="0"> <tr> <td>Number of groups</td> <td>1</td> </tr> <tr> <td>Number of observations</td> <td>27802</td> </tr> <tr> <td>Number of dependent variables</td> <td>4</td> </tr> <tr> <td>Number of independent variables</td> <td>0</td> </tr> <tr> <td>Number of continuous latent variables</td> <td>0</td> </tr> <tr> <td>Number of categorical latent variables</td> <td>1</td> </tr> </table> <p>Observed dependent variables</p> <table border="0"> <tr> <td>Continuous</td> <td>SLPNSSNC</td> <td>WORRYNC</td> <td>SOLC</td> <td>SLPDURC</td> </tr> </table> <p>Categorical latent variables</p> <table border="0"> <tr> <td>C</td> </tr> </table> <p>Variables with special functions</p> <table border="0"> <tr> <td>ID variable</td> <td>ID</td> </tr> <tr> <td>Estimator</td> <td>MLR</td> </tr> </table>	Number of groups	1	Number of observations	27802	Number of dependent variables	4	Number of independent variables	0	Number of continuous latent variables	0	Number of categorical latent variables	1	Continuous	SLPNSSNC	WORRYNC	SOLC	SLPDURC	C	ID variable	ID	Estimator	MLR
Number of groups	1																						
Number of observations	27802																						
Number of dependent variables	4																						
Number of independent variables	0																						
Number of continuous latent variables	0																						
Number of categorical latent variables	1																						
Continuous	SLPNSSNC	WORRYNC	SOLC	SLPDURC																			
C																							
ID variable	ID																						
Estimator	MLR																						

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Information matrix	OBSERVED	Variable/ Percentiles	Mean/ Sample Size	Skewness/ Variance	Minimum/ Kurtosis	% with Maximum
Optimization Specifications for the Quasi-Newton Algorithm for		Min/Max	20%/60%	40%/80%	Median	
Continuous Outcomes						
Maximum number of iterations	100					
Convergence criterion	0.100D-05					
Optimization Specifications for the EM Algorithm		SLPNSSNC	0.933	0.086	0.000	27.10%
Maximum number of iterations	500	0.000 1.000	1.000			
Convergence criteria		27802.000	0.471	-0.878	2.000	20.40%
Loglikelihood change	0.100D-06	1.000 2.000				
Relative loglikelihood change	0.100D-06	WORRYNC	0.000	-1.036	-2.825	1.71%
Derivative	0.100D-05	-0.928 0.020	0.495			
Optimization Specifications for the M step of the EM Algorithm for		27802.000	1.000	0.165	0.969	28.55%
Categorical Latent variables		0.495 0.969				
Number of M step iterations	1	SOLC	0.000	0.967	-1.054	31.63%
M step convergence criterion	0.100D-05	1.054 -0.072	-0.072			
Basis for M step termination	ITERATION	27123.000	1.000	0.467	2.875	2.77%
Optimization Specifications for the M step of the EM Algorithm for		-0.072 0.910				
Censored, Binary or Ordered Categorical (Ordinal), Unordered		SLPDURC	0.000	0.568	-1.205	25.70%
Categorical (Nominal) and Count Outcomes		-1.205 -0.292	-0.292			
Number of M step iterations	1	26763.000	1.000	-0.470	2.445	3.44%
M step convergence criterion	0.100D-05	-0.292 0.620				
Basis for M step termination	ITERATION					
Maximum value for logit thresholds	15					
Minimum value for logit thresholds	-15					
Minimum expected cell size for chi-square	0.100D-01					
Maximum number of iterations for H1	2000					
Convergence criterion for H1	0.100D-03					
Optimization algorithm	EMA					
Random Starts Specifications						
Number of initial stage random starts	1000					
Number of final stage optimizations	250					
Number of initial stage iterations	500					
Initial stage convergence criterion	0.100D+01					
Random starts scale	0.500D+01					
Random seed for generating random starts	0					
Input data file(s)						
mplus2.csv						
Input data format	FREE					
SUMMARY OF DATA						
Number of missing data patterns	4					
Number of y missing data patterns	4					
Number of u missing data patterns	0					
COVARIANCE COVERAGE OF DATA						
Minimum covariance coverage value	0.100					
PROPORTION OF DATA PRESENT FOR Y						
Covariance Coverage						
SLPNSSNC						
WORRYNC						
SOLC						
SLPDURC						
UNIVARIATE SAMPLE STATISTICS						
UNIVARIATE HIGHER-ORDER MOMENT						
DESCRIPTIVE STATISTICS						

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-125994.415 200041	810	-125994.415 679832	302
-125994.415 130541	896	-125994.415 809240	543
-125994.415 616917	261	-125994.415 172913	557
-125994.415 126371	526	-125994.415 830392	35
-125994.415 849670	347	-125994.415 97300	640
-125994.415 237332	661	-125994.415 597614	284
-125994.415 114433	708	-125994.415 723035	538
-125994.415 903633	553	-125994.415 682718	733
-125994.415 202790	198	-125994.415 81951	907
-125994.415 833196	715	-125994.415 879338	309
-125994.415 137377	397	-125994.415 966499	963
-125994.415 399508	415	-125994.415 669634	335
-125994.415 862607	468	-125994.415 863691	481
-125994.415 544048	87	-125994.415 717754	180
-125994.415 414284	158	-125994.415 654136	167
-125994.415 490123	995	-125994.415 193569	440
-125994.415 937885	426	-125994.415 298201	903
-125994.415 105435	265	-125994.415 347515	24
-125994.415 46502	714	-125994.415 545108	667
-125994.415 992389	77	-125994.415 980970	894
-125994.415 308582	752	-125994.415 695155	150
-125994.415 373815	618	-125994.415 766903	505
-125994.415 856536	144	-125994.415 922596	456
-125994.415 679448	937	-125994.415 51375	148
-125994.415 505879	424	-125994.415 158612	581
-125994.415 224950	455	-125994.415 364676	27
-125994.415 260953	589	-125994.415 566739	575
-125994.415 339073	841	-125994.415 442072	625
-125994.415 575700	100	-125994.415 322790	636
-125994.415 549244	756	-125994.415 852283	616
-125994.415 391949	295	-125994.415 677062	680
-125994.415 701525	239	-125994.415 696830	668
-125994.415 576596	99	-125994.415 792389	746
-125994.415 794236	127	-125994.415 535303	923
-125994.415 484687	306	-125994.415 17896	592
-125994.415 972430	491	-125994.415 857122	889
-125994.415 971693	470	-125994.415 735928	916
-125994.415 453915	975	-125994.415 961454	665
-125994.415 710154	831	-125994.415 881886	608
-125994.415 850112	922	-125994.415 370481	742
-125994.415 443442	380	-125994.415 195353	225
-125994.415 444228	860	-125994.415 405371	569
-125994.415 179022	687	-125994.415 415931	10
-125994.415 50887	389	-125994.415 148918	682
-125994.415 70118	104	-125994.415 119513	821
-125994.415 204959	695	-125994.415 303634	169
-125994.415 570908	98	-125994.415 465160	862
-125994.415 907810	795	-125994.415 414828	322
-125994.415 705224	953	-125994.415 333082	578
-125994.415 963967	941	-125994.415 928624	981
-125994.415 923437	398	-125994.415 548245	818
-125994.415 138695	783	-125994.415 599729	658
-125994.415 124999	96	-125994.415 207896	25
-125994.415 634782	979	-125994.415 485483	498
-125994.415 563584	657	-125994.415 640833	434
-125994.415 742609	531	-125994.415 762461	425
-125994.415 836515	341	-125994.415 136842	58
-125994.415 848356	602	-125994.415 311214	64
-125994.415 298553	773	-125994.415 72662	729
-125994.415 745972	521	-125994.415 297518	166
-125994.415 965639	463	-125994.415 61587	400
-125994.415 264081	186	-125994.415 569833	85
-125994.415 213532	503	-125994.415 662718	460
-125994.415 284384	600	-125994.415 484501	163
-125994.415 673496	218	-125994.415 432513	803
-125994.415 440368	797	-125994.415 843555	952
-125994.415 628143	854	-125994.415 unperturbed	0
-125994.415 847088	750	-125994.415 626208	698
-125994.415 954914	911	-125994.415 599136	811
-125994.415 271809	846	-125994.415 721392	768

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-125994.415	153394	429	-125994.415	621542	375
-125994.415	178031	720	-125994.415	326091	759
-125994.415	728038	177	-125994.415	691041	590
-125994.415	484406	421			
-125994.415	224151	973			
-125994.415	589483	950	THE BEST LOGLIKELIHOOD VALUE HAS BEEN		
-125994.415	782821	272	REPLICATED. RERUN WITH AT LEAST TWICE THE		
-125994.415	352277	42	RANDOM STARTS TO CHECK THAT THE BEST		
-125994.415	238906	430	LOGLIKELIHOOD IS STILL OBTAINED AND		
-125994.415	848331	137	REPLICATED.		
-125994.415	855760	593			
-125994.415	891347	504	THE MODEL ESTIMATION TERMINATED NORMALLY		
-125994.415	468036	131			
-125994.415	437181	135	MODEL FIT INFORMATION		
-125994.415	552272	654			
-125994.415	520177	262	Number of Free Parameters	36	
-125994.415	268217	83			
-125994.415	580405	286	Loglikelihood		
-125994.415	87586	871			
-125994.415	638977	643	H0 Value	-125994.415	
-125994.415	55115	408	H0 Scaling Correction Factor	1.3296	
-125994.415	248742	556	for MLR		
-125994.415	838615	677			
-125994.415	751153	110	Information Criteria		
-125994.415	384199	882			
-125994.415	178181	753	Akaike (AIC)	252060.830	
-125994.415	22362	365	Bayesian (BIC)	252357.213	
-125994.415	792993	859	Sample-Size Adjusted BIC	252242.806	
-125994.415	778953	635	(n* = (n + 2) / 24)		
-125994.415	286735	175			
-125994.415	617243	237	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-125994.415	605161	409	LATENT CLASSES		
-125994.415	846194	93	BASED ON THE ESTIMATED MODEL		
-125994.415	357866	968			
-125994.415	726035	191	Latent		
-125994.415	483369	270	Classes		
-125994.415	399380	436			
-125994.415	434915	552	1	5737.65936	0.20638
-125994.415	782200	84	2	18891.89637	0.67952
-125994.415	538872	949	3	3172.44427	0.11411
-125994.415	871851	257			
-125994.415	814975	129	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-125994.415	157351	579	LATENT CLASSES		
-125994.415	81233	825	BASED ON ESTIMATED POSTERIOR PROBABILITIES		
-125994.415	526324	178			
-125994.415	366706	29	Latent		
-125994.415	211281	292	Classes		
-125994.415	848969	173			
-125994.415	937588	293	1	5737.65936	0.20638
-125994.415	788796	145	2	18891.89637	0.67952
-125994.415	574942	558	3	3172.44427	0.11411
-125994.415	35191	703			
-125994.415	195873	6	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-125994.415	118438	601	LATENT CLASSES		
-125994.415	576726	280	BASED ON THEIR MOST LIKELY LATENT CLASS		
-125994.415	345974	622	MEMBERSHIP		
-125994.415	467339	66			
-125994.415	566687	597	Class Counts and Proportions		
-125994.415	407108	366			
-125994.415	791678	974	Latent		
-125994.415	608460	244	Classes		
-125994.415	860029	760			
-125994.415	14262	781	1	6141	0.22088
-125994.415	50983	834	2	18658	0.67110
-125994.415	592219	119	3	3003	0.10801
-125994.415	891531	206			
-125994.415	265218	924	CLASSIFICATION QUALITY		
-125994.415	370957	554			
-125994.415	264935	281	Entropy	0.854	
-125994.415	127362	757			

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)					SLPDURC	-0.187	0.010	-18.506	0.000
					SOLC WITH				
					SLPDURC	0.290	0.011	27.050	0.000
					Means				
1					SLPNSSNC	0.904	0.005	171.611	0.000
2					WORRYNC	0.570	0.006	101.171	0.000
3					SOLC	-0.364	0.009	-40.558	0.000
					SLPDURC	-0.300	0.007	-40.716	0.000
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					Variates				
					SLPNSSNC	0.468	0.003	152.809	0.000
					WORRYNC	0.210	0.009	23.619	0.000
					SOLC	0.626	0.013	47.707	0.000
					SLPDURC	0.798	0.010	76.790	0.000
1					Latent Class 3				
2					SLPNSSNC WITH				
3					WORRYNC	0.009	0.006	1.533	0.125
					SOLC	0.046	0.011	4.198	0.000
					SLPDURC	0.013	0.012	1.153	0.249
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					WORRYNC WITH				
					SOLC	-0.006	0.006	-0.926	0.354
					SLPDURC	-0.070	0.009	-7.748	0.000
1					SOLC WITH				
2					SLPDURC	0.050	0.015	3.310	0.001
3					Means				
					SLPNSSNC	1.112	0.015	76.093	0.000
					WORRYNC	-1.994	0.027	-74.945	0.000
					SOLC	1.572	0.036	43.121	0.000
					SLPDURC	1.108	0.027	40.596	0.000
MODEL RESULTS					Variates				
					SLPNSSNC	0.468	0.003	152.809	0.000
					WORRYNC	0.210	0.009	23.619	0.000
					SOLC	0.626	0.013	47.707	0.000
					SLPDURC	0.798	0.010	76.790	0.000
					Categorical Latent Variables				
					Means				
Latent Class 1					C#1	0.593	0.046	12.891	0.000
					C#2	1.784	0.040	44.896	0.000
					SLPNSSNC WITH				
					WORRYNC	0.015	0.006	2.633	0.008
					SOLC	0.106	0.008	14.059	0.000
					SLPDURC	0.019	0.009	2.191	0.028
					WORRYNC WITH				
					SOLC	0.088	0.017	5.295	0.000
					SLPDURC	-0.048	0.010	-4.925	0.000
					SOLC WITH				
					SLPDURC	0.051	0.013	3.916	0.000
					Means				
					SLPNSSNC	0.931	0.012	78.488	0.000
					WORRYNC	-0.775	0.027	-28.725	0.000
					SOLC	0.353	0.018	19.838	0.000
					SLPDURC	0.391	0.023	17.144	0.000
					Variates				
					SLPNSSNC	0.468	0.003	152.809	0.000
					WORRYNC	0.210	0.009	23.619	0.000
					SOLC	0.626	0.013	47.707	0.000
					SLPDURC	0.798	0.010	76.790	0.000
Latent Class 2					QUALITY OF NUMERICAL RESULTS				
					Condition Number for the Information Matrix				
					0.903E-04				
					(ratio of smallest to largest eigenvalue)				
					TECHNICAL 11 OUTPUT				
					Random Starts Specifications for the k-1 Class Analysis				
					Model				
					Number of initial stage random starts			1000	
					Number of final stage optimizations			250	
					VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO				
					TEST FOR 2 (H0) VERSUS 3 CLASSES				
					H0 Loglikelihood Value			-127763.908	
					2 Times the Loglikelihood Difference			3538.985	
					Difference in the Number of Parameters			11	
					Mean			61.969	
					Standard Deviation			60.777	
1									
2									
3									

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

P-Value	0.0000	Sample and estimated means
LO-MENDELL-RUBIN ADJUSTED LRT TEST		Observed individual values
		Estimated means and observed individual values
		Estimated overall and class-specific distributions
Value	3507.822	
P-Value	0.0000	SAVEDATA INFORMATION
TECHNICAL 14 OUTPUT		Save file
		Step1_LPA3M5.dat
Random Starts Specifications for the k-1 Class Analysis Model		Order and format of variables
Number of initial stage random starts	1000	SLPNSSNC F10.3
Number of final stage optimizations	250	WORRYNC F10.3
Random Starts Specification for the k-1 Class Model for Generated Data		SOLC F10.3
Number of initial stage random starts	2	SLPDURC F10.3
Number of final stage optimizations	1	CPROB1 F10.3
Random Starts Specification for the k Class Model for Generated Data		CPROB2 F10.3
Number of initial stage random starts	50	CPROB3 F10.3
Number of final stage optimizations	10	C F10.3
Number of bootstrap draws requested	250	ID I6
PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES		Save file format
H0 Loglikelihood Value	-127763.908	8F10.3 I6
2 Times the Loglikelihood Difference	3538.985	Save file record length 10000
Difference in the Number of Parameters	11	Save missing symbol *
Approximate P-Value	0.0000	DIAGRAM INFORMATION
Successful Bootstrap Draws	250	Mplus diagrams are currently not available for Mixture analysis.
WARNING: OF THE 250 BOOTSTRAP DRAWS, 230 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 3-CLASS MODEL.		No diagram output was produced.
THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA.		Beginning Time: 10:37:29
INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.		Ending Time: 12:21:31
PLOT INFORMATION		Elapsed Time: 01:44:02
The following plots are available:		MUTHEN & MUTHEN
Histograms (sample values)		3463 Stoner Ave.
Scatterplots (sample values)		Los Angeles, CA 90066
Sample means		Tel: (310) 391-9971
Estimated means, medians, modes, and percentiles		Fax: (310) 391-8971
		Web: www.StatModel.com
		Support: Support@StatModel.com
		Copyright (c) 1998-2021 Muthen & Muthen

Parametrization Model 6 - Varying variances and varying covariances

Mplus VERSION 8.7
MUTHEN & MUTHEN
07/11/2023 5:32 PM

INPUT INSTRUCTIONS

TITLE: LPA 3 profile syntax: M6: Varying means, varying variances and varying covariances

DATA:

FILE IS mplus2.csv;

! Specifies file location for data file. Make sure data is in format appropriate for Mplus

! per Mplus manual. This data file is in individual format (one row of data per participant

VARIABLE:

NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol
slpDur gender genderN yrgrp yrgrpN SCI_02C
slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be listed here

IDVARIABLE IS ID;

CLASSES = c (3);

! This is where you instruct Mplus on how many classes/profiles are being estimated. Init

! model contains only one class/profile, thus it would be CLASSES = c (1). Above specifies

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```

!profiles, and for each further iterative models the number
in parentheses increases by on
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(9999);
! Used to communicate how missing data is coded in data
file. Here shown with a “.” which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 1000 250;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifie
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will sto
!before reaching 500 iterations.
!These values can be increased ... see “Four-Profile Final
Model with Covariate Analysis
!Syntax” for an example.
LRTSTARTS = 2 1 50 10;
LRTBOOTSTRAP = 250;
!The above start values are for the defaults for the LRT
statistic being run to compare th
!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statemen
!specifies the number of bootstrap draws to inform Mplus’
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances
!profiles to be equal.

%c#1%

[slpnssNC worryNC solC slpDurC];

slpnssNC worryNC solC slpDurC;
!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#2%

[slpnssNC worryNC solC slpDurC];

slpnssNC worryNC solC slpDurC;

```

```

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#3%

[slpnssNC worryNC solC slpDurC];

slpnssNC worryNC solC slpDurC;
!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

OUTPUT:
TECH11 TECH14;
! TECH1 provides parameter specifications and starting
values for the analysis
! TECH8 provides optimization history for this analysis type
!TECH11 provides LRT results
!TECH14 provides bootstrapped LRT test
PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);
TYPE=PLOT3;
SAVEDATA:
FILE IS Step1_LPA3M5.dat;
! Tells Mplus where to save the output files from the
analysis
SAVE = CPROBABILITIES;
! The above command lines are to save the most likely
profile membership for each particip
! and the posterior probabilities for their membership in
each latent profile.

*** WARNING in MODEL command
All variables are uncorrelated with all other variables within
class.
Check that this is what is intended.

LPA 3 profile syntax: M6: Varying means, varying variances,
and varying covariances

SUMMARY OF ANALYSIS

Number of groups 1
Number of observations 27802

Number of dependent variables 4
Number of independent variables 0
Number of continuous latent variables 0
Number of categorical latent variables 1

