

Table S1. Items of the Kid-KINDL

PHYSICAL WELL-BEING	FAMILY
P1 ...felt ill	Fa1 ...got on well with my parents
P2 ...headache or stomachache	Fa2 ...felt fine at home
P3 ...tired and sleepy	Fa3 ...quarreled at home
P4 ...strong and full of energy	Fa4 ...stopped from doing certain things
EMOTIONAL WELL-BEING	FRIENDS
E1 ...had fun and laughed a lot	Fr1 ...played with friends
E2 ...bored	Fr2 ...other kids liked me
E3 ...felt alone	Fr3 ...got along well with my friends
E4 ...scared	Fr4 ...felt different from other children*
SELF-ESTEEM	SCHOOL
SE1 ...proud of myself	S1 ...doing my school work was easy
SE2 ...felt on top of the world	S2 ...enjoyed my lessons
SE3 ...felt pleased with myself	S3 ...worried about my future
SE4 ...had lots of good ideas	S4 ...was afraid of bad marks or grades

Negatively worded items have been reverse-coded and reported in **bold** font.

* Item Fr4 was not analyzed in this current study.

Table S2. Baseline factor loadings^a on Chinese Kid-KINDL for four groups

Item #	Unstandardized factor loadings (Standard error)/Standardized factor loadings			
	Boys (n=208)	Girls (n=235)	Parents (n=241)	Children (n=443)
P1	1.000(--)/0.458	1.000(--)/0.464	1.000(--)/0.810	1.000(--)/0.472
P2	1.147(0.253)/0.544	1.381(0.264)/0.634	1.009(0.094)/0.772	1.290(0.180)/0.615
P3	1.922(0.401)/0.714	1.667(0.320)/0.626	0.865(0.088)/0.664	1.746(0.238)/0.667
P4	0.624(0.293)/0.190	1.323(0.320)/0.401	0.438(0.117)/0.258	0.953(0.207)/0.296
E1	1.000(--)/0.208	1.000(--)/0.456	1.000(--)/0.460	1.000(--)/0.328
E2	2.729(1.124)/0.578	1.534(0.237)/0.741	0.732(0.219)/0.322	1.882(0.312)/0.645
E3	1.947(0.845)/0.479	1.609(0.243)/0.793	1.522(0.313)/0.661	2.095(0.335)/0.780
E4	1.274(0.625)/0.321	1.201(0.202)/0.580	1.192(0.256)/0.543	1.665(0.276)/0.618
SE1	1.000(--)/0.620	1.000(--)/0.712	1.000(--)/0.240	1.000(--)/0.620
SE2	1.360(0.143)/0.859	1.029(0.093)/0.733	2.481(0.891)/0.605	1.253(0.099)/0.784
SE3	1.282(0.134)/0.790	1.004(0.089)/0.732	1.717(0.587)/0.473	1.214(0.092)/0.761
SE4	0.792(0.115)/0.522	0.557(0.081)/0.453	1.284(0.479)/0.324	0.648(0.074)/0.442
Fa1	1.000(--)/0.325	1.000(--)/0.744	1.000(--)/0.693	1.000(--)/0.616
Fa2	0.507(0.269)/0.161	0.996(0.170)/0.598	1.055(0.119)/0.777	0.872(0.136)/0.483
Fa3	1.944(0.547)/0.626	0.569(0.141)/0.345	0.600(0.109)/0.428	0.725(0.138)/0.406
Fa4	2.080(0.590)/0.541	0.404(0.153)/0.210	0.485(0.131)/0.278	0.681(0.151)/0.317
Fr1	1.000(--)/0.457	1.000(--)/0.429	1.000(--)/0.403	1.000(--)/0.445
Fr2	1.454(0.270)/0.683	1.476(0.258)/0.637	1.075(0.185)/0.494	1.330(0.182)/0.601
Fr3	1.147(0.214)/0.550	1.315(0.231)/0.644	1.797(0.358)/0.870	1.269(0.171)/0.618
S1	1.000(--)/0.215	1.000(--)/0.641	1.000(--)/0.897	1.000(--)/0.584
S2	1.130(0.449)/0.215	0.940(0.158)/0.535	0.693(0.119)/0.615	0.965(0.160)/0.498
S3	2.713(0.896)/0.480	0.503(0.160)/0.262	0.103(0.085)/0.084*	0.469(0.152)/0.222
S4	5.735(2.102)/0.919	0.548(0.167)/0.276	0.342(0.097)/0.263	0.634(0.170)/0.279

^a Based on correlated traits and uncorrelated methods (CTUM) framework without any constraints.

* p = 0.19; all other p < 0.05

Negatively worded items are in **bold** font.

Table S3. Fit indices of baseline model^a

	Gender group			Rater group			Rater group ^b		
	Boys (n=208)	Girls (n=235)	Configural	Parents (n=241)	Children (n=443) ^c	Configural	Mothers (n=176)	Children (n=443) ^c	Configural
χ^2	306.078*	316.354*	622.433*	288.466*	375.337*	663.802*	278.503*	375.337*	653.840*
<i>df</i>	192	192	384	192	192	384	192	192	384
GFI	0.894	0.900	0.900	0.908	0.934	0.934	0.882	0.934	0.934
NNFI	0.948	0.965	0.959	0.970	0.968	0.969	0.965	0.968	0.967
CFI	0.960	0.974	0.969	0.977	0.976	0.976	0.974	0.976	0.975
RMSEA	0.0476	0.0491	0.0484	0.0438	0.0447	0.0444	0.0478	0.0447	0.0456
SRMR	0.0609	0.0547	0.0547	0.0535	0.0452	0.0452	0.0639	0.0452	0.0452

^aBased on correlated traits and uncorrelated methods (CTUM) framework without any constraints.

^bParent-proxy using mothers' reports only.

^cThe results were the same because they were the same sample.

GFI=goodness of fit index; NNFI=non-normed fit index; CFI=comparable fit index; RMSEA= root mean square error of approximation;

SRMR=standardized root mean square residual.