Observed dependent variables

Continuous
SLPNSSNC WORRYNC SOLC SLPDURC

Categorical latent variables
C

Variables with special functions

ID variable ID

Estimator MLR
Information matrix OBSERVED
Optimization Specifications for the Quasi-Newton Algorithm
for
Continuous Outcomes
Maximum number of iterations 100

```

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Convergence criterion	0.100D-05						
Optimization Specifications for the EM Algorithm		SLPNSSNC	0.933	0.086	0.000	27.10%	
Maximum number of iterations	500	0.000	1.000	1.000			
Convergence criteria		27802.000	0.471	-0.878	2.000	20.40%	
Loglikelihood change	0.100D-06	1.000	2.000				
Relative loglikelihood change	0.100D-06	WORRYNC	0.000	-1.036	-2.825	1.71%	
Derivative	0.100D-05	-0.928	0.020	0.495			
Optimization Specifications for the M step of the EM Algorithm for		27802.000	1.000	0.165	0.969	28.55%	
Categorical Latent variables		0.495	0.969				
Number of M step iterations	1	SOLC	0.000	0.967	-1.054	31.63%	-
M step convergence criterion	0.100D-05	1.054	-0.072	-0.072			
Basis for M step termination	ITERATION	27123.000	1.000	0.467	2.875	2.77%	
Optimization Specifications for the M step of the EM Algorithm for		-0.072	0.910				
Censored, Binary or Ordered Categorical (Ordinal), Unordered		SLPDURC	0.000	0.568	-1.205	25.70%	
Categorical (Nominal) and Count Outcomes		-1.205	-0.292	-0.292			
Number of M step iterations	1	26763.000	1.000	-0.470	2.445	3.44%	
M step convergence criterion	0.100D-05	-0.292	0.620				
Basis for M step termination	ITERATION						
Maximum value for logit thresholds	15						
Minimum value for logit thresholds	-15						
Minimum expected cell size for chi-square	0.100D-01						
Maximum number of iterations for H1	2000						
Convergence criterion for H1	0.100D-03						
Optimization algorithm	EMA						
Random Starts Specifications							
Number of initial stage random starts	1000						
Number of final stage optimizations	250						
Number of initial stage iterations	500						
Initial stage convergence criterion	0.100D+01						
Random starts scale	0.500D+01						
Random seed for generating random starts	0						
Input data file(s)							
mplus2.csv							
Input data format	FREE						
SUMMARY OF DATA							
Number of missing data patterns	4						
Number of y missing data patterns	4						
Number of u missing data patterns	0						
COVARIANCE COVERAGE OF DATA							
Minimum covariance coverage value	0.100						
PROPORTION OF DATA PRESENT FOR Y							
Covariance Coverage							
SLPNSSNC							
WORRYNC							
SOLC							
SLPDURC							
SLPNSSNC	1.000						
WORRYNC	1.000	1.000					
SOLC	0.976	0.976	0.976				
SLPDURC	0.963	0.963	0.940	0.963			
UNIVARIATE SAMPLE STATISTICS							
UNIVARIATE HIGHER-ORDER MOMENT							
DESCRIPTIVE STATISTICS							
Variable/ Percentiles	Mean/	Skewness/	Minimum/	% with			
Sample Size	Variance	Kurtosis	Maximum				
Min/Max	20%/60%	40%/80%	Median				

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-121739.377	723035	538	-121739.377	798821	423
-121739.377	863094	147	-121739.377	507218	613
-121739.377	685268	596	-121739.377	903633	553
-121739.377	467561	819	-121739.377	424223	900
-121739.377	471398	74	-121739.377	81117	305
-121739.377	646573	741	-121739.377	484116	915
-121739.377	996231	310	-121739.377	86698	893
-121739.377	679832	302	-121739.377	518828	432
-121739.377	251680	794	-121739.377	70118	104
-121739.377	404042	675	-121739.377	461866	722
-121739.377	546943	865	-121739.377	850840	232
-121739.377	903420	5	-121739.377	25127	107
-121739.377	148918	682	-121739.377	131856	90
-121739.377	77571	980	-121739.377	794236	127
-121739.377	238906	430	-121739.377	402699	604
-121739.377	930872	277	-121739.377	464179	106
-121739.377	278661	674	-121739.377	178031	720
-121739.377	156536	245	-121739.377	61587	400
-121739.377	758647	951	-121739.377	797594	234
-121739.377	268217	83	-121739.377	404426	344
-121739.377	549244	756	-121739.377	545108	667
-121739.377	495366	964	-121739.377	562716	300
-121739.377	329127	185	-121739.377	391368	802
-121739.377	645052	910	-121739.377	153053	378
-121739.377	84013	598	-121739.377	995249	525
-121739.377	853781	716	-121739.377	926797	406
-121739.377	584397	428	-121739.377	458181	189
-121739.377	685657	69	-121739.377	714455	476
-121739.377	775881	778	-121739.377	829540	324
-121739.377	178475	231	-121739.377	216565	474
-121739.377	622173	992	-121739.377	462953	7
-121739.377	548245	818	-121739.377	49293	707
-121739.377	36714	201	-121739.377	478341	731
-121739.377	455617	242	-121739.377	914505	838
-121739.377	94573	983	-121739.377	266038	672
-121739.377	926762	704	-121739.377	730868	977
-121739.377	792993	859	-121739.377	478421	311
-121739.377	291149	536	-121739.377	677062	680
-121739.377	421731	886	-121739.377	544048	87
-121739.377	732596	320	-121739.377	130011	587
-121739.377	124999	96	-121739.377	690596	858
-121739.377	85734	411	-121739.377	327140	678
-121739.377	484501	163	-121739.377	484687	306
-121739.377	696830	668	-121739.377	875667	367
-121739.377	399671	13	-121739.377	42523	730
-121739.377	741888	138	-121739.377	275475	413
-121739.377	3307	735	-121739.377	168888	817
-121739.377	436806	883	-121739.377	92564	583
-121739.377	283492	435	-121739.377	534193	689
-121739.377	210139	991	-121739.377	444228	860
-121739.377	399848	220	-121739.377	344422	296
-121739.377	848163	47	-121739.377	972873	157
-121739.377	297518	166	-121739.377	264081	186
-121739.377	856536	144	-121739.377	496710	386
-121739.377	587946	120	-121739.377	535804	111
-121739.377	741484	441	-121739.377	137377	397
-121739.377	967237	48	-121739.377	626891	32
-121739.377	836515	341	-121739.377	373815	618
-121739.377	317640	437	-121739.377	534864	307
-121739.377	965639	463	-121739.377	738393	619
-121739.377	520177	262	-121739.377	376411	473
-121739.377	922596	456	-121739.377	213532	503
-121739.377	860772	174	-121739.377	153394	429
-121739.377	193042	316	-121739.377	155749	960
-121739.377	114433	708	-121739.377	136842	58
-121739.377	292884	103	-121739.377	442072	625
-121739.377	898745	466	-121739.377	724519	333
-121739.377	373505	88	-121739.377	997222	229
-121739.377	150531	154	-121739.377	650371	14
-121739.377	760599	832	-121739.377	696773	80

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-121739.377	863691	481	-121739.377	769907	457
-121739.377	287107	864	-121739.377	954914	911
-121739.377	349360	464			
-121739.377	66276	217	THE BEST LOGLIKELIHOOD VALUE HAS BEEN REPLICATED. RERUN WITH AT LEAST TWICE THE RANDOM STARTS TO CHECK THAT THE BEST LOGLIKELIHOOD IS STILL OBTAINED AND REPLICATED.		
-121739.377	121425	511			
-121739.377	761302	744			
-121739.377	804660	260			
-121739.377	950604	172			
-121739.377	824956	607			
-121739.377	94610	884	THE MODEL ESTIMATION TERMINATED NORMALLY		
-121739.377	848969	173			
-121739.377	350608	334	MODEL FIT INFORMATION		
-121739.377	813779	92			
-121739.377	483369	270	Number of Free Parameters		44
-121739.377	655497	376			
-121739.377	814975	129	Loglikelihood		
-121739.377	659832	754			
-121739.377	891531	206	H0 Value		-121739.377
-121739.377	97300	640	H0 Scaling Correction Factor		1.0687
-121739.377	312754	562			for MLR
-121739.377	125727	899			
-121739.377	802644	874	Information Criteria		
-121739.377	565819	65			
-121739.377	501995	791	Akaike (AIC)		243566.755
-121739.377	830392	35	Bayesian (BIC)		243929.001
-121739.377	931874	141	Sample-Size Adjusted BIC		243789.170
-121739.377	566739	575			(n* = (n + 2) / 24)
-121739.377	576596	99			
-121739.377	202790	198	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THE ESTIMATED MODEL		
-121739.377	317868	740			
-121739.377	566687	597	Latent Classes		
-121739.377	535063	329			
-121739.377	576783	866	1	9661.77408	0.34752
-121739.377	782179	835	2	10017.29963	0.36031
-121739.377	749453	33	3	8122.92629	0.29217
-121739.377	840031	276			
-121739.377	636396	168	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON ESTIMATED POSTERIOR PROBABILITIES		
-121739.377	354559	73			
-121739.377	197223	332	Latent Classes		
-121739.377	303834	798			
-121739.377	746978	410	1	9661.77408	0.34752
-121739.377	194143	510	2	10017.29963	0.36031
-121739.377	227563	63	3	8122.92629	0.29217
-121739.377	39756	857			
-121739.377	107446	12	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THEIR MOST LIKELY LATENT CLASS MEMBERSHIP		
-121739.377	570681	777	Class Counts and Proportions		
-121739.377	298553	773			
-121739.377	403801	762	Latent Classes		
-121739.377	872743	374			
-121739.377	851945	18	1	11507	0.41389
-121739.377	539751	459	2	9377	0.33728
-121739.377	468036	131	3	6918	0.24883
-121739.377	358488	264	CLASSIFICATION QUALITY		
-121739.377	991329	182			
-121739.377	263221	447	Entropy		0.651
-121739.377	379823	905	Average Latent Class Probabilities for Most Likely Latent Class Membership (Row)		
-121739.377	783165	170			
-121739.377	422103	62			
-121739.377	241299	912			
-121739.377	536551	696			
-121739.377	354395	486			
-121739.377	631413	439			
-121739.377	865906	641			
-121739.377	383902	673			
-121739.377	39136	226			
-121739.377	244349	736			
-121739.377	967902	52			
-121739.377	347222	533			
-121739.377	476295	969			
-121739.377	534483	290			

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by Latent Class (Column)					SOLC WITH				
	1	2	3		SLPDURC	0.221	0.018	12.366	0.000
1	0.812	0.021	0.167	Means					
2	0.010	0.908	0.082	SLPNSSNC	1.018	0.010	103.100	0.000	
3	0.032	0.182	0.786	WORRYNC	-0.992	0.021	-47.491	0.000	
Classification Probabilities for the Most Likely Latent Class Membership (Column)					SOLC	0.821	0.019	42.708	0.000
by Latent Class (Row)					SLPDURC	0.643	0.015	43.949	0.000
	1	2	3	Variances					
1	0.968	0.010	0.023	SLPNSSNC	0.488	0.005	95.892	0.000	
2	0.024	0.850	0.126	WORRYNC	0.897	0.019	46.060	0.000	
3	0.236	0.095	0.669	SOLC	1.167	0.016	72.683	0.000	
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column)					SLPDURC	1.068	0.013	82.995	0.000
by Latent Class (Row)					Latent Class 3				
	1	2	3	SLPNSSNC WITH					
1	3.753	-0.856	0.000	WORRYNC	0.010	0.006	1.651	0.099	
2	-1.653	1.909	0.000	SOLC	0.083	0.006	13.391	0.000	
3	-1.043	-1.954	0.000	SLPDURC	0.002	0.010	0.176	0.860	
MODEL RESULTS					WORRYNC WITH				
				SOLC	-0.046	0.011	-4.199	0.000	
	Estimate	S.E.	Two-Tailed Est./S.E.	P-Value	SLPDURC	0.053	0.020	2.679	0.007
Latent Class 1					SOLC WITH				
SLPNSSNC WITH					SLPDURC	0.028	0.016	1.716	0.086
WORRYNC	-0.006	0.002	-2.581	0.010	Means				
SOLC	0.043	0.006	7.601	0.000	SLPNSSNC	0.921	0.015	61.046	0.000
SLPDURC	-0.008	0.005	-1.854	0.064	WORRYNC	0.259	0.034	7.565	0.000
WORRYNC WITH					SOLC	-0.239	0.030	-7.868	0.000
SOLC	-0.047	0.005	-10.231	0.000	SLPDURC	0.103	0.036	2.858	0.004
SLPDURC	-0.016	0.002	-6.617	0.000	Variances				
SOLC WITH					SLPNSSNC	0.477	0.006	83.275	0.000
SLPDURC	0.046	0.006	7.140	0.000	WORRYNC	0.193	0.013	14.489	0.000
Means					SOLC	0.319	0.011	29.756	0.000
SLPNSSNC	0.855	0.011	75.023	0.000	SLPDURC	0.648	0.016	40.366	0.000
WORRYNC	0.810	0.012	66.773	0.000	Categorical Latent Variables				
SOLC	-0.636	0.030	-21.139	0.000	Means				
SLPDURC	-0.744	0.016	-47.613	0.000	C#1	0.173	0.110	1.581	0.114
Variances					C#2	0.210	0.067	3.148	0.002
SLPNSSNC	0.434	0.006	76.347	0.000	QUALITY OF NUMERICAL RESULTS				
WORRYNC	0.051	0.002	21.918	0.000	Condition Number for the Information Matrix				
SOLC	0.259	0.015	17.583	0.000	0.661E-05				
SLPDURC	0.236	0.005	50.873	0.000	(ratio of smallest to largest eigenvalue)				
Latent Class 2					TECHNICAL 11 OUTPUT				
SLPNSSNC WITH					Random Starts Specifications for the k-1 Class Analysis Model				
WORRYNC	-0.013	0.009	-1.441	0.150	Number of initial stage random starts				
SOLC	0.157	0.010	15.261	0.000	1000				
SLPDURC	0.032	0.010	3.297	0.001	Number of final stage optimizations				
WORRYNC WITH					250				
SOLC	-0.245	0.012	-20.354	0.000	VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES				
SLPDURC	-0.336	0.014	-23.324	0.000	H0 Loglikelihood Value				
					-123451.625				
					2 Times the Loglikelihood Difference				
					3424.495				
					Difference in the Number of Parameters				
					15				
					Mean				
					54.570				
					Standard Deviation				
					60.046				
					P-Value				
					0.0000				

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<p>LO-MENDELL-RUBIN ADJUSTED LRT TEST</p> <p>Value 3402.329 P-Value 0.0000</p> <p>TECHNICAL 14 OUTPUT</p> <p>Random Starts Specifications for the k-1 Class Analysis Model</p> <p>Number of initial stage random starts 1000 Number of final stage optimizations 250</p> <p>Random Starts Specification for the k-1 Class Model for Generated Data</p> <p>Number of initial stage random starts 2 Number of final stage optimizations 1</p> <p>Random Starts Specification for the k Class Model for Generated Data</p> <p>Number of initial stage random starts 50 Number of final stage optimizations 10 Number of bootstrap draws requested 250</p> <p>PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES</p> <p>H0 Loglikelihood Value -123451.625 2 Times the Loglikelihood Difference 3424.495 Difference in the Number of Parameters 15 Approximate P-Value 0.0000 Successful Bootstrap Draws 250</p> <p>WARNING: OF THE 250 BOOTSTRAP DRAWS, 224 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 3-CLASS MODEL. THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA. INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.</p> <p>PLOT INFORMATION</p> <p>The following plots are available:</p> <p>Histograms (sample values) Scatterplots (sample values) Sample means Estimated means, medians, modes, and percentiles Sample and estimated means</p>	<p>Observed individual values Estimated means and observed individual values Estimated overall and class-specific distributions</p> <p>SAVEDATA INFORMATION</p> <p>Save file Step1_LPA3M5.dat</p> <p>Order and format of variables</p> <p>SLPNSSNC F10.3 WORRYNC F10.3 SOLC F10.3 SLPDURC F10.3 CPROB1 F10.3 CPROB2 F10.3 CPROB3 F10.3 C F10.3 ID I6</p> <p>Save file format 8F10.3 I6</p> <p>Save file record length 10000</p> <p>Save missing symbol *</p> <p>DIAGRAM INFORMATION</p> <p>Mplus diagrams are currently not available for Mixture analysis. No diagram output was produced.</p> <p>Beginning Time: 17:32:22 Ending Time: 22:33:21 Elapsed Time: 05:00:59</p> <p>MUTHEN & MUTHEN 3463 Stoner Ave. Los Angeles, CA 90066</p> <p>Tel: (310) 391-9971 Fax: (310) 391-8971 Web: www.StatModel.com Support: Support@StatModel.com</p> <p>Copyright (c) 1998-2021 Muthen & Muthen</p>
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Step 2: Profile Selection

Series of LPA models were conducted to assess the number of profiles based on participants' SOL, sleep duration, daytime sleepiness and worry disrupting sleep. Each model was compared against the previous model (k-1 classes) in an iterative process to decide on the number of latent profiles (one to six) to be included. Two main indices were used, in line with existing LPA guidelines, to determine the appropriate number of profiles and whether additional profiles in the LPA model improved the model fit: Bayesian Information Criterion (BIC), and the Vuong-Lo-Mendell-Rubin adjusted likelihood ratio test (VLMR-LRT). A lower BIC value represented the preferred model. The VLMR-LRT test assists in determining when additional profiles are not an improvement and compares significance between an estimated model versus a model with one fewer profile. A non-significant VLMR-LRT

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test suggests that the more parsimonious model (fewer profiles) is the better fitting and representative model. Other model fit indicators including entropy index and smallest class size were also examined as diagnostic criteria. An entropy value of > .80 is desired, but values between .60 and .80 were treated as acceptable. Smallest profile size was examined and $n < 30$ was considered not sufficient to support generalizability.

Two-profile model

Mplus VERSION 8.7
MUTHEN & MUTHEN
07/13/2023 2:09 PM

INPUT INSTRUCTIONS

TITLE: LPA 2 profile syntax: M5: Varying means, equal variances, and varying covariances

DATA:

FILE IS mplus2.csv;

! Specifies file location for data file. Make sure data is in format appropriate for Mplus

! per Mplus manual. This data file is in individual format (one row of data per participant)

VARIABLE:

NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol
slpDur gender genderN yrgrp yrgrpN SCI_02C
slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be listed here

IDVARIABLE IS ID;

CLASSES = c (2);

! This is where you instruct Mplus on how many classes/profiles are being estimated. Initi

! model contains only one class/profile, thus it would be CLASSES = c (1). Above specifies

!profiles, and for each further iterative models the number in parentheses increases by one

!three profiles/classes would be c (3), and so on.

MISSING ARE ALL(9999);

! Used to communicate how missing data is coded in data file. Here shown with a “.” which

! all that is included in each cell with missing data in the data file

ANALYSIS:

TYPE = MIXTURE;

! LPA is a version of mixture modeling, and this instructs Mplus to analyze in this way

ESTIMATOR = MLR;

!FIML robust to non-normal data

STARTS = 1000 250;

STITERATIONS = 500;

! Default number of starts for each step of the ML estimation. First STARTS value specifies

!number of unique start values to start with, the 250 represents the 250 best unique start

!carrying forward to completion. The STITERATIONS specifies the number of ML iteration

!steps for those 250 selected start values to go through to be able to converge. This is a

!maximum number of iteration; if a model converges in less than 500 iterations it will stop

!before reaching 500 iterations.

!These values can be increased ... see “Four-Profile Final Model with Covariate Analysis

!Syntax” for an example.

LRTSTARTS = 2 1 50 10;

LRTBOOTSTRAP = 250;

!The above start values are for the defaults for the LRT statistic being run to compare th

!model fit with the model fit of a model with one less class (k-1). The BOOTSTRAP statemen

!specifies the number of bootstrap draws to inform Mplus’ bootstrapped LRT results.

MODEL:

!For a default Mplus model the LPA model does not need to be specified. However, it can be

!The model can also be modified from the Mplus default of estimating the indicator means

!(uniquely across profiles) and variances (constrained across profiles), as well as the la

!profile mean.

%OVERALL%

[slpnssNC worryNC solC slpDurC]; !estimates the indicators means for each profile. Without

!the means are freely estimated in each profile, not constrained.

slpnssNC worryNC solC slpDurC; !Label Var1-Var5 constrains the estimates of the variances

!profiles to be equal.

%c#1%

[slpnssNC worryNC solC slpDurC];

!covariances

slpnssNC WITH worryNC solC slpDurC;

worryNC WITH solC slpDurC;

solC WITH slpDurC;

%c#2%

[slpnssNC worryNC solC slpDurC];

!covariances

slpnssNC WITH worryNC solC slpDurC;

worryNC WITH solC slpDurC;

solC WITH slpDurC;

OUTPUT:

TECH11 TECH14;

! TECH1 provides parameter specifications and starting values for the analysis

! TECH8 provides optimization history for this analysis type

!TECH11 provides LRT results

!TECH14 provides bootstrapped LRT test

PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);

TYPE=PLOT3;

SAVEDATA:

FILE IS Step2_LPA2M5.dat;

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<p>! Tells Mplus where to save the output files from the analysis SAVE = CPROBABILITIES; ! The above command lines are to save the most likely profile membership for each participant and the posterior probabilities for their membership in each latent profile.</p> <p>*** WARNING in MODEL command All variables are uncorrelated with all other variables within class. Check that this is what is intended.</p> <p>LPA 2 profile syntax: M5: Varying means, equal variances, and varying covariances</p> <p>SUMMARY OF ANALYSIS</p> <table border="0"> <tr> <td>Number of groups</td> <td>1</td> </tr> <tr> <td>Number of observations</td> <td>27802</td> </tr> <tr> <td>Number of dependent variables</td> <td>4</td> </tr> <tr> <td>Number of independent variables</td> <td>0</td> </tr> <tr> <td>Number of continuous latent variables</td> <td>0</td> </tr> <tr> <td>Number of categorical latent variables</td> <td>1</td> </tr> </table> <p>Observed dependent variables</p> <table border="0"> <tr> <td>Continuous</td> <td>SLPNSSNC</td> <td>WORRYNC</td> <td>SOLC</td> <td>SLPDURC</td> </tr> </table> <p>Categorical latent variables</p> <table border="0"> <tr> <td>C</td> </tr> </table> <p>Variables with special functions</p> <table border="0"> <tr> <td>ID variable</td> <td>ID</td> </tr> </table> <p>Estimator MLR Information matrix OBSERVED Optimization Specifications for the 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<p>UNIVARIATE SAMPLE STATISTICS</p> <p>UNIVARIATE HIGHER-ORDER MOMENT DESCRIPTIVE STATISTICS</p> <table border="0"> <tr> <th>Variable/ Percentiles</th> <th>Mean/ Sample Size</th> <th>Skewness/ Variance</th> <th>Minimum/ Kurtosis</th> <th>% with Maximum</th> </tr> <tr> <td>SLPNSSNC</td> <td>0.933</td> <td>0.086</td> <td>0.000</td> <td>27.10%</td> </tr> <tr> <td>WORRYNC</td> <td>0.000</td> <td>1.000</td> <td>1.000</td> <td></td> </tr> <tr> <td>SOLC</td> <td>0.471</td> <td>-0.878</td> <td>2.000</td> <td>20.40%</td> </tr> <tr> <td>SLPDURC</td> <td>0.000</td> <td>0.967</td> <td>-1.054</td> <td>31.63%</td> </tr> <tr> <td>WORRYNC</td> <td>0.000</td> <td>-1.036</td> <td>-2.825</td> <td>1.71%</td> </tr> <tr> <td>SOLC</td> <td>0.000</td> <td>0.967</td> <td>-1.054</td> <td>31.63%</td> </tr> <tr> <td>SLPDURC</td> <td>0.000</td> <td>0.568</td> <td>-1.205</td> <td>25.70%</td> </tr> <tr> <td>WORRYNC</td> <td>0.000</td> <td>-0.470</td> <td>2.445</td> <td>3.44%</td> </tr> <tr> <td>SOLC</td> 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Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:		-127763.908	448829	938
		-127763.908	392418	28
		-127763.908	485483	498
-127763.908	512836	289		878
-127763.908	327140	678		939
-127763.908	341519	653		81
-127763.908	185071	370		416
-127763.908	265218	924		316
-127763.908	96617	955		262
-127763.908	847088	750		959
-127763.908	117391	710		627
-127763.908	580181	691		218
-127763.908	345726	461		597
-127763.908	995913	787		166
-127763.908	845580	805		900
-127763.908	728038	177		315
-127763.908	86651	993		761
-127763.908	458181	189		528
-127763.908	792993	859		142
-127763.908	715255	523		733
-127763.908	17359	227		696
-127763.908	574942	558		326
-127763.908	488125	368		273
-127763.908	599729	658		63
-127763.908	399848	220		388
-127763.908	779820	877		559
-127763.908	327927	908		704
-127763.908	496344	808		847
-127763.908	369602	146		312
-127763.908	985387	381		363
-127763.908	50887	389		305
-127763.908	546943	865		706
-127763.908	692169	308		293
-127763.908	370481	742		690
-127763.908	573096	20		526
-127763.908	629320	222		127
-127763.908	742609	531		334
-127763.908	881886	608		915
-127763.908	902278	21		889
-127763.908	150531	154		280
-127763.908	939870	655		592
-127763.908	403801	762		432
-127763.908	152496	123		511
-127763.908	440395	917		86
-127763.908	903633	553		763
-127763.908	596257	405		680
-127763.908	136842	58		314
-127763.908	77571	980		843
-127763.908	863094	147		664
-127763.908	833196	715		68
-127763.908	344422	296		232
-127763.908	529496	343		362
-127763.908	232559	136		152
-127763.908	137305	379		372
-127763.908	848890	95		604
-127763.908	723775	97		747
-127763.908	549244	756		43
-127763.908	379729	512		265
-127763.908	533738	11		700
-127763.908	699834	723		335
-127763.908	628143	854		371
-127763.908	738393	619		654
-127763.908	231281	542		501
-127763.908	645664	39		51
-127763.908	508445	946		7
-127763.908	298553	773		647
-127763.908	416463	467		53
-127763.908	690596	858		125
-127763.908	160326	546		260
-127763.908	928287	197		48

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-127763.908	625191	350	-127763.908	712702	684
-127763.908	954914	911	-127763.908	217130	443
-127763.908	288738	940	-127763.908	462662	872
-127763.908	802682	419	-127763.908	466971	109
-127763.908	622173	992	-127763.908	836515	341
-127763.908	85114	385	-127763.908	360419	356
-127763.908	30098	209	-127763.908	163110	584
-127763.908	231400	738	-127763.908	287107	864
-127763.908	96941	890	-127763.908	264521	929
-127763.908	407108	366	-127763.908	354395	486
-127763.908	972430	491	-127763.908	220492	743
-127763.908	760878	249	-127763.908	11397	944
-127763.908	838615	677	-127763.908	180889	920
-127763.908	872743	374	-127763.908	238906	430
-127763.908	742688	594	-127763.908	358488	264
-127763.908	882750	646	-127763.908	648035	836
-127763.908	425982	485	-127763.908	563584	657
-127763.908	400319	887	-127763.908	618760	489
-127763.908	915107	54	-127763.908	758647	951
-127763.908	761302	744	-127763.908	862607	468
-127763.908	695453	902	-127763.908	314757	345
-127763.908	622290	880	-127763.908	193847	354
-127763.908	134830	780	-127763.908	914505	838
-127763.908	22874	588	-127763.908	327475	518
-127763.908	995249	525	-127763.908	264901	634
-127763.908	992389	77	-127763.908	373815	618
-127763.908	650371	14	-127763.908	957392	79
-127763.908	685657	69	-127763.908	521575	313
-127763.908	399508	415	-127763.908	971853	402
-127763.908	907810	795	-127763.908	548245	818
-127763.908	264081	186	-127763.908	474357	789
-127763.908	298201	903	-127763.908	268896	124
-127763.908	751153	110	-127763.908	856536	144
-127763.908	347222	533	-127763.908	963967	941
-127763.908	527050	948	-127763.908	42523	730
-127763.908	535063	329	-127763.908	411615	171
-127763.908	735928	916	-127763.908	804561	59
-127763.908	399380	436	-127763.908	372176	23
-127763.908	246575	291	-127763.908	425929	508
-127763.908	147440	514	-127763.908	376411	473
-127763.908	739214	807	-127763.908	569131	26
-127763.908	197223	332	-127763.908	748692	204
-127763.908	871438	561	-127763.908	562716	300
-127763.908	659832	754			
-127763.908	63231	935			
-127763.908	414828	322	THE BEST LOGLIKELIHOOD VALUE HAS BEEN		
-127763.908	153053	378	REPLICATED. RERUN WITH AT LEAST TWICE THE		
-127763.908	345974	622	RANDOM STARTS TO CHECK THAT THE BEST		
-127763.908	942358	644	LOGLIKELIHOOD IS STILL OBTAINED AND		
-127763.908	848331	137	REPLICATED.		
-127763.908	481835	57			
-127763.908	459221	652	THE MODEL ESTIMATION TERMINATED NORMALLY		
-127763.908	210139	991			
-127763.908	848969	173	MODEL FIT INFORMATION		
-127763.908	213532	503			
-127763.908	939021	8	Number of Free Parameters	25	
-127763.908	326091	759	Loglikelihood		
-127763.908	829540	324			
-127763.908	442072	625	H0 Value	-127763.908	
-127763.908	860772	174	H0 Scaling Correction Factor	1.1651	
-127763.908	505244	582	for MLR		
-127763.908	580539	786			
-127763.908	97300	640	Information Criteria		
-127763.908	790452	303			
-127763.908	830292	527	Akaike (AIC)	255577.815	
-127763.908	852283	616	Bayesian (BIC)	255783.637	
-127763.908	421731	886	Sample-Size Adjusted BIC	255704.188	
-127763.908	897782	545	(n* = (n + 2) / 24)		
-127763.908	595153	230			
-127763.908	422103	62			

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THE ESTIMATED MODEL

Latent Classes		
1	6498.62894	0.23375
2	21303.37106	0.76625

FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON ESTIMATED POSTERIOR PROBABILITIES

Latent Classes		
1	6498.62894	0.23375
2	21303.37106	0.76625

FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THEIR MOST LIKELY LATENT CLASS MEMBERSHIP

Class Counts and Proportions

Latent Classes		
1	6336	0.22790
2	21466	0.77210

CLASSIFICATION QUALITY

Entropy 0.878

Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)

	1	2
1	0.940	0.060
2	0.025	0.975

Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)

	1	2
1	0.917	0.083
2	0.018	0.982

Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)

	1	2
1	2.399	0.000
2	-4.013	0.000

MODEL RESULTS

Estimate	Two-Tailed S.E.	Est./S.E.	P-Value
----------	-----------------	-----------	---------

Latent Class 1

SLPNSSNC WITH				
WORRYNC	-0.024	0.005	-5.002	0.000
SOLC	0.113	0.007	16.449	0.000
SLPDURC	0.042	0.008	5.542	0.000

WORRYNC WITH				
SOLC	-0.175	0.007	-25.662	0.000
SLPDURC	-0.139	0.007	-20.205	0.000

SOLC WITH				
SLPDURC	0.169	0.010	17.225	0.000

Means				
SLPNSSNC	0.989	0.010	100.111	0.000
WORRYNC	-1.520	0.012	-122.617	0.000
SOLC	0.816	0.018	45.861	0.000
SLPDURC	0.743	0.016	46.134	0.000

Variiances				
SLPNSSNC	0.471	0.003	155.408	0.000
WORRYNC	0.305	0.003	99.427	0.000
SOLC	0.820	0.009	92.599	0.000
SLPDURC	0.859	0.008	105.661	0.000

Latent Class 2

SLPNSSNC WITH				
WORRYNC	-0.039	0.003	-11.465	0.000
SOLC	0.136	0.005	26.590	0.000
SLPDURC	0.042	0.005	8.336	0.000

WORRYNC WITH				
SOLC	-0.283	0.005	-52.165	0.000
SLPDURC	-0.252	0.005	-46.861	0.000

SOLC WITH				
SLPDURC	0.357	0.008	42.220	0.000

Means				
SLPNSSNC	0.916	0.005	189.197	0.000
WORRYNC	0.464	0.005	102.502	0.000
SOLC	-0.243	0.006	-39.804	0.000
SLPDURC	-0.222	0.007	-33.454	0.000

Variiances				
SLPNSSNC	0.471	0.003	155.408	0.000
WORRYNC	0.305	0.003	99.427	0.000
SOLC	0.820	0.009	92.599	0.000
SLPDURC	0.859	0.008	105.661	0.000

Categorical Latent Variables

Means				
C#1	-1.187	0.018	-67.816	0.000

QUALITY OF NUMERICAL RESULTS

Condition Number for the Information Matrix
0.563E-02
(ratio of smallest to largest eigenvalue)

TECHNICAL 11 OUTPUT

Random Starts Specifications for the k-1 Class Analysis Model

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<p>Number of initial stage random starts 1000 Number of final stage optimizations 250</p> <p>VUONG-LO-MENDELLE-RUBIN LIKELIHOOD RATIO TEST FOR 1 (H0) VERSUS 2 CLASSES</p> <p>H0 Loglikelihood Value -132903.306 2 Times the Loglikelihood Difference 10278.796 Difference in the Number of Parameters 11 Mean 17.872 Standard Deviation 10.707 P-Value 0.0000</p> <p>LO-MENDELLE-RUBIN ADJUSTED LRT TEST</p> <p>Value 10188.283 P-Value 0.0000</p> <p>TECHNICAL 14 OUTPUT</p> <p>Random Starts Specifications for the k-1 Class Analysis Model</p> <p>Number of initial stage random starts 1000 Number of final stage optimizations 250</p> <p>Random Starts Specification for the k-1 Class Model for Generated Data</p> <p>Number of initial stage random starts 2 Number of final stage optimizations 1</p> <p>Random Starts Specification for the k Class Model for Generated Data</p> <p>Number of initial stage random starts 50 Number of final stage optimizations 10 Number of bootstrap draws requested 250</p> <p>PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 1 (H0) VERSUS 2 CLASSES</p> <p>H0 Loglikelihood Value -132903.306 2 Times the Loglikelihood Difference 10278.796 Difference in the Number of Parameters 11 Approximate P-Value 0.0000 Successful Bootstrap Draws 249</p> <p>WARNING: OF THE 249 BOOTSTRAP DRAWS, 225 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 2-CLASS MODEL. THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA. INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.</p> <p>WARNING: 1 OUT OF 250 BOOTSTRAP DRAWS DID NOT CONVERGE. INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.</p>	<p>PLOT INFORMATION</p> <p>The following plots are available:</p> <p>Histograms (sample values) Scatterplots (sample values) Sample means Estimated means, medians, modes, and percentiles Sample and estimated means Observed individual values Estimated means and observed individual values Estimated overall and class-specific distributions</p> <p>SAVEDATA INFORMATION</p> <p>Save file Step2_LPA2M5.dat</p> <p>Order and format of variables</p> <p>SLPNSSNC F10.3 WORRYNC F10.3 SOLC F10.3 SLPDURC F10.3 CPROB1 F10.3 CPROB2 F10.3 C F10.3 ID I6</p> <p>Save file format 7F10.3 I6</p> <p>Save file record length 10000</p> <p>Save missing symbol *</p> <p>DIAGRAM INFORMATION</p> <p>Mplus diagrams are currently not available for Mixture analysis. No diagram output was produced.</p> <p>Beginning Time: 14:09:11 Ending Time: 15:19:05 Elapsed Time: 01:09:54</p> <p>MUTHEN & MUTHEN 3463 Stoner Ave. Los Angeles, CA 90066</p> <p>Tel: (310) 391-9971 Fax: (310) 391-8971 Web: www.StatModel.com Support: Support@StatModel.com</p> <p>Copyright (c) 1998-2021 Muthen & Muthen</p>
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Three-profile model

<p>Mplus VERSION 8.7 MUTHEN & MUTHEN 07/20/2023 9:24 AM</p> <p>INPUT INSTRUCTIONS</p> <p>TITLE: LPA 3 profile syntax: M5: Varying means, equal variances, and varying covariances</p>	<p>DATA: FILE IS mplus2.csv; ! Specifies file location for data file. Make sure data is in format appropriate for Mplus ! per Mplus manual. This data file is in individual format (one row of data per participant) VARIABLE:</p>
---	--

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```

NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol
slpDur gender genderN yrgrp yrgrpN SCI_02C
slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be
listed here
IDVARIABLE IS ID;
CLASSES = c (3);
! This is where you instruct Mplus on how many
classes/profiles are being estimated. Initi
! model contains only one class/profile, thus it would be
CLASSES = c (1). Above specifies
!profiles, and for each further iterative models the number
in parentheses increases by on
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(9999);
! Used to communicate how missing data is coded in data
file. Here shown with a “.” which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 1000 250;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifie
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will sto
!before reaching 500 iterations.
!These values can be increased ... see “Four-Profile Final
Model with Covariate Analysis
!Syntax” for an example.
LRTSTARTS = 2 1 50 10;
LRTBOOTSTRAP = 250;
!The above start values are for the defaults for the LRT
statistic being run to compare th
!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statemen
!specifies the number of bootstrap draws to inform Mplus’
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances
!profiles to be equal.
%c#1%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#2%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#3%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

OUTPUT:
TECH11 TECH14;
! TECH1 provides parameter specifications and starting
values for the analysis
! TECH8 provides optimization history for this analysis type
!TECH11 provides LRT results
!TECH14 provides bootstrapped LRT test
PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);
TYPE=PLOT3;
SAVEDATA:
FILE IS Step1_LPA3M5.dat;
! Tells Mplus where to save the output files from the
analysis
SAVE = CPROBABILITIES;
! The above command lines are to save the most likely
profile membership for each particip
! and the posterior probabilities for their membership in
each latent profile.

*** WARNING in MODEL command
All variables are uncorrelated with all other variables within
class.
Check that this is what is intended.

LPA 3 profile syntax: M5: Varying means, equal variances,
and varying covariances

SUMMARY OF ANALYSIS

Number of groups 1
Number of observations 27802

Number of dependent variables 4
Number of independent variables 0
Number of continuous latent variables 0
Number of categorical latent variables 1

Observed dependent variables

Continuous
SLPNSSNC WORRYNC SOLC SLPDURC

Categorical latent variables
C

```

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Variables with special functions		SLPDURC	0.963	0.963	0.940	0.963
UNIVARIATE SAMPLE STATISTICS						
UNIVARIATE HIGHER-ORDER MOMENT DESCRIPTIVE STATISTICS						
ID variable	ID	Variable/	Mean/	Skewness/	Minimum/	% with
Estimator	MLR	Percentiles				
Information matrix	OBSERVED	Sample Size	Variance	Kurtosis	Maximum	
Optimization Specifications for the Quasi-Newton Algorithm for		Min/Max	20%/60%	40%/80%	Median	
Continuous Outcomes						
Maximum number of iterations	100	SLPNSSNC	0.933	0.086	0.000	27.10%
Convergence criterion	0.100D-05	0.000	1.000	1.000		
Optimization Specifications for the EM Algorithm		27802.000	0.471	-0.878	2.000	20.40%
Maximum number of iterations	500	1.000	2.000			
Convergence criteria		WORRYNC	0.000	-1.036	-2.825	1.71%
Loglikelihood change	0.100D-06	-0.928	0.020	0.495		
Relative loglikelihood change	0.100D-06	27802.000	1.000	0.165	0.969	28.55%
Derivative	0.100D-05	0.495	0.969			
Optimization Specifications for the M step of the EM Algorithm for		0.495	0.969			
Categorical Latent variables		SOLC	0.000	0.967	-1.054	31.63%
Number of M step iterations	1	1.054	-0.072	-0.072		
M step convergence criterion	0.100D-05	27123.000	1.000	0.467	2.875	2.77%
Basis for M step termination	ITERATION	-0.072	0.910			
Optimization Specifications for the M step of the EM Algorithm for		SLPDURC	0.000	0.568	-1.205	25.70%
Censored, Binary or Ordered Categorical (Ordinal), Unordered		-1.205	-0.292	-0.292		
Categorical (Nominal) and Count Outcomes		26763.000	1.000	-0.470	2.445	3.44%
Number of M step iterations	1	-0.292	0.620			
M step convergence criterion	0.100D-05					
Basis for M step termination	ITERATION					
Maximum value for logit thresholds	15					
Minimum value for logit thresholds	-15					
Minimum expected cell size for chi-square	0.100D-01					
Maximum number of iterations for H1	2000					
Convergence criterion for H1	0.100D-03					
Optimization algorithm	EMA					
Random Starts Specifications						
Number of initial stage random starts	1000	-125994.415	691234	250		
Number of final stage optimizations	250	-125994.415	246794	801		
Number of initial stage iterations	500	-125994.415	917702	693		
Initial stage convergence criterion	0.100D+01	-125994.415	970689	266		
Random starts scale	0.500D+01	-125994.415	244349	736		
Random seed for generating random starts	0	-125994.415	478421	311		
Input data file(s)		-125994.415	278661	674		
mplus2.csv		-125994.415	930872	277		
Input data format	FREE	-125994.415	618000	190		
		-125994.415	49910	829		
		-125994.415	314034	513		
		-125994.415	91231	727		
		-125994.415	851945	18		
		-125994.415	926797	406		
Number of missing data patterns	4	-125994.415	791285	416		
Number of y missing data patterns	4	-125994.415	544077	809		
Number of u missing data patterns	0	-125994.415	268896	124		
		-125994.415	27071	15		
		-125994.415	177175	851		
		-125994.415	485256	371		
Minimum covariance coverage value	0.100	-125994.415	495490	990		
		-125994.415	349263	263		
		-125994.415	3607	873		
		-125994.415	168762	200		
		-125994.415	109357	765		
		-125994.415	544009	842		
		-125994.415	180889	920		
		-125994.415	636396	168		
		-125994.415	772131	407		
		-125994.415	383979	603		
		-125994.415	373702	669		
SLPNSSNC	1.000					
WORRYNC	1.000	1.000				
SOLC	0.976	0.976	0.976			

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-125994.415 551340	766	-125994.415 673496	218
-125994.415 303834	798	-125994.415 440368	797
-125994.415 294811	637	-125994.415 628143	854
-125994.415 383902	673	-125994.415 847088	750
-125994.415 462228	298	-125994.415 954914	911
-125994.415 215353	164	-125994.415 271809	846
-125994.415 200041	810	-125994.415 679832	302
-125994.415 130541	896	-125994.415 809240	543
-125994.415 616917	261	-125994.415 172913	557
-125994.415 126371	526	-125994.415 830392	35
-125994.415 849670	347	-125994.415 97300	640
-125994.415 237332	661	-125994.415 597614	284
-125994.415 114433	708	-125994.415 723035	538
-125994.415 903633	553	-125994.415 682718	733
-125994.415 202790	198	-125994.415 81951	907
-125994.415 833196	715	-125994.415 879338	309
-125994.415 137377	397	-125994.415 966499	963
-125994.415 399508	415	-125994.415 669634	335
-125994.415 862607	468	-125994.415 863691	481
-125994.415 544048	87	-125994.415 717754	180
-125994.415 414284	158	-125994.415 654136	167
-125994.415 490123	995	-125994.415 193569	440
-125994.415 937885	426	-125994.415 298201	903
-125994.415 105435	265	-125994.415 347515	24
-125994.415 46502	714	-125994.415 545108	667
-125994.415 992389	77	-125994.415 980970	894
-125994.415 308582	752	-125994.415 695155	150
-125994.415 373815	618	-125994.415 766903	505
-125994.415 856536	144	-125994.415 922596	456
-125994.415 679448	937	-125994.415 51375	148
-125994.415 505879	424	-125994.415 158612	581
-125994.415 224950	455	-125994.415 364676	27
-125994.415 260953	589	-125994.415 566739	575
-125994.415 339073	841	-125994.415 442072	625
-125994.415 575700	100	-125994.415 322790	636
-125994.415 549244	756	-125994.415 852283	616
-125994.415 391949	295	-125994.415 677062	680
-125994.415 701525	239	-125994.415 696830	668
-125994.415 576596	99	-125994.415 792389	746
-125994.415 794236	127	-125994.415 535303	923
-125994.415 484687	306	-125994.415 17896	592
-125994.415 972430	491	-125994.415 857122	889
-125994.415 971693	470	-125994.415 735928	916
-125994.415 453915	975	-125994.415 961454	665
-125994.415 710154	831	-125994.415 881886	608
-125994.415 850112	922	-125994.415 370481	742
-125994.415 443442	380	-125994.415 195353	225
-125994.415 444228	860	-125994.415 405371	569
-125994.415 179022	687	-125994.415 415931	10
-125994.415 50887	389	-125994.415 148918	682
-125994.415 70118	104	-125994.415 119513	821
-125994.415 204959	695	-125994.415 303634	169
-125994.415 570908	98	-125994.415 465160	862
-125994.415 907810	795	-125994.415 414828	322
-125994.415 705224	953	-125994.415 333082	578
-125994.415 963967	941	-125994.415 928624	981
-125994.415 923437	398	-125994.415 548245	818
-125994.415 138695	783	-125994.415 599729	658
-125994.415 124999	96	-125994.415 207896	25
-125994.415 634782	979	-125994.415 485483	498
-125994.415 563584	657	-125994.415 640833	434
-125994.415 742609	531	-125994.415 762461	425
-125994.415 836515	341	-125994.415 136842	58
-125994.415 848356	602	-125994.415 311214	64
-125994.415 298553	773	-125994.415 72662	729
-125994.415 745972	521	-125994.415 297518	166
-125994.415 965639	463	-125994.415 61587	400
-125994.415 264081	186	-125994.415 569833	85
-125994.415 213532	503	-125994.415 662718	460
-125994.415 284384	600	-125994.415 484501	163

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-125994.415	432513	803	-125994.415	592219	119
-125994.415	843555	952	-125994.415	891531	206
-125994.415	unperturbed	0	-125994.415	265218	924
-125994.415	626208	698	-125994.415	370957	554
-125994.415	599136	811	-125994.415	264935	281
-125994.415	721392	768	-125994.415	127362	757
-125994.415	153394	429	-125994.415	621542	375
-125994.415	178031	720	-125994.415	326091	759
-125994.415	728038	177	-125994.415	691041	590
-125994.415	484406	421			
-125994.415	224151	973			
-125994.415	589483	950	THE BEST LOGLIKELIHOOD VALUE HAS BEEN		
-125994.415	782821	272	REPLICATED. RERUN WITH AT LEAST TWICE THE		
-125994.415	352277	42	RANDOM STARTS TO CHECK THAT THE BEST		
-125994.415	238906	430	LOGLIKELIHOOD IS STILL OBTAINED AND		
-125994.415	848331	137	REPLICATED.		
-125994.415	855760	593			
-125994.415	891347	504	THE MODEL ESTIMATION TERMINATED NORMALLY		
-125994.415	468036	131			
-125994.415	437181	135	MODEL FIT INFORMATION		
-125994.415	552272	654			
-125994.415	520177	262	Number of Free Parameters	36	
-125994.415	268217	83			
-125994.415	580405	286	Loglikelihood		
-125994.415	87586	871			
-125994.415	638977	643	H0 Value	-125994.415	
-125994.415	55115	408	H0 Scaling Correction Factor	1.3296	
-125994.415	248742	556	for MLR		
-125994.415	838615	677			
-125994.415	751153	110	Information Criteria		
-125994.415	384199	882			
-125994.415	178181	753	Akaike (AIC)	252060.830	
-125994.415	22362	365	Bayesian (BIC)	252357.213	
-125994.415	792993	859	Sample-Size Adjusted BIC	252242.806	
-125994.415	778953	635	(n* = (n + 2) / 24)		
-125994.415	286735	175			
-125994.415	617243	237	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-125994.415	605161	409	LATENT CLASSES		
-125994.415	846194	93	BASED ON THE ESTIMATED MODEL		
-125994.415	357866	968			
-125994.415	726035	191	Latent		
-125994.415	483369	270	Classes		
-125994.415	399380	436			
-125994.415	434915	552	1	5737.65936	0.20638
-125994.415	782200	84	2	18891.89637	0.67952
-125994.415	538872	949	3	3172.44427	0.11411
-125994.415	871851	257			
-125994.415	814975	129	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-125994.415	157351	579	LATENT CLASSES		
-125994.415	81233	825	BASED ON ESTIMATED POSTERIOR PROBABILITIES		
-125994.415	526324	178			
-125994.415	366706	29	Latent		
-125994.415	211281	292	Classes		
-125994.415	848969	173			
-125994.415	937588	293	1	5737.65936	0.20638
-125994.415	788796	145	2	18891.89637	0.67952
-125994.415	574942	558	3	3172.44427	0.11411
-125994.415	35191	703			
-125994.415	195873	6	FINAL CLASS COUNTS AND PROPORTIONS FOR THE		
-125994.415	118438	601	LATENT CLASSES		
-125994.415	576726	280	BASED ON THEIR MOST LIKELY LATENT CLASS		
-125994.415	345974	622	MEMBERSHIP		
-125994.415	467339	66			
-125994.415	566687	597	Class Counts and Proportions		
-125994.415	407108	366			
-125994.415	791678	974	Latent		
-125994.415	608460	244	Classes		
-125994.415	860029	760			
-125994.415	14262	781	1	6141	0.22088
-125994.415	50983	834	2	18658	0.67110
			3	3003	0.10801

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

CLASSIFICATION QUALITY					WORRYNC	-0.030	0.004	-8.232	0.000
Entropy					SOLC	0.118	0.005	22.000	0.000
0.854					SLPDURC	0.037	0.006	6.407	0.000
Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)					WORRYNC WITH				
1 2 3					SOLC	-0.192	0.008	-25.245	0.000
1 0.822 0.116 0.063					SLPDURC	-0.187	0.010	-18.506	0.000
2 0.026 0.974 0.000					SOLC WITH				
3 0.071 0.001 0.928					SLPDURC	0.290	0.011	27.050	0.000
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					Means				
1 2 3					SLPNSSNC	0.904	0.005	171.611	0.000
1 0.879 0.083 0.037					WORRYNC	0.570	0.006	101.171	0.000
2 0.038 0.962 0.000					SOLC	-0.364	0.009	-40.558	0.000
3 0.122 0.000 0.878					SLPDURC	-0.300	0.007	-40.716	0.000
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					Variances				
1 2 3					SLPNSSNC	0.468	0.003	152.809	0.000
1 3.164 0.809 0.000					WORRYNC	0.210	0.009	23.619	0.000
2 5.521 8.764 0.000					SOLC	0.626	0.013	47.707	0.000
3 -1.978 -10.865 0.000					SLPDURC	0.798	0.010	76.790	0.000
MODEL RESULTS					Latent Class 3				
Estimate					SLPNSSNC WITH				
Two-Tailed					WORRYNC	0.009	0.006	1.533	0.125
S.E. Est./S.E. P-Value					SOLC	0.046	0.011	4.198	0.000
Latent Class 1					SLPDURC	0.013	0.012	1.153	0.249
SLPNSSNC WITH					WORRYNC WITH				
WORRYNC 0.015 0.006 2.633 0.008					SOLC	-0.006	0.006	-0.926	0.354
SOLC 0.106 0.008 14.059 0.000					SLPDURC	-0.070	0.009	-7.748	0.000
SLPDURC 0.019 0.009 2.191 0.028					SOLC WITH				
WORRYNC WITH					SLPDURC	0.050	0.015	3.310	0.001
SOLC 0.088 0.017 5.295 0.000					Means				
SLPDURC -0.048 0.010 -4.925 0.000					SLPNSSNC	1.112	0.015	76.093	0.000
SOLC WITH					WORRYNC	-1.994	0.027	-74.945	0.000
SLPDURC 0.051 0.013 3.916 0.000					SOLC	1.572	0.036	43.121	0.000
Means					SLPDURC	1.108	0.027	40.596	0.000
SLPNSSNC 0.931 0.012 78.488 0.000					Variances				
WORRYNC -0.775 0.027 -28.725 0.000					SLPNSSNC	0.468	0.003	152.809	0.000
SOLC 0.353 0.018 19.838 0.000					WORRYNC	0.210	0.009	23.619	0.000
SLPDURC 0.391 0.023 17.144 0.000					SOLC	0.626	0.013	47.707	0.000
Variances					SLPDURC	0.798	0.010	76.790	0.000
SLPNSSNC 0.468 0.003 152.809 0.000					Categorical Latent Variables				
WORRYNC 0.210 0.009 23.619 0.000					Means				
SOLC 0.626 0.013 47.707 0.000					C#1	0.593	0.046	12.891	0.000
SLPDURC 0.798 0.010 76.790 0.000					C#2	1.784	0.040	44.896	0.000
Latent Class 2					QUALITY OF NUMERICAL RESULTS				
SLPNSSNC WITH					Condition Number for the Information Matrix				
					0.903E-04				
					(ratio of smallest to largest eigenvalue)				
					TECHNICAL 11 OUTPUT				
					Random Starts Specifications for the k-1 Class Analysis Model				
					Number of initial stage random starts 1000				
					Number of final stage optimizations 250				
					VUONG-LO-MENDELLE-RUBIN LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES				

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<table border="0"> <tr> <td>H0 Loglikelihood Value</td> <td>-127763.908</td> </tr> <tr> <td>2 Times the Loglikelihood Difference</td> <td>3538.985</td> </tr> <tr> <td>Difference in the Number of Parameters</td> <td>11</td> </tr> <tr> <td>Mean</td> <td>61.969</td> </tr> <tr> <td>Standard Deviation</td> <td>60.777</td> </tr> <tr> <td>P-Value</td> <td>0.0000</td> </tr> </table> <p>LO-MENDELL-RUBIN ADJUSTED LRT TEST</p> <table border="0"> <tr> <td>Value</td> <td>3507.822</td> </tr> <tr> <td>P-Value</td> <td>0.0000</td> </tr> </table> <p>TECHNICAL 14 OUTPUT</p> <p>Random Starts Specifications for the k-1 Class Analysis Model</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>1000</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>250</td> </tr> </table> <p>Random Starts Specification for the k-1 Class Model for Generated Data</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>2</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>1</td> </tr> </table> <p>Random Starts Specification for the k Class Model for Generated Data</p> <table border="0"> <tr> <td>Number of initial stage random starts</td> <td>50</td> </tr> <tr> <td>Number of final stage optimizations</td> <td>10</td> </tr> <tr> <td>Number of bootstrap draws requested</td> <td>250</td> </tr> </table> <p>PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 2 (H0) VERSUS 3 CLASSES</p> <table border="0"> <tr> <td>H0 Loglikelihood Value</td> <td>-127763.908</td> </tr> <tr> <td>2 Times the Loglikelihood Difference</td> <td>3538.985</td> </tr> <tr> <td>Difference in the Number of Parameters</td> <td>11</td> </tr> <tr> <td>Approximate P-Value</td> <td>0.0000</td> </tr> <tr> <td>Successful Bootstrap Draws</td> <td>250</td> </tr> </table> <p>WARNING: OF THE 250 BOOTSTRAP DRAWS, 230 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 3-CLASS MODEL. THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA. INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.</p> <p>PLOT INFORMATION</p> <p>The following plots are available:</p> <p>Histograms (sample values)</p>	H0 Loglikelihood Value	-127763.908	2 Times the Loglikelihood Difference	3538.985	Difference in the Number of Parameters	11	Mean	61.969	Standard Deviation	60.777	P-Value	0.0000	Value	3507.822	P-Value	0.0000	Number of initial stage random starts	1000	Number of final stage optimizations	250	Number of initial stage random starts	2	Number of final stage optimizations	1	Number of initial stage random starts	50	Number of final stage optimizations	10	Number of bootstrap draws requested	250	H0 Loglikelihood Value	-127763.908	2 Times the Loglikelihood Difference	3538.985	Difference in the Number of Parameters	11	Approximate P-Value	0.0000	Successful Bootstrap Draws	250	<p>Scatterplots (sample values)</p> <p>Sample means</p> <p>Estimated means, medians, modes, and percentiles</p> <p>Sample and estimated means</p> <p>Observed individual values</p> <p>Estimated means and observed individual values</p> <p>Estimated overall and class-specific distributions</p> <p>SAVE DATA INFORMATION</p> <p>Save file</p> <p>Step1_LPA3M5.dat</p> <p>Order and format of variables</p> <table border="0"> <tr> <td>SLPNSSNC</td> <td>F10.3</td> </tr> <tr> <td>WORRYNC</td> <td>F10.3</td> </tr> <tr> <td>SOLC</td> <td>F10.3</td> </tr> <tr> <td>SLPDURC</td> <td>F10.3</td> </tr> <tr> <td>CPROB1</td> <td>F10.3</td> </tr> <tr> <td>CPROB2</td> <td>F10.3</td> </tr> <tr> <td>CPROB3</td> <td>F10.3</td> </tr> <tr> <td>C</td> <td>F10.3</td> </tr> <tr> <td>ID</td> <td>I6</td> </tr> </table> <p>Save file format</p> <p>8F10.3 I6</p> <p>Save file record length 10000</p> <p>Save missing symbol *</p> <p>DIAGRAM INFORMATION</p> <p>Mplus diagrams are currently not available for Mixture analysis. No diagram output was produced.</p> <p>Beginning Time: 09:24:01 Ending Time: 11:07:04 Elapsed Time: 01:43:03</p> <p>MUTHEN & MUTHEN 3463 Stoner Ave. Los Angeles, CA 90066</p> <p>Tel: (310) 391-9971 Fax: (310) 391-8971 Web: www.StatModel.com Support: Support@StatModel.com</p> <p>Copyright (c) 1998-2021 Muthen & Muthen</p>	SLPNSSNC	F10.3	WORRYNC	F10.3	SOLC	F10.3	SLPDURC	F10.3	CPROB1	F10.3	CPROB2	F10.3	CPROB3	F10.3	C	F10.3	ID	I6
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Four-profile model

<p>Mplus VERSION 8.7 MUTHEN & MUTHEN 07/25/2023 1:46 PM</p> <p>INPUT INSTRUCTIONS</p> <p>TITLE: LPA 4 profile syntax: M5: Varying means, equal variances, and varying covariances</p> <p>DATA: FILE IS mplus2.csv; ! Specifies file location for data file. Make sure data is in format appropriate for Mplus</p>	<p>! per Mplus manual. This data file is in individual format (one row of data per participant)</p> <p>VARIABLE: NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol slpDur gender genderN yrgrp yrgrpN SCI_02C slpnssNC worryNC solC slpDurC;</p> <p>! All variables included in data file should be named here. USEVARIABLES ARE slpnssNC worryNC solC slpDurC; ! Only variables intended for use in the analysis should be listed here IDVARIABLE IS ID; CLASSES = c (4);</p>
---	---

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```

! This is where you instruct Mplus on how many
classes/profiles are being estimated. Init
! model contains only one class/profile, thus it would be
CLASSES = c (1). Above specifies
!profiles, and for each further iterative models the number
in parentheses increases by on
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(999999);
! Used to communicate how missing data is coded in data
file. Here shown with a “.” which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 1000 250;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifie
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will sto
!before reaching 500 iterations.
!These values can be increased ... see “Four-Profile Final
Model with Covariate Analysis
!Syntax” for an example.
LRTSTARTS = 2 1 50 10;
LRTBOOTSTRAP = 250;
!The above start values are for the defaults for the LRT
statistic being run to compare th
!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statemen
!specifies the number of bootstrap draws to inform Mplus’
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances
!profiles to be equal.

%c#1%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#2%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#3%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#4%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

OUTPUT:
TECH11 TECH14;
! TECH1 provides parameter specifications and starting
values for the analysis
! TECH8 provides optimization history for this analysis type
!TECH11 provides LRT results
!TECH14 provides bootstrapped LRT test
PLOT: SERIES=slpnssNC worryNC solC slpDurC(*);
TYPE=PLOT3;
SAVEDATA:
FILE IS Step2_LPA4M5.dat;
! Tells Mplus where to save the output files from the
analysis
SAVE = CPROBABILITIES;
! The above command lines are to save the most likely
profile membership for each particip
! and the posterior probabilities for their membership in
each latent profile.

*** WARNING in MODEL command
All variables are uncorrelated with all other variables within
class.
Check that this is what is intended.

LPA 4 profile syntax: M5: Varying means, equal variances,
and varying covariances

SUMMARY OF ANALYSIS

Number of groups 1
Number of observations 27802

Number of dependent variables 4
Number of independent variables 0
Number of continuous latent variables 0
Number of categorical latent variables 1

Observed dependent variables

Continuous
SLPNSSNC WORRYNC SOLC SLPDURC

Categorical latent variables
C

```

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Variables with special functions				UNIVARIATE SAMPLE STATISTICS						
ID variable	ID			UNIVARIATE HIGHER-ORDER MOMENT DESCRIPTIVE STATISTICS						
Estimator		MLR								
Information matrix		OBSERVED								
Optimization Specifications for the Quasi-Newton Algorithm for				Variable/	Mean/	Skewness/	Minimum/	% with		
Continuous Outcomes				Percentiles						
Maximum number of iterations		100		Sample Size	Variance	Kurtosis	Maximum			
Convergence criterion		0.100D-05		Min/Max	20%/60%	40%/80%	Median			
Optimization Specifications for the EM Algorithm				SLPNSSNC	0.933	0.086	0.000	27.10%		
Maximum number of iterations		500		0.000	1.000	1.000				
Convergence criteria				27802.000	0.471	-0.878	2.000	20.40%		
Loglikelihood change		0.100D-06		1.000	2.000					
Relative loglikelihood change		0.100D-06		WORRYNC	0.000	-1.036	-2.825	1.71%		
Derivative		0.100D-05		-0.928	0.020	0.495				
Optimization Specifications for the M step of the EM Algorithm for				27802.000	1.000	0.165	0.969	28.55%		
Categorical Latent variables				0.495	0.969					
Number of M step iterations		1		SOLC	0.000	0.967	-1.054	31.63%		
M step convergence criterion		0.100D-05		1.054	-0.072	-0.072				
Basis for M step termination		ITERATION		27123.000	1.000	0.467	2.875	2.77%		
Optimization Specifications for the M step of the EM Algorithm for				-0.072	0.910					
Censored, Binary or Ordered Categorical (Ordinal), Unordered				SLPDURC	0.000	0.568	-1.205	25.70%		
Categorical (Nominal) and Count Outcomes				-1.205	-0.292	-0.292				
Number of M step iterations		1		26763.000	1.000	-0.470	2.445	3.44%		
M step convergence criterion		0.100D-05		-0.292	0.620					
Basis for M step termination		ITERATION								
Maximum value for logit thresholds		15		RANDOM STARTS RESULTS RANKED FROM THE BEST TO THE WORST LOGLIKELIHOOD VALUES						
Minimum value for logit thresholds		-15		Unperturbed starting value run did not converge in the initial stage						
Minimum expected cell size for chi-square		0.100D-01		optimizations.						
Maximum number of iterations for H1		2000		696 perturbed starting value run(s) did not converge in the initial stage						
Convergence criterion for H1		0.100D-03		optimizations.						
Optimization algorithm		EMA		Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:						
Random Starts Specifications										
Number of initial stage random starts		1000		-124709.900	182524	373				
Number of final stage optimizations		250		-124709.900	42523	730				
Number of initial stage iterations		500		-124709.900	512836	289				
Initial stage convergence criterion		0.100D+01		-124709.900	373702	669				
Random starts scale		0.500D+01		-124709.900	497522	502				
Random seed for generating random starts		0		-124709.900	299977	956				
Input data file(s)				-124709.900	298201	903				
mplus2.csv				-124709.900	944186	541				
Input data format		FREE		-124709.900	319144	176				
SUMMARY OF DATA				-124709.900	402224	91				
Number of missing data patterns		4		-124709.900	73576	213				
Number of y missing data patterns		4		-124709.900	863691	481				
Number of u missing data patterns		0		-124709.900	246261	38				
COVARIANCE COVERAGE OF DATA				-124709.900	790452	303				
Minimum covariance coverage value		0.100		-124709.900	252346	528				
PROPORTION OF DATA PRESENT FOR Y				-124709.900	407108	366				
Covariance Coverage				-124709.900	461687	833				
SLPNSSNC				-124709.900	798821	423				
WORRYNC				-124709.900	997222	229				
SOLC				-124709.900	782179	835				
SLPDURC				-124709.900	848969	173				
SLPNSSNC	1.000			-124709.900	739214	807				
WORRYNC	1.000	1.000		-124709.900	317868	740				
SOLC	0.976	0.976	0.976	-124709.900	415931	10				
SLPDURC	0.963	0.963	0.940	0.963	-124709.900	190339	102			
					-124709.900	781489	627			
					-124709.900	875667	367			
					-124709.900	685657	69			

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-124709.900	383986	159	-124709.900	484406	421
-124709.900	566739	575	-124709.900	97300	640
-124709.900	579995	183	-124709.900	966499	963
-124709.900	443917	60	-124709.900	584397	428
-124709.900	266038	672	-124709.900	746978	410
-124709.900	301428	885	-124709.900	259507	53
-124709.900	384199	882	-124709.900	876056	1000
-124709.900	50887	389	-124709.900	965994	396
-124709.900	724519	333	-124709.900	173191	422
-124709.900	853195	431	-124709.900	406935	958
-124709.900	165853	105	-124709.900	266340	683
-124709.900	464179	106	-124709.900	604861	800
-124709.900	699554	427	-124709.900	471398	74
-124709.900	207896	25	-124709.900	644297	340
-124709.900	637095	207	-124709.900	804104	566
-124709.900	679448	937	-124709.900	726035	191
-124709.900	87586	871	-124709.900	907810	795
-124709.900	529455	268	-124709.900	195873	6
-124709.900	779820	877	-124709.900	694303	282
-124709.900	769907	457	-124709.900	284384	600
-124709.900	535303	923	-124709.900	544048	87
-124709.900	574412	679	-124709.900	153394	429
-124709.900	3307	735	-124709.900	563584	657
-124709.900	276102	599	-124709.900	847088	750
-124709.900	486622	522	-124709.900	954354	840
-124709.900	349360	464	-124709.900	688839	273
-124709.900	603842	61	-124709.900	950604	172
-124709.900	575700	100	-124709.900	436806	883
-124709.900	232559	136	-124709.900	576726	280
-124709.900	481835	57	-124709.900	539389	544
-124709.900	544009	842	-124709.900	673496	218
-124709.900	928624	981	-124709.900	552272	654
-124709.900	547702	767	-124709.900	561664	392
-124709.900	59963	670	-124709.900	66276	217
-124709.900	246794	801	-124709.900	371246	101
-124709.900	291112	645	-124709.900	647617	725
-124709.900	728038	177	-124709.900	177936	726
-124709.900	357866	968	-124709.900	549244	756
-124709.900	197223	332	-124709.900	227563	63
-124709.900	741888	138	-124709.900	392418	28
-124709.900	192191	629	-124709.900	49293	707
-124709.900	780698	337	-124709.900	392407	221
-124709.900	891347	504	-124709.900	94610	884
-124709.900	349263	263	-124709.900	931558	898
-124709.900	576783	866	-124709.900	178181	753
-124709.900	81951	907	-124709.900	501995	791
-124709.900	879338	309	-124709.900	158612	581
-124709.900	499150	216	-124709.900	833196	715
-124709.900	204959	695	-124709.900	995648	936
-124709.900	3607	873	-124709.900	858804	534
-124709.900	607072	717	-124709.900	364676	27
-124709.900	321390	133	-124709.900	602032	648
-124709.900	84013	598	-124709.900	850112	922
-124709.900	814975	129	-124709.900	869564	869
-124709.900	640833	434	-124709.900	948615	140
-124709.900	375590	438	-124709.900	76974	16
-124709.900	872743	374	-124709.900	427006	839
-124709.900	140442	500	-124709.900	508482	446
-124709.900	315029	471	-124709.900	263268	165
-124709.900	467339	66	-124709.900	46437	153
-124709.900	59674	240	-124709.900	758647	951
-124709.900	473343	844	-124709.900	414284	158
-124709.900	526324	178	-124709.900	118421	139
-124709.900	424223	900	-124709.900	995875	547
-124709.900	11397	944	-124709.900	188498	258
-124709.900	404042	675	-124709.900	545108	667
-124709.900	391949	295	-124709.900	645052	910
-124709.900	937588	293	-124709.900	862607	468
-124709.900	648035	836	-124709.900	840031	276
-124709.900	496703	758	-124709.900	569833	85

Scripts & Outputs *“Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”*

-124709.900	863094	147	-124709.900	399848	220
-124709.900	506886	576	-124709.900	961454	665
-124709.900	638611	524	-124709.900	314757	345
-124709.900	271809	846	-124709.900	618760	489
-124709.900	244349	736	-124709.900	760878	249
-124709.900	597614	284	-124709.900	172913	557
-124709.900	579138	706	-124709.900	345726	461
-124709.900	455617	242	-124709.900	638977	643
-124709.900	852283	616	-124709.900	856536	144
-124709.900	193042	316	-124709.900	860772	174
-124709.900	370466	41	-124709.900	781190	219
-124709.900	23012	352	-124709.900	967902	52
-124709.900	407168	44			
-124709.900	178031	720			
-124709.900	789985	67			
-124709.900	509733	130			
-124709.900	339073	841			
-124709.900	804660	260			
-124709.900	947714	573			
-124709.900	692169	308			
-124709.900	922596	456			
-124709.900	53621	483			
-124709.900	505244	582			
-124709.900	915107	54			
-124709.900	638577	769			
-124709.900	715255	523			
-124709.900	475419	987			
-124709.900	125727	899			
-124709.900	954914	911			
-124709.900	442072	625			
-124709.900	851945	18			
-124709.900	978781	497			
-124709.900	432148	30			
-124709.900	297518	166			
-124709.900	177175	851			
-124709.900	376411	473			
-124709.900	650371	14			
-124709.900	215353	164			
-124709.900	930872	277			
-124709.900	478421	311			
-124709.900	766903	505			
-124709.900	301180	236			
-124709.900	848163	47			
-124709.900	326091	759			
-124709.900	700846	867			
-124709.900	39136	226			
-124709.900	306768	978			
-124709.900	466971	109			
-124709.900	292884	103			
-124709.900	926283	269			
-124709.900	629320	222			
-124709.900	871438	561			
-124709.900	574942	558			
-124709.900	311214	64			
-124709.900	440395	917			
-124709.900	843555	952			
-124709.900	626891	32			
-124709.900	403801	762			
-124709.900	677720	681			
-124709.900	717754	180			
-124709.900	655497	376			
-124709.900	387701	275			
-124709.900	391368	802			
-124709.900	798839	312			
-124709.900	996231	310			
-124709.900	636396	168			
-124709.900	193847	354			
-124709.900	393232	152			
-124709.900	61587	400			
-124709.900	98068	998			

THE BEST LOGLIKELIHOOD VALUE HAS BEEN REPLICATED. RERUN WITH AT LEAST TWICE THE RANDOM STARTS TO CHECK THAT THE BEST LOGLIKELIHOOD IS STILL OBTAINED AND REPLICATED.					
THE MODEL ESTIMATION TERMINATED NORMALLY					
MODEL FIT INFORMATION					
Number of Free Parameters		47			
Loglikelihood					
H0 Value		-124709.900			
H0 Scaling Correction Factor for MLR		1.1775			
Information Criteria					
Akaike (AIC)		249513.801			
Bayesian (BIC)		249900.745			
Sample-Size Adjusted BIC		249751.380			
(n* = (n + 2) / 24)					
FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THE ESTIMATED MODEL					
Latent Classes					
1		4456.90910	0.16031		
2		2657.87540	0.09560		
3		1910.44951	0.06872		
4		18776.76599	0.67537		
FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON ESTIMATED POSTERIOR PROBABILITIES					
Latent Classes					
1		4456.90910	0.16031		
2		2657.87540	0.09560		
3		1910.44951	0.06872		
4		18776.76599	0.67537		
FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THEIR MOST LIKELY LATENT CLASS MEMBERSHIP					
Class Counts and Proportions					

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Latent Classes					Variances				
					SLPNSSNC	0.460	0.003	150.050	0.000
					WORRYNC	0.245	0.004	55.585	0.000
1	4162	0.14970			SOLC	0.405	0.006	72.186	0.000
2	2502	0.08999			SLPDURC	0.797	0.008	98.354	0.000
3	1451	0.05219							
4	19687	0.70811			Latent Class 2				
CLASSIFICATION QUALITY					SLPNSSNC WITH				
Entropy		0.855			WORRYNC	0.003	0.007	0.398	0.691
Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)					SOLC	0.007	0.011	0.686	0.492
	1	2	3	4	SLPDURC	0.012	0.012	0.968	0.333
1	0.862	0.055	0.011	0.072	WORRYNC WITH				
2	0.071	0.912	0.016	0.001	SOLC	-0.042	0.008	-5.604	0.000
3	0.023	0.097	0.797	0.083	SLPDURC	-0.089	0.010	-8.966	0.000
4	0.034	0.000	0.034	0.932	SOLC WITH				
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					SLPDURC	0.029	0.013	2.263	0.024
	1	2	3	4	Means				
1	0.805	0.040	0.007	0.148	SLPNSSNC	1.148	0.015	75.059	0.000
2	0.087	0.859	0.053	0.001	WORRYNC	-1.962	0.022	-88.615	0.000
3	0.025	0.021	0.605	0.350	SOLC	1.874	0.022	85.791	0.000
4	0.016	0.000	0.006	0.978	SLPDURC	1.137	0.025	46.280	0.000
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					Variances				
	1	2	3	4	SLPNSSNC	0.460	0.003	150.050	0.000
1	1.691	-1.315	-3.000	0.000	WORRYNC	0.245	0.004	55.585	0.000
2	4.229	6.523	3.740	0.000	SOLC	0.405	0.006	72.186	0.000
3	-2.648	-2.830	0.548	0.000	SLPDURC	0.797	0.008	98.354	0.000
4	-4.119	-8.984	-5.021	0.000	Latent Class 3				
MODEL RESULTS					SLPNSSNC WITH				
	Estimate		Two-Tailed		WORRYNC	-0.019	0.010	-1.946	0.052
		S.E.	Est./S.E.	P-Value	SOLC	0.010	0.015	0.653	0.514
Latent Class 1					SLPDURC	-0.022	0.017	-1.290	0.197
SLPNSSNC WITH					WORRYNC WITH				
WORRYNC	-0.002	0.006	-0.280	0.780	SOLC	-0.075	0.012	-6.420	0.000
SOLC	0.081	0.007	10.882	0.000	SLPDURC	-0.102	0.014	-7.163	0.000
SLPDURC	0.030	0.009	3.201	0.001	SOLC WITH				
WORRYNC WITH					SLPDURC	0.049	0.020	2.429	0.015
SOLC	-0.007	0.005	-1.446	0.148	Means				
SLPDURC	-0.076	0.009	-8.952	0.000	SLPNSSNC	1.189	0.021	56.070	0.000
SOLC WITH					WORRYNC	0.017	0.049	0.341	0.733
SLPDURC	0.063	0.010	6.438	0.000	SOLC	1.520	0.047	32.307	0.000
Means					SLPDURC	0.303	0.047	6.470	0.000
SLPNSSNC	0.872	0.014	64.494	0.000	Variances				
WORRYNC	-1.107	0.021	-52.403	0.000	SLPNSSNC	0.460	0.003	150.050	0.000
SOLC	0.029	0.017	1.766	0.077	WORRYNC	0.245	0.004	55.585	0.000
SLPDURC	0.455	0.022	21.037	0.000	SOLC	0.405	0.006	72.186	0.000
					SLPDURC	0.797	0.008	98.354	0.000
					Latent Class 4				
					SLPNSSNC WITH				
					WORRYNC	-0.022	0.004	-6.262	0.000
					SOLC	0.075	0.004	18.458	0.000
					SLPDURC	0.022	0.005	4.056	0.000
					WORRYNC WITH				
					SOLC	-0.170	0.005	-37.049	0.000
					SLPDURC	-0.207	0.006	-33.958	0.000
					SOLC WITH				

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SLPDURC	0.217	0.007	33.011	0.000	2 Times the Loglikelihood Difference	2569.029
					Difference in the Number of Parameters	11
Means					Approximate P-Value	0.0000
SLPNSSNC	0.891	0.005	164.073	0.000	Successful Bootstrap Draws	250
WORRYNC	0.539	0.005	99.557	0.000		
SOLC	-0.420	0.008	-54.829	0.000		
SLPDURC	-0.294	0.007	-41.098	0.000		
Variiances					WARNING: OF THE 250 BOOTSTRAP DRAWS, 233 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 4-CLASS MODEL.	
SLPNSSNC	0.460	0.003	150.050	0.000	THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA.	
WORRYNC	0.245	0.004	55.585	0.000	INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.	
SOLC	0.405	0.006	72.186	0.000		
SLPDURC	0.797	0.008	98.354	0.000		
Categorical Latent Variables						
Means					PLOT INFORMATION	
C#1	-1.438	0.027	-54.013	0.000	The following plots are available:	
C#2	-1.955	0.033	-60.126	0.000	Histograms (sample values)	
C#3	-2.285	0.055	-41.873	0.000	Scatterplots (sample values)	
QUALITY OF NUMERICAL RESULTS					Sample means	
Condition Number for the Information Matrix					Estimated means, medians, modes, and percentiles	
0.201E-04					Sample and estimated means	
(ratio of smallest to largest eigenvalue)					Observed individual values	
TECHNICAL 11 OUTPUT					Estimated means and observed individual values	
Random Starts Specifications for the k-1 Class Analysis Model					Estimated overall and class-specific distributions	
Number of initial stage random starts			1000		SAVEDATA INFORMATION	
Number of final stage optimizations			250		Save file	
VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO TEST FOR 3 (H0) VERSUS 4 CLASSES					Step2_LPA4M5.dat	
H0 Loglikelihood Value			-125994.415		Order and format of variables	
2 Times the Loglikelihood Difference			2569.029		SLPNSSNC F10.3	
Difference in the Number of Parameters			11		WORRYNC F10.3	
Mean			-20.016		SOLC F10.3	
Standard Deviation			62.940		SLPDURC F10.3	
P-Value			0.0000		CPROB1 F10.3	
LO-MENDELL-RUBIN ADJUSTED LRT TEST					CPROB2 F10.3	
Value			2546.407		CPROB3 F10.3	
P-Value			0.0000		CPROB4 F10.3	
TECHNICAL 14 OUTPUT					C F10.3	
Random Starts Specifications for the k-1 Class Analysis Model					ID I6	
Number of initial stage random starts			1000		Save file format	
Number of final stage optimizations			250		9F10.3 I6	
Random Starts Specification for the k-1 Class Model for Generated Data					Save file record length	10000
Number of initial stage random starts			2		Save missing symbol	*
Number of final stage optimizations			1		DIAGRAM INFORMATION	
Random Starts Specification for the k Class Model for Generated Data					Mplus diagrams are currently not available for Mixture analysis.	
Number of initial stage random starts			50		No diagram output was produced.	
Number of final stage optimizations			10		Beginning Time: 13:46:02	
Number of bootstrap draws requested			250		Ending Time: 16:52:39	
PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 3 (H0) VERSUS 4 CLASSES					Elapsed Time: 03:06:37	
H0 Loglikelihood Value			-125994.415		MUTHEN & MUTHEN	
					3463 Stoner Ave.	
					Los Angeles, CA 90066	
					Tel: (310) 391-9971	
					Fax: (310) 391-8971	
					Web: www.StatModel.com	

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

Support: Support@StatModel.com

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Five-profile model

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Mplus VERSION 8.7
MUTHEN & MUTHEN
07/13/2023 8:14 PM

INPUT INSTRUCTIONS

TITLE: LPA 5 profile syntax: M5: Varying means, equal
variances, and varying covariances

DATA:
FILE IS mplus2.csv;
! Specifies file location for data file. Make sure data is in
format appropriate for Mplus
! per Mplus manual. This data file is in individual format
(one row of data per participan
VARIABLE:
NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol
slpDur gender genderN yrgrp yrgrpN SCI_02C
slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be
listed here
IDVARIABLE IS ID;
CLASSES = c (5);
! This is where you instruct Mplus on how many
classes/profiles are being estimated. Initi
! model contains only one class/profile, thus it would be
CLASSES = c (1). Above specifies
!profiles, and for each further iterative models the number
in parentheses increases by on
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(9999);
! Used to communicate how missing data is coded in data
file. Here shown with a "." which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 2000 500;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifie
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!Steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will sto
!before reaching 500 iterations.
!These values can be increased ... see “Four-Profile Final
Model with Covariate Analysis
!Syntax” for an example.
LRTSTARTS = 4 2 100 20;
LRTBOOTSTRAP = 500;
!The above start values are for the defaults for the LRT
statistic being run to compare th

!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statemen
!specifies the number of bootstrap draws to inform Mplus’
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances
!profiles to be equal.

%c#1%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#2%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#3%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#4%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#5%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;

```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

<pre>solC WITH slpDurC; OUTPUT: TECH11 TECH14; ! TECH1 provides parameter specifications and starting values for the analysis ! TECH8 provides optimization history for this analysis type !TECH11 provides LRT results !TECH14 provides bootstrapped LRT test PLOT: SERIES=slpnssNC worryNC solC slpDurC(*); TYPE=PLOT3; SAVEDATA: FILE IS Step2_LPA5M5.dat; ! Tells Mplus where to save the output files from the analysis SAVE = CPROBABILITIES; ! The above command lines are to save the most likely profile membership for each particip ! and the posterior probabilities for their membership in each latent profile. *** WARNING in MODEL command All variables are uncorrelated with all other variables within class. Check that this is what is intended. LPA 5 profile syntax: M5: Varying means, equal variances, and varying covariances SUMMARY OF ANALYSIS Number of groups 1 Number of observations 27802 Number of dependent variables 4 Number of independent variables 0 Number of continuous latent variables 0 Number of categorical latent variables 1 Observed dependent variables Continuous SLPNSSNC WORRYNC SOLC SLPDURC Categorical latent variables C Variables with special functions ID variable ID Estimator MLR Information matrix OBSERVED Optimization Specifications for the Quasi-Newton Algorithm for Continuous Outcomes Maximum number of iterations 100 Convergence criterion 0.100D-05 Optimization Specifications for the EM Algorithm Maximum number of iterations 500 Convergence criteria Loglikelihood change 0.100D-06 Relative loglikelihood change 0.100D-06 Derivative 0.100D-05 Optimization Specifications for the M step of the EM Algorithm for Categorical Latent variables Number of M step iterations 1 M step convergence criterion 0.100D-05 Basis for M step termination ITERATION</pre>	<pre>Optimization Specifications for the M step of the EM Algorithm for Censored, Binary or Ordered Categorical (Ordinal), Unordered Categorical (Nominal) and Count Outcomes Number of M step iterations 1 M step convergence criterion 0.100D-05 Basis for M step termination ITERATION Maximum value for logit thresholds 15 Minimum value for logit thresholds -15 Minimum expected cell size for chi-square 0.100D- 01 Maximum number of iterations for H1 2000 Convergence criterion for H1 0.100D-03 Optimization algorithm EMA Random Starts Specifications Number of initial stage random starts 2000 Number of final stage optimizations 500 Number of initial stage iterations 500 Initial stage convergence criterion 0.100D+01 Random starts scale 0.500D+01 Random seed for generating random starts 0 Input data file(s) mplus2.csv Input data format FREE SUMMARY OF DATA Number of missing data patterns 4 Number of y missing data patterns 4 Number of u missing data patterns 0 COVARIANCE COVERAGE OF DATA Minimum covariance coverage value 0.100 PROPORTION OF DATA PRESENT FOR Y Covariance Coverage SLPNSSNC WORRYNC SOLC SLPDURC ----- SLPNSSNC 1.000 WORRYNC 1.000 1.000 SOLC 0.976 0.976 0.976 SLPDURC 0.963 0.963 0.940 0.963 UNIVARIATE SAMPLE STATISTICS UNIVARIATE HIGHER-ORDER MOMENT DESCRIPTIVE STATISTICS Variable/ Mean/ Skewness/ Minimum/ % with Percentiles Sample Size Variance Kurtosis Maximum Min/Max 20%/60% 40%/80% Median SLPNSSNC 0.933 0.086 0.000 27.10% 0.000 1.000 1.000 27802.000 0.471 -0.878 2.000 20.40% 1.000 2.000 WORRYNC 0.000 -1.036 -2.825 1.71% -0.928 0.020 0.495 27802.000 1.000 0.165 0.969 28.55% 0.495 0.969 SOLC 0.000 0.967 -1.054 31.63% 1.054 -0.072 -0.072 27123.000 1.000 0.467 2.875 2.77% -0.072 0.910</pre>
---	--

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

SLPDURC	0.000	0.568	-1.205	25.70%	-124083.504	300844	1405	
-1.205	-0.292	-0.292			-124083.504	301180	236	
	26763.000	1.000	-0.470	2.445	3.44%	-124083.504	583281	1907
-0.292	0.620				-124083.504	256197	1923	
					-124083.504	908674	1748	
RANDOM STARTS RESULTS RANKED FROM THE BEST TO THE WORST LOGLIKELIHOOD VALUES					-124083.504	793487	595	
					-124083.504	992389	77	
					-124083.504	150531	154	
Unperturbed starting value run did not converge in the initial stage optimizations.					-124083.504	929361	1971	
					-124083.504	964999	1522	
					-124083.504	325724	1043	
					-124083.504	876943	650	
1760 perturbed starting value run(s) did not converge in the initial stage optimizations.					-124083.504	568279	1580	
					-124083.504	306768	978	
					-124083.504	547702	767	
					-124083.504	549244	756	
Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:					-124083.504	526742	1546	
					-124083.504	922042	492	
					-124083.504	848163	47	
-124083.504	285763	1486			-124083.504	105435	265	
-124083.504	530559	1845			-124083.504	393232	152	
-124083.504	464810	1424			-124083.504	501995	791	
-124083.504	453260	1106			-124083.504	222513	1088	
-124083.504	627151	1601			-124083.504	59963	670	
-124083.504	317868	740			-124083.504	919217	1501	
-124083.504	628997	1188			-124083.504	238317	812	
-124083.504	393453	1355			-124083.504	354208	196	
-124083.504	153942	31			-124083.504	290210	1620	
-124083.504	884619	1441			-124083.504	855963	1408	
-124083.504	848331	137			-124083.504	990504	827	
-124083.504	929823	1205			-124083.504	81951	907	
-124083.504	922596	456			-124083.504	251680	794	
-124083.504	716616	1159			-124083.504	51375	148	
-124083.504	718876	1104			-124083.504	856347	1927	
-124083.504	539389	544			-124083.504	760850	739	
-124083.504	268664	1784			-124083.504	981480	1025	
-124083.504	225932	1285			-124096.895	702409	1227	
-124083.504	712531	631			-124096.895	117891	1818	
-124083.504	626208	698			-124096.895	669634	335	
-124083.504	210870	383			-124096.895	131063	843	
-124083.504	185331	1162			-124096.895	126922	1124	
-124083.504	778331	1280			-124096.895	331689	1872	
-124083.504	881886	608			-124096.895	303429	1016	
-124083.504	930872	277			-124096.895	765589	1140	
-124083.504	685657	69			-124096.895	165853	105	
-124083.504	526265	1962			-124096.895	570681	777	
-124083.504	760599	832			-124096.895	84927	1511	
-124083.504	175820	1040			-124096.895	349263	263	
-124083.504	696830	668			-124096.895	928085	1946	
-124083.504	879211	453			-124096.895	956446	1081	
-124083.504	676560	1250			-124096.895	164643	1941	
-124083.504	507218	613			-124096.895	358115	1724	
-124083.504	327927	908			-124096.895	197904	1782	
-124083.504	927045	1994			-124096.895	769907	457	
-124083.504	273669	1118			-124096.895	341041	34	
-124083.504	188498	258			-124096.895	765372	1123	
-124083.504	291112	645			-124096.895	159216	1423	
-124083.504	276243	1768			-124096.895	441118	1091	
-124083.504	782821	272			-124096.895	63231	935	
-124083.504	266349	1989			-124096.895	964389	1223	
-124083.504	645052	910			-124096.895	813351	1507	
-124083.504	399071	1007			-124096.895	95624	1605	
-124083.504	547354	1357			-124096.895	674786	1496	
-124083.504	165822	1803			-124096.895	879338	309	
-124083.504	61710	1327			-124096.895	629320	222	
-124083.504	16096	1147			-124096.895	662718	460	
-124083.504	967902	52			-124096.895	644297	340	
-124083.504	384859	1530			-124096.895	558208	1403	
-124083.504	424280	1245			-124096.895	53089	1660	
-124083.504	680987	1926			-124096.895	471398	74	

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-124096.895	933578	506	Bayesian (BIC)	248760.513				
-124096.895	814975	129	Sample-Size Adjusted BIC	248576.190				
-124096.895	518828	432	($n^* = (n + 2) / 24$)					
-124096.895	166220	1502						
-124096.895	96617	955						
-124096.895	496710	386	FINAL CLASS COUNTS AND PROPORTIONS FOR THE					
-124096.895	510491	1275	LATENT CLASSES					
-124096.895	752847	1109	BASED ON THE ESTIMATED MODEL					
-124096.895	848356	602	Latent					
-124096.895	654377	1684	Classes					
-124096.895	271809	846						
-124096.895	699748	1120	1	1300.32515 0.04677				
-124096.895	678813	1881	2	4426.85090 0.15923				
-124096.895	621542	375	3	1645.56536 0.05919				
-124096.895	358688	1573	4	2323.29222 0.08357				
-124096.895	937885	426	5	18105.96637 0.65125				
-124096.895	435506	988						
-124096.895	521770	945	FINAL CLASS COUNTS AND PROPORTIONS FOR THE					
-124096.895	188490	1460	LATENT CLASSES					
-124096.895	493775	1402	BASED ON ESTIMATED POSTERIOR PROBABILITIES					
-124096.895	910119	1886	Latent					
-124096.895	246261	38	Classes					
-124096.895	792993	859						
-124096.895	51011	1111	1	1300.32515 0.04677				
-124096.895	153053	378	2	4426.85090 0.15923				
-124096.895	970689	266	3	1645.56536 0.05919				
-124096.895	528938	1898	4	2323.29222 0.08357				
-124709.900	945711	1419	5	18105.96637 0.65125				
-124709.900	996231	310						
-124709.900	611687	1933	FINAL CLASS COUNTS AND PROPORTIONS FOR THE					
-124709.900	569833	85	LATENT CLASSES					
-124709.900	469158	1145	BASED ON THEIR MOST LIKELY LATENT CLASS					
-124709.900	451258	848	MEMBERSHIP					
-124709.900	137305	379	Class Counts and Proportions					
-124709.900	531546	702	Latent					
-124709.900	207299	1039	Classes					
-124709.900	938409	1144						
-124709.900	135076	1692	1	1037 0.03730				
-125293.398	842091	1373	2	4825 0.17355				
-125293.398	677720	681	3	1250 0.04496				
-125394.935	232559	136	4	2335 0.08399				
-125394.935	881268	1299	5	18355 0.66020				
-125994.415	489927	520						
-125994.415	181178	1069						
-126943.690	91231	727						
334 perturbed starting value run(s) did not converge or were rejected in the third stage.			CLASSIFICATION QUALITY					
			Entropy	0.860				
THE BEST LOGLIKELIHOOD VALUE HAS BEEN REPLICATED. RERUN WITH AT LEAST TWICE THE RANDOM STARTS TO CHECK THAT THE BEST LOGLIKELIHOOD IS STILL OBTAINED AND REPLICATED.			Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)					
			1	2	3	4	5	
THE MODEL ESTIMATION TERMINATED NORMALLY			1	0.782	0.142	0.001	0.075	0.000
MODEL FIT INFORMATION			2	0.061	0.778	0.052	0.023	0.085
Number of Free Parameters	58		3	0.002	0.036	0.905	0.043	0.014
Loglikelihood			4	0.081	0.019	0.008	0.891	0.000
			5	0.000	0.024	0.013	0.000	0.963
			Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)					
			1	2	3	4	5	
Information Criteria			1	0.624	0.228	0.002	0.146	0.000
Akaike (AIC)	248283.007		2	0.033	0.848	0.010	0.010	0.098

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

3	0.000	0.152	0.687	0.012	0.148	SOLC	0.414	0.004	99.215	0.000
4	0.033	0.048	0.023	0.896	0.000	SLPDURC	0.752	0.009	82.198	0.000
5	0.000	0.023	0.001	0.000	0.976					
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)						Latent Class 3				
	1	2	3	4	5	SLPNSSNC WITH WORRYNC	-0.002	0.007	-0.314	0.753
						SOLC	-0.015	0.019	-0.760	0.447
						SLPDURC	-0.030	0.016	-1.880	0.060
1	9.660	8.655	3.760	8.210	0.000	WORRYNC WITH SOLC	-0.004	0.008	-0.475	0.635
2	-1.078	2.158	-2.261	-2.280	0.000	SLPDURC	-0.076	0.010	-7.544	0.000
3	-5.926	0.029	1.535	-2.541	0.000	SOLC WITH SLPDURC	0.012	0.020	0.592	0.554
4	9.727	10.084	9.366	13.018	0.000					
5	-13.507	-3.761	-6.925	-12.691	0.000					
MODEL RESULTS						Means				
	Estimate	Two-Tailed S.E. Est./S.E.		P-Value		SLPNSSNC	1.209	0.022	54.036	0.000
Latent Class 1						WORRYNC	-0.224	0.037	-5.997	0.000
SLPNSSNC WITH WORRYNC		0.028	0.014	2.097	0.036	SOLC	1.723	0.027	63.438	0.000
SOLC		0.054	0.019	2.863	0.004	SLPDURC	0.439	0.044	9.899	0.000
SLPDURC		-0.004	0.021	-0.173	0.863	Variances				
WORRYNC WITH SOLC		0.018	0.023	0.760	0.447	SLPNSSNC	0.460	0.003	150.607	0.000
SLPDURC		0.011	0.017	0.603	0.546	WORRYNC	0.183	0.003	66.958	0.000
SOLC WITH SLPDURC		0.029	0.023	1.279	0.201	SOLC	0.414	0.004	99.215	0.000
SLPDURC						SLPDURC	0.752	0.009	82.198	0.000
Means						Latent Class 4				
SLPNSSNC		0.941	0.041	23.047	0.000	SLPNSSNC WITH WORRYNC	0.004	0.006	0.633	0.527
WORRYNC		-1.812	0.077	-23.556	0.000	SOLC	-0.007	0.014	-0.517	0.605
SOLC		0.058	0.078	0.741	0.459	SLPDURC	0.013	0.013	1.000	0.317
SLPDURC		0.950	0.069	13.849	0.000	WORRYNC WITH SOLC	-0.050	0.007	-6.747	0.000
Variances						SLPDURC	-0.072	0.008	-9.171	0.000
SLPNSSNC		0.460	0.003	150.607	0.000	SOLC WITH SLPDURC	0.024	0.014	1.765	0.078
WORRYNC		0.183	0.003	66.958	0.000	Means				
SOLC		0.414	0.004	99.215	0.000	SLPNSSNC	1.167	0.017	69.498	0.000
SLPDURC		0.752	0.009	82.198	0.000	WORRYNC	-2.030	0.021	-97.318	0.000
Latent Class 2						SOLC	1.944	0.024	81.105	0.000
SLPNSSNC WITH WORRYNC		0.009	0.009	1.072	0.284	SLPDURC	1.173	0.026	45.965	0.000
SOLC		0.088	0.008	10.716	0.000	Variances				
SLPDURC		0.029	0.010	2.814	0.005	SLPNSSNC	0.460	0.003	150.607	0.000
WORRYNC WITH SOLC		-0.074	0.009	-8.090	0.000	WORRYNC	0.183	0.003	66.958	0.000
SLPDURC		-0.007	0.014	-0.457	0.648	SOLC	0.414	0.004	99.215	0.000
SOLC WITH SLPDURC		0.096	0.013	7.505	0.000	SLPDURC	0.752	0.009	82.198	0.000
Means						Latent Class 5				
SLPNSSNC		0.875	0.015	58.846	0.000	SLPNSSNC WITH WORRYNC	-0.016	0.003	-5.075	0.000
WORRYNC		-0.749	0.035	-21.694	0.000	SOLC	0.074	0.004	19.537	0.000
SOLC		0.032	0.029	1.135	0.256	SLPDURC	0.017	0.005	3.071	0.002
SLPDURC		0.332	0.033	9.996	0.000	WORRYNC WITH SOLC	-0.136	0.003	-43.443	0.000
Variances						SLPDURC	-0.150	0.005	-32.481	0.000
SLPNSSNC		0.460	0.003	150.607	0.000	SOLC WITH SLPDURC	0.195	0.006	33.001	0.000
WORRYNC		0.183	0.003	66.958	0.000	Means				

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

SLPNSSNC	0.892	0.005	163.537	0.000	Difference in the Number of Parameters	11
WORRYNC	0.594	0.005	123.078	0.000	Approximate P-Value	0.0000
SOLC	-0.411	0.006	-74.068	0.000	Successful Bootstrap Draws	500
SLPDURC	-0.334	0.007	-45.482	0.000		
Variances					WARNING: OF THE 500 BOOTSTRAP DRAWS, 459 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 5-CLASS MODEL.	
SLPNSSNC	0.460	0.003	150.607	0.000	THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA.	
WORRYNC	0.183	0.003	66.958	0.000	INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.	
SOLC	0.414	0.004	99.215	0.000		
SLPDURC	0.752	0.009	82.198	0.000		
Categorical Latent Variables						
Means						
C#1	-2.634	0.144	-18.330	0.000	PLOT INFORMATION	
C#2	-1.409	0.032	-44.014	0.000	The following plots are available:	
C#3	-2.398	0.042	-56.710	0.000	Histograms (sample values)	
C#4	-2.053	0.032	-63.375	0.000	Scatterplots (sample values)	
QUALITY OF NUMERICAL RESULTS						
Condition Number for the Information Matrix						
0.996E-05						
(ratio of smallest to largest eigenvalue)						
TECHNICAL 11 OUTPUT						
Random Starts Specifications for the k-1 Class Analysis						
Model						
Number of initial stage random starts 2000						
Number of final stage optimizations 500						
VUONG-LO-MENDELLE-RUBIN LIKELIHOOD RATIO						
TEST FOR 4 (H0) VERSUS 5 CLASSES						
H0 Loglikelihood Value -124709.900						
2 Times the Loglikelihood Difference 1252.794						
Difference in the Number of Parameters 11						
Mean 26.542						
Standard Deviation 38.528						
P-Value 0.0000						
LO-MENDELLE-RUBIN ADJUSTED LRT TEST						
Value 1241.762						
P-Value 0.0000						
TECHNICAL 14 OUTPUT						
Random Starts Specifications for the k-1 Class Analysis						
Model						
Number of initial stage random starts 2000						
Number of final stage optimizations 500						
Random Starts Specification for the k-1 Class Model for Generated Data						
Number of initial stage random starts 4						
Number of final stage optimizations 2						
Random Starts Specification for the k Class Model for Generated Data						
Number of initial stage random starts 100						
Number of final stage optimizations 20						
Number of bootstrap draws requested 500						
PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO						
TEST FOR 4 (H0) VERSUS 5 CLASSES						
H0 Loglikelihood Value -124709.900						
2 Times the Loglikelihood Difference 1252.794						
SAVEDATA INFORMATION						
Save file						
Step2_LPA5M5.dat						
Order and format of variables						
SLPNSSNC F10.3						
WORRYNC F10.3						
SOLC F10.3						
SLPDURC F10.3						
CPROB1 F10.3						
CPROB2 F10.3						
CPROB3 F10.3						
CPROB4 F10.3						
CPROB5 F10.3						
C F10.3						
ID I6						
Save file format						
10F10.3 I6						
Save file record length 10000						
Save missing symbol *						
DIAGRAM INFORMATION						
Mplus diagrams are currently not available for Mixture analysis.						
No diagram output was produced.						
Beginning Time: 20:14:35						
Ending Time: 20:57:03						
Elapsed Time: 24:42:28						
MUTHEN & MUTHEN						
3463 Stoner Ave.						
Los Angeles, CA 90066						
Tel: (310) 391-9971						
Fax: (310) 391-8971						
Web: www.StatModel.com						

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

Support: Support@StatModel.com

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Six-profile model

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Mplus VERSION 8.7
MUTHEN & MUTHEN
07/15/2023 1:26 PM

INPUT INSTRUCTIONS

TITLE: LPA 6 profile syntax: M5: Varying means, equal
variances, and varying covariances

DATA:
FILE IS mplus2.csv;
! Specifies file location for data file. Make sure data is in
format appropriate for Mplus
! per Mplus manual. This data file is in individual format
(one row of data per participan
VARIABLE:
NAMES ARE ID SCI_02 SCI_02B slpnssN worryN sol
slpDur gender genderN yrgrp yrgrpN SCI_02C
slpnssNC worryNC solC slpDurC;

! All variables included in data file should be named here.
USEVARIABLES ARE slpnssNC worryNC solC slpDurC;
! Only variables intended for use in the analysis should be
listed here
IDVARIABLE IS ID;
CLASSES = c (6);
! This is where you instruct Mplus on how many
classes/profiles are being estimated. Initi
! model contains only one class/profile, thus it would be
CLASSES = c (1). Above specifies
!profiles, and for each further iterative models the number
in parentheses increases by on
!three profiles/classes would be c (3), and so on.
MISSING ARE ALL(9999);
! Used to communicate how missing data is coded in data
file. Here shown with a "." which
! all that is included in each cell with missing data in the
data file
ANALYSIS:
TYPE = MIXTURE;
! LPA is a version of mixture modeling, and this instructs
Mplus to analyze in this way
ESTIMATOR = MLR;
!FIML robust to non-normal data
STARTS = 8000 2000;
STITERATIONS = 500;
! Default number of starts for each step of the ML
estimation. First STARTS value specifie
!number of unique start values to start with, the 250
represents the 250 best unique start
!carrying forward to completion. The STITERATIONS
specifies the number of ML iteration
!Steps for those 250 selected start values to go through to
be able to converge. This is a
!maximum number of iteration; if a model converges in less
than 500 iterations it will sto
!before reaching 500 iterations.
!These values can be increased ... see "Four-Profile Final
Model with Covariate Analysis
!Syntax" for an example.
LRTSTARTS = 16 8 400 80;
LRTBOOTSTRAP = 500;
!The above start values are for the defaults for the LRT
statistic being run to compare th

!model fit with the model fit of a model with one less class
(k-1). The BOOTSTRAP statemen
!specifies the number of bootstrap draws to inform Mplus'
bootstrapped LRT results.
MODEL:
!For a default Mplus model the LPA model does not need
to be specified. However, it can be
!The model can also be modified from the Mplus default of
estimating the indicator means
!(uniquely across profiles) and variances (constrained
across profiles), as well as the la
!profile mean.
%OVERALL%
[slpnssNC worryNC solC slpDurC]; !estimates the
indicators means for each profile. Without
!the means are freely estimated in each profile, not
constrained.
slpnssNC worryNC solC slpDurC; !Label Var1-Var5
constrains the estimates of the variances
!profiles to be equal.

%c#1%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#2%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#3%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#4%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
solC WITH slpDurC;

%c#5%

[slpnssNC worryNC solC slpDurC];

!covariances
slpnssNC WITH worryNC solC slpDurC;
worryNC WITH solC slpDurC;
```

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

<pre>solC WITH slpDurC; %c#6% [slpnssNC worryNC solC slpDurC]; !covariances slpnssNC WITH worryNC solC slpDurC; worryNC WITH solC slpDurC; solC WITH slpDurC; OUTPUT: TECH11 TECH14; ! TECH1 provides parameter specifications and starting values for the analysis ! TECH8 provides optimization history for this analysis type ! TECH11 provides LRT results ! TECH14 provides bootstrapped LRT test PLOT: SERIES=slpnssNC worryNC solC slpDurC(*); TYPE=PLOT3; SAVEDATA: FILE IS Step2_LPA6M5.dat; ! Tells Mplus where to save the output files from the analysis SAVE = CPROBABILITIES; ! The above command lines are to save the most likely profile membership for each particip ! and the posterior probabilities for their membership in each latent profile. *** WARNING in MODEL command All variables are uncorrelated with all other variables within class. Check that this is what is intended. LPA 6 profile syntax: M5: Varying means, equal variances, and varying covariances SUMMARY OF ANALYSIS Number of groups 1 Number of observations 27802 Number of dependent variables 4 Number of independent variables 0 Number of continuous latent variables 0 Number of categorical latent variables 1 Observed dependent variables Continuous SLPNSSNC WORRYNC SOLC SLPDURC Categorical latent variables C Variables with special functions ID variable ID Estimator MLR Information matrix OBSERVED Optimization Specifications for the Quasi-Newton Algorithm for Continuous Outcomes Maximum number of iterations 100 Convergence criterion 0.100D-05 Optimization Specifications for the EM Algorithm Maximum number of iterations 500 Convergence criteria</pre>	<pre>Loglikelihood change 0.100D-06 Relative loglikelihood change 0.100D-06 Derivative 0.100D-05 Optimization Specifications for the M step of the EM Algorithm for Categorical Latent variables Number of M step iterations 1 M step convergence criterion 0.100D-05 Basis for M step termination ITERATION Optimization Specifications for the M step of the EM Algorithm for Censored, Binary or Ordered Categorical (Ordinal), Unordered Categorical (Nominal) and Count Outcomes Number of M step iterations 1 M step convergence criterion 0.100D-05 Basis for M step termination ITERATION Maximum value for logit thresholds 15 Minimum value for logit thresholds -15 Minimum expected cell size for chi-square 0.100D- 01 Maximum number of iterations for H1 2000 Convergence criterion for H1 0.100D-03 Optimization algorithm EMA Random Starts Specifications Number of initial stage random starts 8000 Number of final stage optimizations 2000 Number of initial stage iterations 500 Initial stage convergence criterion 0.100D+01 Random starts scale 0.500D+01 Random seed for generating random starts 0 Input data file(s) mplus2.csv Input data format FREE SUMMARY OF DATA Number of missing data patterns 4 Number of y missing data patterns 4 Number of u missing data patterns 0 COVARIANCE COVERAGE OF DATA Minimum covariance coverage value 0.100 PROPORTION OF DATA PRESENT FOR Y Covariance Coverage SLPNSSNC WORRYNC SOLC SLPDURC SLPNSSNC 1.000 WORRYNC 1.000 1.000 SOLC 0.976 0.976 0.976 SLPDURC 0.963 0.963 0.940 0.963 UNIVARIATE SAMPLE STATISTICS UNIVARIATE HIGHER-ORDER MOMENT DESCRIPTIVE STATISTICS Variable/ Mean/ Skewness/ Minimum/ % with Percentiles Sample Size Variance Kurtosis Maximum Min/Max 20%/60% 40%/80% Median SLPNSSNC 0.933 0.086 0.000 27.10% 0.000 1.000 1.000</pre>
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Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

27802.000	0.471	-0.878	2.000	20.40%	-123380.204	279725	5288
1.000	2.000				-123380.204	548301	5906
WORRYNC	0.000	-1.036	-2.825	1.71%	-123380.204	270257	5176
-0.928	0.020	0.495			-123380.204	682047	2666
27802.000	1.000	0.165	0.969	28.55%	-123380.204	9427	4643
0.495	0.969				-123380.204	880807	6726
SOLC	0.000	0.967	-1.054	31.63%	-123380.204	550354	4790
1.054	-0.072	-0.072			-123380.204	120368	6622
27123.000	1.000	0.467	2.875	2.77%	-123380.204	797872	7655
-0.072	0.910				-123380.204	463894	1733
SLPDURC	0.000	0.568	-1.205	25.70%	-123380.204	876943	650
-1.205	-0.292	-0.292			-123380.204	400299	4782
26763.000	1.000	-0.470	2.445	3.44%	-123380.204	854941	6096
-0.292	0.620				-123380.204	217990	1077
					-123380.204	907407	2624
RANDOM STARTS RESULTS RANKED FROM THE BEST TO THE WORST LOGLIKELIHOOD VALUES					-123380.204	54085	5617
Unperturbed starting value run did not converge in the initial stage optimizations.					-123380.204	755611	3182
					-123380.204	551340	766
7502 perturbed starting value run(s) did not converge in the initial stage optimizations.					-123380.204	542577	1004
					-123380.204	766178	1937
					-123380.204	512048	3172
					-123380.204	678595	5688
					-123380.204	153053	378
					-123380.204	444228	860
					-123380.204	129163	4601
					-123380.204	139941	3271
Final stage loglikelihood values at local maxima, seeds, and initial stage start numbers:					-123380.204	563452	7834
					-123380.204	846789	7904
					-123380.204	715466	5857
-105840.802	102504	7215			-123380.204	728282	5901
-123380.204	347110	3761			-123380.204	430691	7901
-123380.204	366872	1169			-123380.204	8370	4917
-123380.204	630834	2964			-123380.204	776607	7866
-123380.204	45556	3294			-123380.204	254297	4551
-123380.204	530305	6477			-123380.204	420298	1222
-123380.204	209545	7123			-123380.204	407520	2854
-123380.204	689589	3947			-123380.204	294397	7351
-123380.204	419979	5094			-123380.204	352277	42
-123380.204	632565	4724			-123380.204	62571	3923
-123380.204	460630	3542			-123380.204	937651	5704
-123380.204	470921	2993			-123380.204	280104	1875
-123380.204	929499	3974			-123380.204	848163	47
-123380.204	166700	7162			-123380.204	953815	1347
-123380.204	231366	1137			-123380.204	510814	3219
-123380.204	640797	4654			-123380.204	988209	7713
-123380.204	474013	7485			-123380.204	530559	1845
-123380.204	977507	1716			-123380.204	354693	2038
-123380.204	110260	2535			-123380.204	736635	7286
-123380.204	231986	7210			-123380.204	666720	671
-123380.204	81598	7240			-123380.204	970838	3615
-123380.204	855715	6992			-123380.204	357565	7348
-123380.204	607148	6862			-123380.204	697272	2010
-123380.204	29275	6028			-123380.204	480926	1824
-123380.204	863480	5377			-123380.204	748083	4737
-123380.204	25584	7402			-123380.204	413363	2884
-123380.204	466035	6850			-123380.204	751592	3458
-123380.204	933578	506			-123380.204	876447	6025
-123380.204	690608	1800			-123380.204	37806	4535
-123380.204	848969	173			-123380.204	634782	979
-123380.204	220633	5604			-123380.204	485948	3188
-123380.204	567252	1843			-123380.204	908524	6156
-123380.204	978671	5259			-123380.204	314551	3955
-123380.204	789564	2275			-123380.204	619187	7233
-123380.204	719104	6684			-123380.204	937476	7543
-123380.204	12814	4104			-123380.204	760850	739
-123380.204	899062	3493			-123380.204	717754	180
-123380.204	165822	1803			-123380.204	390788	3191
-123380.204	42278	1305			-123380.204	564582	7832
-123380.204	88178	5619			-123380.204	938366	6817
-123380.204	473869	3707			-123380.204	356418	7958

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-123380.204 787782	3419	-123380.204 692399	3763
-123380.204 634094	7507	-123380.204 463203	5945
-123380.204 829063	2276	-123380.204 120100	5289
-123380.204 739458	1963	-123380.204 832850	2779
-123380.204 362529	2736	-123380.204 173191	422
-123380.204 321363	1095	-123380.204 885436	4377
-123380.204 337438	4320	-123380.204 626490	4476
-123380.204 734978	5042	-123380.204 749808	3344
-123380.204 286505	2416	-123380.204 140694	5735
-123380.204 794411	5790	-123380.204 323788	3307
-123380.204 721907	4479	-123380.204 573976	1217
-123380.204 76422	5258	-123380.204 42579	4794
-123380.204 310591	4901	-123380.204 882244	6464
-123380.204 925741	1567	-123380.204 703618	7115
-123380.204 450584	3772	-123380.204 566784	6618
-123380.204 244349	736	-123380.204 469648	6565
-123380.204 513053	5961	-123380.204 670998	1876
-123380.204 900041	3785	-123380.204 655088	7777
-123380.204 821111	6824	-123380.204 412313	6615
-123380.204 402776	7710	-123380.204 477242	4717
-123380.204 687000	5863	-123380.204 627224	4579
-123380.204 891347	504	-123380.204 231581	1048
-123380.204 76204	2003	-123380.204 824956	607
-123380.204 701586	5384	-123380.204 995881	4723
-123380.204 7510	2482	-123380.204 177175	851
-123380.204 962359	4118	-123380.204 529627	1542
-123380.204 187001	4588	-123380.204 900432	2808
-123380.204 81648	5344	-123380.204 470570	3598
-123380.204 252552	2489	-123380.204 880440	5602
-123380.204 476538	5063	-123380.204 963975	6544
-123380.204 712066	2494	-123380.204 976112	4060
-123380.204 950604	172	-123380.204 417391	7371
-123380.204 475953	3687	-123380.204 603765	1006
-123380.204 861413	1154	-123380.204 863378	2556
-123380.204 18840	3536	-123380.204 738142	1970
-123380.204 24045	2066	-123380.204 633988	6030
-123380.204 374630	1869	-123380.204 828340	1559
-123380.204 145690	7951	-123380.204 804616	868
-123380.204 61710	1327	-123380.204 399071	1007
-123380.204 25294	2192	-123380.204 683728	5363
-123380.204 803696	1340	-123380.204 681079	3005
-123380.204 43897	2230	-123380.204 838356	5513
-123380.204 589839	4722	-123380.204 660633	5171
-123380.204 763802	4958	-123380.204 659469	5608
-123380.204 369266	7511	-123380.204 287220	4395
-123380.204 344416	3290	-123380.204 606153	5028
-123380.204 603416	6452	-123380.204 335396	5902
-123380.204 955504	7773	-123380.204 973369	202
-123380.204 573791	6121	-123380.204 526836	6690
-123380.204 164467	2047	-123380.204 576306	3678
-123380.204 501995	791	-123380.204 695314	6891
-123380.204 307045	7189	-123380.204 697866	1018
-123380.204 691544	2183	-123380.204 927045	1994
-123380.204 757590	2804	-123380.204 669634	335
-123380.204 584751	4113	-123380.204 159216	1423
-123380.204 643858	5665	-123380.204 802250	7675
-123380.204 439675	7827	-123380.204 200114	6733
-123380.204 17896	592	-123380.204 860732	2774
-123380.204 166096	4361	-123380.204 587497	5903
-123380.204 389980	7164	-123380.204 978781	497
-123380.204 438631	7526	-123380.204 427273	2524
-123380.204 486133	7364	-123380.204 869821	6237
-123380.204 602960	2226	-123380.204 955439	7882
-123380.204 785023	3454	-123380.204 498982	4136
-123380.204 262076	2015	-123380.204 10606	4208
-123380.204 438144	271	-123380.204 581912	3149
-123380.204 801057	6339	-123380.204 769417	4111
-123380.204 488565	5839	-123380.204 561782	6822
-123380.204 760380	3885	-123380.204 234123	7681
-123380.204 887787	5625	-123380.204 914462	3094

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-123380.204	950970	2963	-123380.204	437885	7147
-123380.204	234995	4728	-123380.204	953205	2247
-123380.204	779878	6421	-123380.204	383658	6186
-123380.204	925265	5017	-123380.204	179332	4881
-123380.204	365050	1835	-123380.204	518828	432
-123380.204	3307	735	-123380.204	579995	183
-123380.204	652179	5703	-123380.204	423665	3152
-123380.204	208803	6223	-123380.204	360476	3647
-123380.204	862495	3657	-123380.204	166220	1502
-123380.204	127619	5679	-123380.204	204860	5437
-123380.204	596257	405	-123380.204	690707	5370
-123380.204	778331	1280	-123380.204	879211	453
-123380.204	171251	5015	-123380.204	626891	32
-123380.204	726787	1550	-123380.204	395914	5801
-123380.204	145560	5410	-123380.204	907342	1133
-123380.204	616776	3283	-123380.204	718876	1104
-123380.204	385116	7397	-123380.204	863691	481
-123380.204	88407	6844	-123380.204	367683	4650
-123380.204	136094	7099	-123380.204	105392	7874
-123380.204	321574	1185	-123380.204	393318	4188
-123380.204	791087	4308	-123380.204	117891	1818
-123380.204	844630	7711	-123380.204	746304	3335
-123380.204	236935	3873	-123380.204	137790	5503
-123380.204	271769	7649	-123380.204	785068	3000
-123380.204	286163	6626	-123380.204	63470	4478
-123380.204	555074	6182	-123380.204	807339	4334
-123380.204	526265	1962	-123380.204	227563	63
-123380.204	922042	492	-123380.204	754616	4405
-123380.204	51375	148	-123380.204	383371	3818
-123380.204	277405	5638	-123380.204	197794	4439
-123380.204	957757	6403	-123380.204	798091	5260
-123380.204	208737	7212	-123380.204	139080	7105
-123380.204	981709	6322	-123380.204	735663	4273
-123380.204	370434	1832	-123380.204	99433	4520
-123380.204	761633	50	-123380.204	309767	3240
-123380.204	992389	77	-123380.204	715588	6020
-123380.204	609004	2777	-123380.204	232226	235
-123380.204	164582	7991	-123380.204	534583	6098
-123380.204	949412	4952	-123380.204	52311	7800
-123380.204	435399	2896	-123380.204	293349	2520
-123380.204	296924	1702	-123380.204	971770	2708
-123380.204	646573	741	-123380.204	735406	4759
-123380.204	46908	3949	-123380.204	957792	2069
-123380.204	789171	2365	-123380.204	478421	311
-123380.204	697914	6806	-123380.204	396943	5911
-123380.204	448012	7413	-123380.204	997222	229
-123380.204	303628	7673	-123380.204	949086	1426
-123380.204	626264	3014	-123380.204	480046	2484
-123380.204	25530	4179	-123380.204	994880	5709
-123380.204	71845	7650	-123380.204	830618	3377
-123380.204	527412	7073	-123380.204	745674	3511
-123380.204	850953	4066	-123646.100	694918	1206
-123380.204	477475	6934	-123646.100	718336	2280
-123380.204	524904	4913	-123646.100	643311	888
-123380.204	832635	3239	-123646.100	123098	2191
-123380.204	602588	5436	-123646.100	789916	4259
-123380.204	417342	6942	-123646.100	570518	6831
-123380.204	422776	5742	-123646.100	10968	1492
-123380.204	115831	3223	-123646.100	447198	7824
-123380.204	612202	4269	-123656.753	471811	5132
-123380.204	322964	3451	-123793.364	571443	4528
-123380.204	392717	4834	-123793.364	311384	5074
-123380.204	988709	2430	-123793.364	600244	3830
-123380.204	612837	3056	-123793.364	393453	1355
-123380.204	548625	7854	-123793.364	943783	5797
-123380.204	911887	6629	-123817.802	188498	258
-123380.204	170485	6091	-123817.802	156983	5262
-123380.204	317070	7733	-123817.802	202790	198
-123380.204	245802	5336	-123817.802	80	2188
-123380.204	662807	4554	-123817.802	84256	2662

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

-123817.802	866824	7117	1541 perturbed starting value run(s) did not converge or were rejected in the third stage.
-123817.802	802771	4878	
-123817.802	896300	5588	
-123817.802	100054	1225	WARNING: THE BEST LOGLIKELIHOOD VALUE WAS NOT REPLICATED. THE SOLUTION MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA. INCREASE THE NUMBER OF RANDOM STARTS.
-123817.802	704526	6256	
-123817.802	247709	6305	
-123817.802	810668	4172	
-123817.802	789701	6909	
-123817.802	654884	1352	
-123817.802	818908	5898	THE STANDARD ERRORS OF THE MODEL PARAMETER ESTIMATES MAY NOT BE TRUSTWORTHY FOR SOME PARAMETERS DUE TO A NON-POSITIVE DEFINITE FIRST-ORDER DERIVATIVE PRODUCT MATRIX. THIS MAY BE DUE TO THE STARTING VALUES BUT MAY ALSO BE AN INDICATION OF MODEL NONIDENTIFICATION. THE CONDITION NUMBER IS 0.573D-10. PROBLEM INVOLVING THE FOLLOWING PARAMETER:
-123817.802	368364	4774	Parameter 62, %C#6%: SLPDURC WITH SLPNSSNC
-123817.802	421731	886	
-123817.802	662359	1954	
-123817.802	218683	5325	
-123817.802	462159	6130	
-123817.802	362536	7653	
-123817.802	402049	4468	
-123817.802	763134	4903	
-123817.802	157736	1524	
-123817.802	115267	2689	
-123817.802	667120	2859	
-123817.802	515742	7693	
-123817.802	940517	2728	THE MODEL ESTIMATION TERMINATED NORMALLY
-123817.802	167220	2608	
-124083.504	937225	394	MODEL FIT INFORMATION
-124083.504	450164	5105	
-124083.504	975027	1552	Number of Free Parameters 69
-124083.504	920593	611	Loglikelihood
-124083.504	429802	3983	
-124083.504	924648	6655	
-124083.504	945925	2583	H0 Value -105840.802
-124083.504	269171	2307	H0 Scaling Correction Factor 1.1409 for MLR
-124096.895	662718	460	
-124096.895	789081	6962	Information Criteria
-124096.895	135288	5248	
-124096.895	819788	4192	
-124096.895	301105	4652	Akaike (AIC) 211819.605
-124096.895	91231	727	Bayesian (BIC) 212387.673
-124096.895	886520	3171	Sample-Size Adjusted BIC 212168.392 (n* = (n + 2) / 24)
-124096.895	429510	4040	
-124096.895	299736	7688	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THE ESTIMATED MODEL
-124096.895	785215	3364	
-124096.895	883355	2140	
-124117.383	311796	4910	
-124117.383	109599	3448	
-124117.383	676560	1250	Latent Classes
-124117.383	272240	6376	
-124241.254	737045	1638	
-124241.254	808639	5924	1 6256.52283 0.22504
-124241.254	887941	2425	2 2512.95967 0.09039
-124241.254	504509	5610	3 3788.57316 0.13627
-124241.254	730263	2641	4 7106.88021 0.25562
-124241.254	187594	3708	5 3150.58120 0.11332
-124367.958	50708	6449	6 4986.48292 0.17936
-124608.382	406734	2605	
-124709.900	954581	3583	FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON ESTIMATED POSTERIOR PROBABILITIES
-124709.900	692357	2081	
-124709.900	903246	3176	
-124709.900	970728	1371	
-124709.900	517917	5157	Latent Classes
-124709.900	933596	2349	
-124709.900	232509	2745	
-124709.900	472141	5833	1 6256.52283 0.22504
-124709.900	723642	3817	2 2512.95967 0.09039
-124709.900	181178	1069	3 3788.57316 0.13627
-124709.900	767778	7158	4 7106.88021 0.25562
-124709.900	124303	7695	5 3150.58120 0.11332
-124709.900	21345	199	6 4986.48292 0.17936

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

FINAL CLASS COUNTS AND PROPORTIONS FOR THE LATENT CLASSES BASED ON THEIR MOST LIKELY LATENT CLASS MEMBERSHIP							SLPDURC	0.199	0.017	11.802	0.000
Class Counts and Proportions							WORRYNC WITH SOLC	0.138	0.002	56.172	0.000
Latent Classes							SLPDURC	-0.911	0.017	-52.271	0.000
1	6431	0.23131					SOLC WITH SLPDURC	-0.131	0.003	-48.309	0.000
2	2290	0.08237					Means				
3	3705	0.13326					SLPNSSNC	0.787	0.009	86.467	0.000
4	7760	0.27912					WORRYNC	0.855	0.004	196.544	0.000
5	3166	0.11388					SOLC	-1.055	0.000	*****	0.000
6	4450	0.16006					SLPDURC	-0.752	0.014	-55.549	0.000
CLASSIFICATION QUALITY							Variances				
Entropy	0.878						SLPNSSNC	0.473	0.005	88.730	0.000
Average Latent Class Probabilities for Most Likely Latent Class Membership (Row) by Latent Class (Column)							WORRYNC	0.988	0.020	49.163	0.000
	1	2	3	4	5	6	SOLC	0.020	0.001	36.573	0.000
1	0.928	0.063	0.001	0.007	0.000	0.001	SLPDURC	1.115	0.022	49.574	0.000
2	0.093	0.900	0.001	0.001	0.001	0.004	Latent Class 2				
3	0.000	0.000	0.996	0.001	0.001	0.001	SLPNSSNC WITH WORRYNC	0.107	0.027	3.924	0.000
4	0.008	0.002	0.004	0.852	0.002	0.132	SOLC	-0.016	0.007	-2.437	0.015
5	0.001	0.001	0.007	0.002	0.981	0.007	SLPDURC	0.066	0.025	2.581	0.010
6	0.002	0.005	0.008	0.099	0.006	0.880	WORRYNC WITH SOLC	-0.076	0.004	-17.336	0.000
Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)							SLPDURC	0.208	0.040	5.254	0.000
	1	2	3	4	5	6	SOLC WITH SLPDURC	-0.058	0.006	-9.098	0.000
1	0.954	0.034	0.000	0.009	0.001	0.001	Means				
2	0.161	0.820	0.001	0.007	0.001	0.010	SLPNSSNC	0.679	0.017	39.866	0.000
3	0.001	0.001	0.974	0.009	0.006	0.009	WORRYNC	-0.080	0.027	-2.949	0.003
4	0.006	0.000	0.000	0.930	0.001	0.062	SOLC	-1.054	0.000	-6158.720	0.000
5	0.000	0.001	0.001	0.004	0.986	0.008	SLPDURC	0.157	0.022	6.983	0.000
6	0.002	0.002	0.001	0.206	0.005	0.785	Variances				
Logits for the Classification Probabilities for the Most Likely Latent Class Membership (Column) by Latent Class (Row)							SLPNSSNC	0.473	0.005	88.730	0.000
	1	2	3	4	5	6	WORRYNC	0.988	0.020	49.163	0.000
1	6.500	3.171	-1.890	1.869	-0.946	0.000	SOLC	0.020	0.001	36.573	0.000
2	2.821	4.447	-2.894	-0.260	-2.318	0.000	SLPDURC	1.115	0.022	49.574	0.000
3	-2.122	-2.648	4.647	-0.068	-0.441	0.000	Latent Class 3				
4	-2.335	-5.695	-4.844	2.710	-4.048	0.000	SLPNSSNC WITH WORRYNC	0.097	0.015	6.565	0.000
5	-3.300	-2.693	-1.970	-0.648	4.827	0.000	SOLC	0.029	0.003	8.616	0.000
6	-6.242	-6.076	-6.711	-1.340	-5.134	0.000	SLPDURC	-0.125	0.016	-7.852	0.000
MODEL RESULTS							WORRYNC WITH SOLC	0.049	0.005	9.155	0.000
Two-Tailed							SLPDURC	-0.497	0.022	-22.271	0.000
Estimate	S.E.	Est./S.E.	P-Value				SOLC WITH SLPDURC	-0.067	0.004	-15.090	0.000
Latent Class 1							Means				
SLPNSSNC WITH WORRYNC	-0.223	0.017	-12.898	0.000			SLPNSSNC	1.101	0.011	101.666	0.000
SOLC	-0.033	0.003	-12.984	0.000			WORRYNC	-0.580	0.015	-37.798	0.000
							SOLC	0.910	0.000	9470.622	0.000
							SLPDURC	0.494	0.016	31.282	0.000
							Variances				
							SLPNSSNC	0.473	0.005	88.730	0.000
							WORRYNC	0.988	0.020	49.163	0.000

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

SOLC	0.020	0.001	36.573	0.000	SLPNSSNC	0.901	0.013	68.010	0.000
SLPDURC	1.115	0.022	49.574	0.000	WORRYNC	-0.509	0.017	-30.758	0.000
Latent Class 4					SOLC	-0.072	0.000	-768.888	0.000
SLPNSSNC WITH					SLPDURC	0.471	0.017	27.896	0.000
WORRYNC	-0.145	0.023	-6.286	0.000	Variances				
SOLC	0.023	0.003	6.466	0.000	SLPNSSNC	0.473	0.005	88.730	0.000
SLPDURC	0.132	0.021	6.144	0.000	WORRYNC	0.988	0.020	49.163	0.000
WORRYNC WITH					SOLC	0.020	0.001	36.573	0.000
SOLC	-0.133	0.002	-54.161	0.000	SLPDURC	1.115	0.022	49.574	0.000
SLPDURC	-0.828	0.018	-46.563	0.000	Categorical Latent Variables				
SOLC WITH					Means				
SLPDURC	0.120	0.003	43.259	0.000	C#1	0.227	0.028	8.058	0.000
Means					C#2	-0.685	0.041	-16.645	0.000
SLPNSSNC	0.972	0.010	94.865	0.000	C#3	-0.275	0.028	-9.843	0.000
WORRYNC	0.535	0.007	80.781	0.000	C#4	0.354	0.035	10.135	0.000
SOLC	-0.071	0.000	-836.073	0.000	C#5	-0.459	0.029	-15.917	0.000
SLPDURC	-0.373	0.013	-27.768	0.000	QUALITY OF NUMERICAL RESULTS				
Variances					Condition Number for the Information Matrix				
SLPNSSNC	0.473	0.005	88.730	0.000	0.573E-10				
WORRYNC	0.988	0.020	49.163	0.000	(ratio of smallest to largest eigenvalue)				
SOLC	0.020	0.001	36.573	0.000	TECHNICAL 11 OUTPUT				
SLPDURC	1.115	0.022	49.574	0.000	Random Starts Specifications for the k-1 Class Analysis Model				
Latent Class 5					Number of initial stage random starts				
SLPNSSNC WITH					8000				
WORRYNC	0.014	0.013	1.147	0.251	Number of final stage optimizations				
SOLC	-0.001	0.001	-2.245	0.025	2000				
SLPDURC	-0.004	0.014	-0.268	0.789	VUONG-LO-MENDELL-RUBIN LIKELIHOOD RATIO TEST FOR 5 (H0) VERSUS 6 CLASSES				
WORRYNC WITH					H0 Loglikelihood Value				
SOLC	-0.011	0.001	-11.157	0.000	-107985.460				
SLPDURC	-0.407	0.020	-20.611	0.000	2 Times the Loglikelihood Difference				
SOLC WITH					4289.315				
SLPDURC	0.004	0.001	4.415	0.000	Difference in the Number of Parameters				
Means					11				
SLPNSSNC	1.184	0.012	97.778	0.000	Mean				
WORRYNC	-1.338	0.019	-71.482	0.000	-106396860.944				
SOLC	2.133	0.008	279.403	0.000	Standard Deviation				
SLPDURC	0.900	0.020	46.072	0.000	150385030.149				
Variances					P-Value				
SLPNSSNC	0.473	0.005	88.730	0.000	0.2396				
WORRYNC	0.988	0.020	49.163	0.000	LO-MENDELL-RUBIN ADJUSTED LRT TEST				
SOLC	0.020	0.001	36.573	0.000	Value				
SLPDURC	1.115	0.022	49.574	0.000	4251.544				
Latent Class 6					P-Value				
SLPNSSNC WITH					0.2396				
WORRYNC	0.024	0.020	1.194	0.233	TECHNICAL 14 OUTPUT				
SOLC	0.004	0.004	1.007	0.314	Random Starts Specifications for the k-1 Class Analysis Model				
SLPDURC	0.002	0.017	0.121	0.904	Number of initial stage random starts				
WORRYNC WITH					8000				
SOLC	0.082	0.003	25.708	0.000	Number of final stage optimizations				
SLPDURC	-0.318	0.024	-13.211	0.000	2000				
SOLC WITH					Random Starts Specification for the k-1 Class Model for Generated Data				
SLPDURC	-0.063	0.005	-13.511	0.000	Number of initial stage random starts				
Means					16				
					Number of final stage optimizations				
					8				
					Random Starts Specification for the k Class Model for Generated Data				
					Number of initial stage random starts				
					400				
					Number of final stage optimizations				
					80				
					Number of bootstrap draws requested				
					500				
					PARAMETRIC BOOTSTRAPPED LIKELIHOOD RATIO TEST FOR 5 (H0) VERSUS 6 CLASSES				
					H0 Loglikelihood Value				
					-107985.460				
					2 Times the Loglikelihood Difference				
					4289.315				

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”

Difference in the Number of Parameters 11
Approximate P-Value 0.0000
Successful Bootstrap Draws 493

3463 Stoner Ave.
Los Angeles, CA 90066

Tel: (310) 391-9971
Fax: (310) 391-8971
Web: www.StatModel.com
Support: Support@StatModel.com

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WARNING: OF THE 493 BOOTSTRAP DRAWS, 452 DRAWS HAD BOTH A SMALLER LRT VALUE THAN THE OBSERVED LRT VALUE AND NOT A REPLICATED BEST LOGLIKELIHOOD VALUE FOR THE 6-CLASS MODEL.

THIS MEANS THAT THE P-VALUE MAY NOT BE TRUSTWORTHY DUE TO LOCAL MAXIMA.

INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.

WARNING: 7 OUT OF 500 BOOTSTRAP DRAWS DID NOT CONVERGE.

INCREASE THE NUMBER OF RANDOM STARTS USING THE LRTSTARTS OPTION.

PLOT INFORMATION

The following plots are available:

- Histograms (sample values)
- Scatterplots (sample values)
- Sample means
- Estimated means, medians, modes, and percentiles
- Sample and estimated means
- Observed individual values
- Estimated means and observed individual values
- Estimated overall and class-specific distributions

SAVEDATA INFORMATION

Save file
Step2_LPA6M5.dat

Order and format of variables

SLPNSSNC F10.3
WORRYNC F10.3
SOLC F10.3
SLPDURC F10.3
CPROB1 F10.3
CPROB2 F10.3
CPROB3 F10.3
CPROB4 F10.3
CPROB5 F10.3
CPROB6 F10.3
C F10.3
ID I6

Save file format
11F10.3 I6

Save file record length 10000

Save missing symbol *

DIAGRAM INFORMATION

Mplus diagrams are currently not available for Mixture analysis.

No diagram output was produced.

Beginning Time: 13:26:45
Ending Time: 08:20:25
Elapsed Time: 114:53:40

MUTHEN & MUTHEN

Secondary Analysis in R

Set graphic theme

```
month_theme <- theme(axis.text.x = element_text(angle = 45, hjust=1, vjust=1, size=7,
  family="Serif"),
  axis.title.x = element_blank(),
  strip.text.x = element_text(size = 10,
  family="Serif"),
  axis.text.y = element_text(size=7,
  family="Serif"),
  axis.title.y = element_text(size=7,
  family="Serif"),
  legend.title = element_blank(),
  legend.text = element_text(size=7,
  family="Serif"),
  legend.position="right",
  panel.grid.major.y = element_blank(),
  panel.grid.minor.y = element_blank())

set_theme(
  geom.outline.color = "white",
  geom.outline.size = 0,
  geom.label.size = 2,
  geom.label.color = "#040540",
  axis.textcolor = "#040540",
  axis.title.color = "#040540",
  legend.color = "#040540",
  legend.title.color = "#040540",
  base = theme_bw()
)
```

Apply classes

```
# Read Mplus data from txt file
classData<-read.table("Step2_LPA5M5.txt", header=FALSE,na.strings=c("", ""))

## Names of columns - taken from MPlus output
names(classData) <- c("SLPNSSNC", "WORRYNC", "SOLC", "SLPDURC", "CPROB1", "CPROB2", "CPROB3", "CPROB4", "CPR
OB5", "C", "ID")

data$ID <- data$X

## Merge the two
merged <- merge(classData, data, by="ID")

## Probability
probability <- merged[,c("CPROB1", "CPROB2", "CPROB3", "CPROB4", "CPROB5")] # select the probability columns
probability$max<-apply(probability, 1, max) # obtain the max value across the columns

# Add the 'max' column into the data as "weight"
merged$weight <- probability$max

## Create class column to text:
merged$Class <- ifelse(merged$C==1,"Class1",
  ifelse(merged$C==2, "Class2",
  ifelse(merged$C==3, "Class3",
  ifelse(merged$C==4,"Class4", "Class5"))))

merged$Class <- as.factor(merged$Class)

table(is.na(merged$Class))

##
## FALSE
## 27802
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```
merged[is.na(merged["Class"]), "ID"]
```

Double check missingness & describe classes

```
predictors <- c('Gender_dummy', 'YEARGROUP_cat_dummy', 'SCI_02_binary')
```

```
for(x in predictors){
  merged[,x] <- as.character(merged[,x])
}
```

```
# get descriptives
```

```
table1::table1(~ Gender_dummy + Gender + YEARGROUP_cat_dummy +
  YEARGROUP_cat + SCI_02_binary +
  SCI_02 +
  sleepiness_num +
  worry_stops_sleep_num +
  sol +
  sch_sleep_duration|Class,
  data = merged,
  topclass="Rtable1-zebra")
```

```
## Get nicer `table1` .docx output by simply installing the `flextable` package
```

```
##
## 1          Class1      Class2      Class3
## 2          (N=1037)    (N=4825)    (N=1250)
## 3      Gender_dummy
## 4          0      681 (65.7%)    2982 (61.8%)    772 (61.8%)
## 5          1      356 (34.3%)    1843 (38.2%)    478 (38.2%)
## 6          Gender
## 7          Female      681 (65.7%)    2982 (61.8%)    772 (61.8%)
## 8          Male      356 (34.3%)    1843 (38.2%)    478 (38.2%)
## 9      YEARGROUP_cat_dummy
## 10         0      367 (35.4%)    1426 (29.6%)    181 (14.5%)
## 11         1      448 (43.2%)    2438 (50.5%)    628 (50.2%)
## 12         2      222 (21.4%)    961 (19.9%)    441 (35.3%)
## 13      YEARGROUP_cat
## 14      Year 10-13      222 (21.4%)    961 (19.9%)    441 (35.3%)
## 15      Year 5-6      367 (35.4%)    1426 (29.6%)    181 (14.5%)
## 16      Year 7-9      448 (43.2%)    2438 (50.5%)    628 (50.2%)
## 17      SCI_02_binary
## 18      good sleep      316 (30.5%)    4823 (100.0%)    1250 (100%)
## 19      probable insomnia      721 (69.5%)    2 (0.0%)    0 (0%)
## 20      SCI_02
## 21      Mean (SD)      1.97 (0.895)    4.27 (0.762)    5.39 (1.24)
## 22      Median [Min, Max]      2.00 [0, 3.00]    4.00 [2.00, 6.00]    5.00 [4.00, 8.00]
## 23      sleepiness_num
## 24      Mean (SD)      1.96 (0.593)    2.21 (0.655)    4.15 (0.378)
## 25      Median [Min, Max]      2.00 [1.00, 3.00]    2.00 [1.00, 3.00]    4.00 [3.00, 5.00]
## 26      Missing      23 (2.2%)    187 (3.9%)    0 (0%)
## 27      worry_stops_sleep_num
## 28      Mean (SD)      3.38 (1.14)    2.73 (1.03)    2.84 (1.14)
## 29      Median [Min, Max]      4.00 [1.00, 5.00]    3.00 [1.00, 5.00]    3.00 [1.00, 5.00]
## 30      Missing      41 (4.0%)    239 (5.0%)    61 (4.9%)
## 31      sol
## 32      Mean (SD)      1.35 (0.565)    1.01 (0.559)    0.848 (0.589)
## 33      Median [Min, Max]      1.49 [0, 2.00]    0.964 [0, 2.00]    0.688 [0, 2.00]
## 34      Missing      40 (3.9%)    198 (4.1%)    57 (4.6%)
## 35      sch_sleep_duration
## 36      Mean (SD)      7.14 (1.69)    7.58 (1.54)    7.27 (1.56)
## 37      Median [Min, Max]      7.19 [3.00, 11.8]    7.63 [3.04, 12.0]    7.27 [3.00, 12.0]
## 38      Missing      144 (13.9%)    649 (13.5%)    161 (12.9%)
## 39
## 40      Class4      Class5      Overall
## 41      (N=2335)    (N=18355)    (N=27802)
## 42
## 43      1701 (72.8%)    9072 (49.4%)    15208 (54.7%)
## 44      634 (27.2%)    9283 (50.6%)    12594 (45.3%)
## 45
## 46      1701 (72.8%)    9072 (49.4%)    15208 (54.7%)
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```
## 7 634 (27.2%) 9283 (50.6%) 12594 (45.3%)
## 8
## 9 319 (13.7%) 5242 (28.6%) 7535 (27.1%)
## 10 1244 (53.3%) 9837 (53.6%) 14595 (52.5%)
## 11 772 (33.1%) 3276 (17.8%) 5672 (20.4%)
## 12
## 13 772 (33.1%) 3276 (17.8%) 5672 (20.4%)
## 14 319 (13.7%) 5242 (28.6%) 7535 (27.1%)
## 15 1244 (53.3%) 9837 (53.6%) 14595 (52.5%)
## 16
## 17 473 (20.3%) 18355 (100%) 25217 (90.7%)
## 18 1862 (79.7%) 0 (0%) 2585 (9.3%)
## 19
## 20 1.57 (0.994) 7.22 (0.763) 5.96 (2.11)
## 21 2.00 [0, 3.00] 7.00 [6.00, 8.00] 7.00 [0, 8.00]
## 22
## 23 4.06 (0.656) 1.65 (0.628) 2.07 (1.02)
## 24 4.00 [3.00, 5.00] 2.00 [1.00, 3.00] 2.00 [1.00, 5.00]
## 25 95 (4.1%) 374 (2.0%) 679 (2.4%)
## 26
## 27 3.65 (1.03) 1.95 (0.873) 2.32 (1.10)
## 28 4.00 [1.00, 5.00] 2.00 [1.00, 5.00] 2.00 [1.00, 5.00]
## 29 89 (3.8%) 609 (3.3%) 1039 (3.7%)
## 30
## 31 1.38 (0.590) 0.632 (0.478) 0.795 (0.572)
## 32 1.51 [0, 2.00] 0.492 [0, 2.00] 0.560 [0, 2.00]
## 33 92 (3.9%) 629 (3.4%) 1016 (3.7%)
## 34
## 35 6.61 (1.62) 8.24 (1.40) 7.90 (1.56)
## 36 6.64 [3.00, 11.9] 8.28 [3.00, 12.0] 7.99 [3.00, 12.0]
## 37 320 (13.7%) 2428 (13.2%) 3702 (13.3%)
```

```
# get descriptives
```

```
table1::table1(~ Class,
  data = merged,
  topclass="Rtable1-zebra")
```

```
## Get nicer `table1` .docx output by simply installing the `flextable` package
```

```
## Overall
## 1 (N=27802)
## 2 Class
## 3 Class1 1037 (3.7%)
## 4 Class2 4825 (17.4%)
## 5 Class3 1250 (4.5%)
## 6 Class4 2335 (8.4%)
## 7 Class5 18355 (66.0%)
```

Graph classes

```
# create a percentage variable
```

```
merged$Class2 <- factor(merged$Class, levels = c("Class1", "Class2", "Class3", "Class4", "Class5"),
  labels = c("Poor sleepers (3.7%)",
    "Moderate sleepers (17.4%)",
    "Moderate sleepy sleepers (4.5%)",
    "Poor sleepy sleepers (8.4%)",
    "Good sleepers (66.0%)"))
```

```
merged$SCI_02_S <- scale(merged$SCI_02, center = TRUE, scale = TRUE)
merged$sleepiness_S <- scale(merged$sleepiness_num, center = TRUE, scale = TRUE)
merged$worry_S <- scale(merged$worry_stops_sleep_num, center = TRUE, scale = TRUE)
merged$sol_S <- scale(merged$sol, center = TRUE, scale = TRUE)
merged$dur_S <- scale(merged$sch_sleep_duration, center = TRUE, scale = TRUE)
```

```
# separate graphs for class
```

```
# Select only columns that are needed:
```

```
data2 <- merged[,c("Class2", "SCI_02_S", "sleepiness_S", "worry_S", "sol_S", "dur_S", "ID")]
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

Summary

```
summary(data2)
```

```
##          Class2      SCI_02_S.V1  sleepiness_S.V1
## Poor sleepers (3.7%)      : 1037  Min.  :-2.8248292  Min.  :-1.0538
## Moderate sleepers (17.4%) : 4825  1st Qu.: -0.4538482  1st Qu.: -1.0538
## Moderate sleepy sleepers (4.5%): 1250  Median : 0.4945442  Median : -0.0717
## Poor sleepy sleepers (8.4%) : 2335  Mean   : 0.0000000  Mean   : 0.0000
## Good sleepers (66.0%)    :18355  3rd Qu.: 0.9687404  3rd Qu.: -0.0717
##                               Max.   : 0.9687404  Max.   : 2.8745
##                               NA's   :679
## worry_S.V1  sol_S.V1    dur_S.V1    ID
## Min.  :-1.2049  Min.  :-1.3912  Min.  :-3.152  Min.  : 1
## 1st Qu.: -1.2049  1st Qu.: -0.7475  1st Qu.: -0.612  1st Qu.: 6951
## Median : -0.2923  Median : -0.4116  Median : 0.056  Median :13902
## Mean   : 0.0000  Mean   : 0.0000  Mean   : 0.000  Mean   :13902
## 3rd Qu.: 0.6202  3rd Qu.: 0.6799  3rd Qu.: 0.683  3rd Qu.:20852
## Max.   : 2.4452  Max.   : 2.1073  Max.   : 2.619  Max.   :27802
## NA's  :1039  NA's  :1016  NA's  :3702
```

```
class(data2)
```

```
## [1] "data.frame"
```

Long data: just indicators

```
long_data <- pivot_longer(data = data2,
  cols = sleepiness_S:dur_S,
  names_to = "scale",
  values_to = "measurement")
```

```
long_data$scale <- factor(long_data$scale, levels = c("sleepiness_S", "worry_S", "sol_S", "dur_S"),
  labels = c("Sleepiness", "Worry disrupts sleep", "SOL", "Sleep duration"))
```

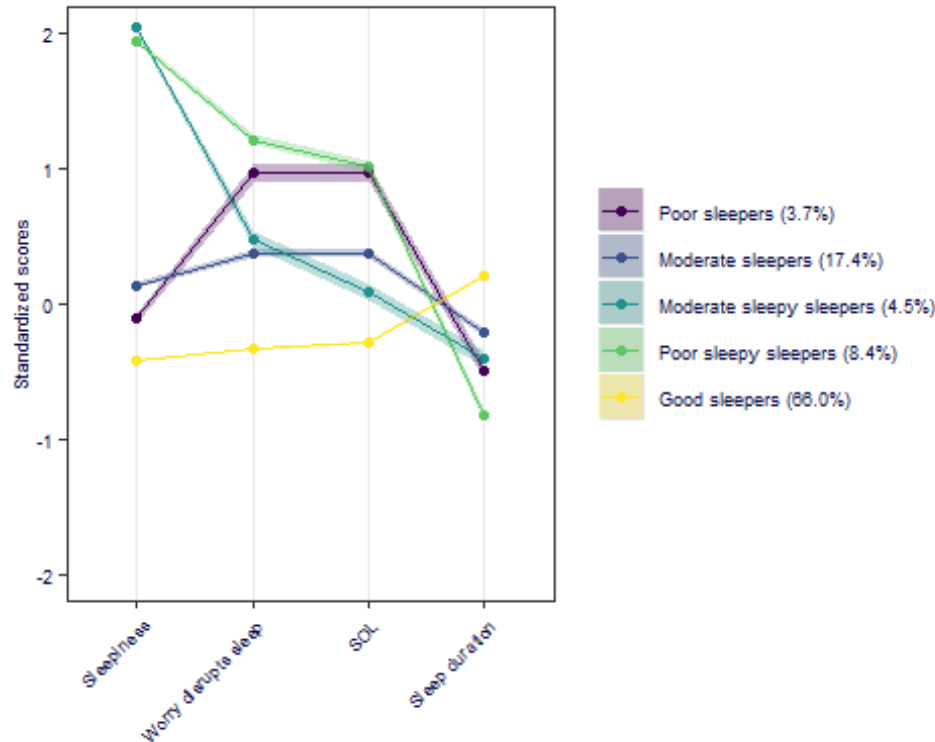
```
summary(long_data)
```

```
##          Class2      SCI_02_S.V1    ID
## Poor sleepers (3.7%)      : 4148  Min.  :-2.8248292  Min.  : 1
## Moderate sleepers (17.4%) :19300  1st Qu.: -0.4538482  1st Qu.: 6951
## Moderate sleepy sleepers (4.5%): 5000  Median : 0.4945442  Median :13902
## Poor sleepy sleepers (8.4%) : 9340  Mean   : 0.0000000  Mean   :13902
## Good sleepers (66.0%)    :73420  3rd Qu.: 0.9687404  3rd Qu.:20852
##                               Max.   : 0.9687404  Max.   :27802
##
##          scale      measurement.V1
## Sleepiness      :27802  Min.  :-3.152
## Worry disrupts sleep:27802  1st Qu.: -0.810
## SOL              :27802  Median : -0.072
## Sleep duration   :27802  Mean   : 0.000
##                 3rd Qu.: 0.620
##                 Max.   : 2.875
##                 NA's   :6436
```

```
lineFig <- ggplot(long_data, aes(x=scale, y=measurement, group=Class2, colour=Class2, fill=Class2)) +
  stat_summary(fun=mean, geom="point") + stat_summary(fun=mean, geom="line") + stat_summary(fun.data = mean_cl_normal
, geom="ribbon", alpha = 0.3, colour=NA) + ylab(label= 'Standardized scores') + scale_colour_viridis_d() + scale_fill_viridis_d()
+ month_theme + coord_cartesian(ylim = c(-2,2))
```

```
lineFig
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”



Investigate relationship with SCI-02

```
# graph
## Long data: with SCI-02
long_data <- pivot_longer(data = data2,
  cols = SCI_02_S:dur_S,
  names_to = "scale",
  values_to = "measurement")

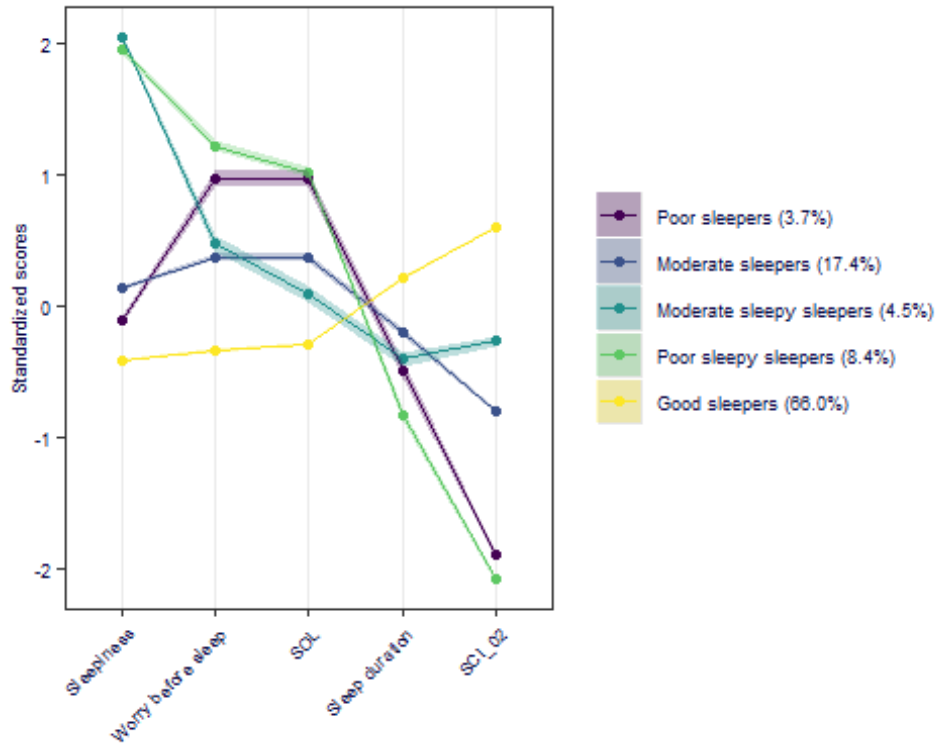
long_data$scale <- factor(long_data$scale, levels = c("sleepiness_S", "worry_S", "sol_S", "dur_S", "SCI_02_S"),
  labels = c("Sleepiness", "Worry before sleep", "SOL", "Sleep duration", "SCI_02"))

summary(long_data)

##           Class2           ID
## Poor sleepers (3.7%)       :5185  Min.   : 1
## Moderate sleepers (17.4%)  :24125 1st Qu.: 6951
## Moderate sleepy sleepers (4.5%): 6250  Median :13902
## Poor sleepy sleepers (8.4%)  :11675  Mean   :13902
## Good sleepers (66.0%)      :91775  3rd Qu.:20852
##                               Max.   :27802
##
##           scale      measurement.V1
## Sleepiness      :27802  Min.   :-3.152
## Worry before sleep:27802 1st Qu.: -0.780
## SOL              :27802  Median :-0.072
## Sleep duration   :27802  Mean    : 0.000
## SCI_02          :27802  3rd Qu.: 0.620
##                               Max.   : 2.875
##                               NA's   :6436

ggplot(long_data, aes(x=scale, y=measurement, group=Class2, colour=Class2, fill=Class2)) +
  stat_summary(fun=mean, geom="point")+stat_summary(fun=mean, geom="line")+ stat_summary(fun.data = mean_cl_normal,
  geom="ribbon", alpha = 0.3, colour=NA) + ylab(label= 'Standardized scores') + scale_colour_viridis_d() + scale_fill_viridis_d() +
  month_theme
```

Scripts & Outputs “*Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)*”



```
# anova
SCI_02_comp <- aov(SCI_02 ~ Class2, data = merged)

summary(SCI_02_comp)

##           Df Sum Sq Mean Sq F value Pr(>F)
## Class2    4 105084  26271  39364 <2e-16 ***
## Residuals 27797 18551     1
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

pairwise.t.test(merged$SCI_02, merged$Class2,
  p.adjust.method = "bonferroni")

##
## Pairwise comparisons using t tests with pooled SD
##
## data: merged$SCI_02 and merged$Class2
##
##           Poor sleepers (3.7%) Moderate sleepers (17.4%)
## Moderate sleepers (17.4%) <2e-16 -
## Moderate sleepy sleepers (4.5%) <2e-16 <2e-16
## Poor sleepy sleepers (8.4%) <2e-16 <2e-16
## Good sleepers (66.0%) <2e-16 <2e-16
##
##           Moderate sleepy sleepers (4.5%)
## Moderate sleepers (17.4%) -
## Moderate sleepy sleepers (4.5%) -
## Poor sleepy sleepers (8.4%) <2e-16
## Good sleepers (66.0%) <2e-16
##
##           Poor sleepy sleepers (8.4%)
## Moderate sleepers (17.4%) -
## Moderate sleepy sleepers (4.5%) -
## Poor sleepy sleepers (8.4%) -
## Good sleepers (66.0%) <2e-16
##
## P value adjustment method: bonferroni
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

Logistic regression

```
# Select only columns that are needed:
data3 <- merged[,c("Class", "Gender_dummy", "YEARGROUP_cat_dummy", "weight", "ID")]

predictors <- c('Gender_dummy', 'YEARGROUP_cat_dummy')

for(x in predictors){
  data3[,x] <- as.character(data3[,x])
}

# Summary
summary(data3)

## Class Gender_dummy YEARGROUP_cat_dummy weight
## Class1: 1037 Length:27802 Length:27802 Min. :0.3510
## Class2: 4825 Class :character Class :character 1st Qu.:0.8850
## Class3: 1250 Mode :character Mode :character Median :0.9950
## Class4: 2335 Mean :0.9157
## Class5:18355 3rd Qu.:1.0000
## Max. :1.0000
## ID
## Min. : 1
## 1st Qu.: 6951
## Median :13902
## Mean :13902
## 3rd Qu.:20852
## Max. :27802

## Long data
data4 = mlogit.data(data3, choice = "Class", shape="wide")

# Model:
model_LR <- mlogit(Class ~ 1 | Gender_dummy + YEARGROUP_cat_dummy, data=data4, reflevel = "Class5")

#summary
summary(model_LR)

##
## Call:
## mlogit(formula = Class ~ 1 | Gender_dummy + YEARGROUP_cat_dummy,
## data = data4, reflevel = "Class5", method = "nr")
##
## Frequencies of alternatives:choice
## Class5 Class1 Class2 Class3 Class4
## 0.660204 0.037299 0.173549 0.044961 0.083987
##
## nr method
## 7 iterations, 0h:0m:7s
## g'(-H)^-1g = 1.04E-05
## successive function values within tolerance limits
##
## Coefficients :
## Estimate Std. Error z-value Pr(>|z|)
## (Intercept):Class1 -2.349727 0.059826 -39.2761 < 2.2e-16 ***
## (Intercept):Class2 -1.061812 0.033331 -31.8564 < 2.2e-16 ***
## (Intercept):Class3 -3.137894 0.079848 -39.2985 < 2.2e-16 ***
## (Intercept):Class4 -2.389418 0.059998 -39.8249 < 2.2e-16 ***
## Gender_dummy1:Class1 -0.684501 0.067160 -10.1921 < 2.2e-16 ***
## Gender_dummy1:Class2 -0.506638 0.033142 -15.2869 < 2.2e-16 ***
## Gender_dummy1:Class3 -0.477822 0.060346 -7.9180 2.442e-15 ***
## Gender_dummy1:Class4 -0.984424 0.049116 -20.0429 < 2.2e-16 ***
## YEARGROUP_cat_dummy1:Class1 -0.458847 0.072667 -6.3144 2.713e-10 ***
## YEARGROUP_cat_dummy1:Class2 -0.114792 0.037669 -3.0474 0.002308 **
## YEARGROUP_cat_dummy1:Class3 0.594139 0.086180 6.8942 5.417e-12 ***
## YEARGROUP_cat_dummy1:Class4 0.691392 0.065404 10.5711 < 2.2e-16 ***
## YEARGROUP_cat_dummy2:Class1 -0.062002 0.088141 -0.7034 0.481785
## YEARGROUP_cat_dummy2:Class2 0.053350 0.047540 1.1222 0.261776
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```
## YEARGROUP_cat_dummy2:Class3 1.339768 0.091165 14.6961 < 2.2e-16 ***
## YEARGROUP_cat_dummy2:Class4 1.313035 0.070679 18.5775 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Log-Likelihood: -28462
## McFadden R^2: 0.023353
## Likelihood ratio test : chisq = 1361.2 (p.value = < 2.22e-16)
```

```
exp(coefficients(model_LR)) ## risk ratio
```

```
## (Intercept):Class1 (Intercept):Class2
## 0.09539524 0.34582845
## (Intercept):Class3 (Intercept):Class4
## 0.04337404 0.09168299
## Gender_dummy1:Class1 Gender_dummy1:Class2
## 0.50434183 0.60251763
## Gender_dummy1:Class3 Gender_dummy1:Class4
## 0.62013258 0.37365447
## YEARGROUP_cat_dummy1:Class1 YEARGROUP_cat_dummy1:Class2
## 0.63201174 0.89155141
## YEARGROUP_cat_dummy1:Class3 YEARGROUP_cat_dummy1:Class4
## 1.81147094 1.99649191
## YEARGROUP_cat_dummy2:Class1 YEARGROUP_cat_dummy2:Class2
## 0.93988144 1.05479873
## YEARGROUP_cat_dummy2:Class3 YEARGROUP_cat_dummy2:Class4
## 3.81815842 3.71743742
```

```
exp(confint.default(model_LR))
```

```
## 2.5 % 97.5 %
## (Intercept):Class1 0.08484043 0.1072632
## (Intercept):Class2 0.32395830 0.3691750
## (Intercept):Class3 0.03709057 0.0507220
## (Intercept):Class4 0.08151139 0.1031239
## Gender_dummy1:Class1 0.44213823 0.5752967
## Gender_dummy1:Class2 0.56462379 0.6429547
## Gender_dummy1:Class3 0.55095677 0.6979938
## Gender_dummy1:Class4 0.33936156 0.4114127
## YEARGROUP_cat_dummy1:Class1 0.54811380 0.7287517
## YEARGROUP_cat_dummy1:Class2 0.82809917 0.9598656
## YEARGROUP_cat_dummy1:Class3 1.52994298 2.1448034
## YEARGROUP_cat_dummy1:Class4 1.75628654 2.2695499
## YEARGROUP_cat_dummy2:Class1 0.79076449 1.1171179
## YEARGROUP_cat_dummy2:Class2 0.96095520 1.1578067
## YEARGROUP_cat_dummy2:Class3 3.19340534 4.5651373
## YEARGROUP_cat_dummy2:Class4 3.23654718 4.2697789
```

Produce a forest graph

```
oddsSleep <- broom::tidy(model_LR) # extract model estimates into a separate DF
oddsSleep$OR <- exp(coefficients(model_LR)) # extract and save ORs
oddsSleep$CI <- exp(confint.default(model_LR)) # extract and save CIs
oddsSleep$Class <- rep(c("C1",
  "C2",
  "C3",
  "C4"), 4) # create a new variable for class

# might want to adjust these but a place holder for now
oddsSleep$Class <- factor(oddsSleep$Class,
  levels = c("C1",
  "C2",
  "C3",
  "C4",
  "C5"),
  labels = c("Poor sleepers",
  "Moderate sleepers",
  "Moderate sleepy sleepers",
  "Poor sleepy sleepers",
```

Scripts & Outputs “Insomnia symptoms in children and adolescents: Screening for sleep problems with the Two-item Sleep Condition Indicator (SCI-02)”

```

"Good sleepers"))
oddsSleep$term <- gsub("\\:.*", "", oddsSleep$term) # remove class name from the effects
oddsSleep <- dplyr::filter(oddsSleep, term!="(Intercept)") # remove intercept values

#write.table(oddsSleep, file="oddsNon-gamers.csv", row.names=F, sep=",")

oddsSleep$effect <- rep(c("Gender: Boy (vs. Girl)",
  "Year: 7-9 (vs. 5-6)",
  "Year: 10-13 (vs. 5-6)"),
  each=4) # create a new variable for class

# Relevel effect factor
oddsSleep$effect <- factor(oddsSleep$effect,
  levels = c("Gender: Boy (vs. Girl)",
  "Year: 7-9 (vs. 5-6)",
  "Year: 10-13 (vs. 5-6)"))

forestStr <- ggplot(oddsSleep, aes(y=effect,x=OR,colour=Class))+
  geom_pointrangeh(aes(xmin=CI[,1],
    xmax=CI[,2]),
  position=position_dodgev(height=0.75)) + scale_y_discrete(limits=rev) + geom_vline(xintercept=1, linetype=2, color = "grey")
+ labs(x = "Odds Ratio", y=element_blank()) + theme_few() + theme(axis.text.x = element_text(hjust=0.5, vjust=1, size=7, family="Serif"), axis.title.x = element_text(size=7, family="Serif"), axis.text.y = element_text(size=7, family="Serif"), axis.title.y = element_text(size=7, family="Serif"), legend.title = element_blank(), legend.text = element_text(size=7, family="Serif"), legend.position="right") + scale_x_log10() + scale_colour_viridis_d()

print(forestStr)

```

