

Tables

Table 1 Descriptive statistics and sex differences in main variables

	Male	Female	Total sample	t	df	p	Cohen's d
Sample (N / %)	584 (38.4)	937 (61.6)	1521 (100)	/	/	/	/
Participants with at least one apnea symptom (N / %)	152 (26.3)	231 (25)	383 (25.5)	/	/	0.566	/
Age (M /SD)	20.40 (1.8)	20.30 (1.7)	20.30 (1.8)	/	/	/	/
Depression symptoms (M /SD)	5.62 (5.3)	6.98 (5.9)	6.46 (5.7)	4.6	1327	<0.001	0.24
Anxiety symptoms (M /SD)	21.11 (12.8)	27.55 (15.6)	25.08 (14.9)	8.8	1402	<0.001	0.46
Externalizing symptoms (M /SD)	7.34 (6.4)	5.96 (5.00)	6.49 (5.6)	4.4	982	<0.001	0.24

Note. df: degrees of freedom; M/SD: mean/standard deviation; p: p-value; t: t-student

Apnea symptoms were measured by 3 single questions referring to principal symptoms of apnea in the past month, ranging from 0="no sign" to 4="always". Those participants who responded to all items as "never" or "do not know" were not classified as having at least one apnea symptom. Depression, anxiety and externalizing behaviors were measured using RCADS (Revised Child Anxiety and Depression Scale), SMFQ (Short Mood and Feelings Questionnaire) items from the scales of rule breaking and aggression of the adult self-report form, respectively. Higher scores represent increased levels of depression, anxiety and externalizing behaviors.

Table 2: Univariate analyses

Model (comparison)	A	D	E	Twin effect	df	-2LL	AIC	P	
95% CI									
Apnea symptoms									
I	0 (0,0)	0.51 (0,0.70)	0.49 (0.30,0.72)	0 (0,0)	1495	1687.70	-1302.30		
II (I)	0 (0,0.44)	0.51 (0,0.70)	0.49 (0.32,0.72)	/	1496	1687.70	-1304.30	0.99	
III (II)	0.40 (0.19,0.59)	/	0.60 (0.41,0.81)	/	1497	1691.41	-1302.59	0.05	
IV(III)	/	/	1 (1,1)	/	1498	1704.80	-1291.20	<0.001	
Externalizing									
I	0 (0,0.34)	0.42 (0.09,0.57)	0.53 (0.43,0.65)	0.05(0,0.19)	1471	4471.24	1529.24		
II (I)	0 (0,0.36)	0.48 (0.07,0.57)	0.52 (0.43,0.64)	/	1472	4471.69	1527.68	0.50	
III (II)	0.40 (0.29,0.50)	/	0.60 (0.50,0.71)	/	1473	4476.87	1530.67	0.02	
IV(III)	/	/	1 (1,1)	/	1474	4521.26	1573.26	<0.001	
Depression									
I	0.06 (0,0.45)	0.37 (0, 0.54)	0.55 (0.46,0.67)	0.02 (0,0.17)	1505	4808.52	1798.52		
II (I)	0.09 (0,0.45)	0.35 (0,0.54)	0.55 (0.46,0.67)	/	1506	4808.55	1796.55	0.87	
III (II)	0.40 (0.29,0.49)	/	0.60 (0.51,0.71)	/	1507	4811.36	1797.36	0.09	
IV(III)	/	/	1 (1,1)	/	1508	4859.61	1843.60	<0.001	
Anxiety									
I	0.34 (0,0.53)	0.12 (0,0.52)	0.53 (0.44,0.64)	0 (0,0.21)	1508	3906.35	890.35		
II (I)	0.35 (0,0.53)	0.12 (0,0.52)	0.53 (0.44,0.64)	/	1509	3906.35	888.35	0.96	
III (II)	0.45 (0.36,0.54)	/	0.55 (0.46,0.64)	/	1510	3906.69	886.69	0.56	
IV(III)	/	/	1 (1,1)	/	1511	3979.30	957.30	<0.001	

Note. A, additive genetic influence; D, dominant genetic influence E, non-shared environmental influence; -2LL, negative 2 log likelihood; AIC, Akaike's information criterion; CI, confidence interval; df, degrees of freedom; P-value, significance value of the likelihood-ratio chi-square test; rDZ, dizygotic correlations; rMZ, monozygotic correlations. TW, twin effect component on the variance. Bold text indicates best fitting models.

Model legend: I: ADE+TW; II: ADE; III: AE; IV: E

Monozygotic intrapair correlations were 0.53, 0.45, 0.44 and 0.49 for apnea symptoms, externalising, depression and anxiety respectively

Dizygotic intrapair correlations were 0.05, 0.12, 0.14 and 0.20 for apnea symptoms, externalising, depression and anxiety respectively

Table 3: Correlations from the multivariate analysis

Phenotypes	rG	rE	rPh	Proportion of the phenotypic correlation due to	
				Genetic factors	Environmental factors
95% CI				%	
AE					
Apnea-Depression	0.60 (0.32,0.92)	0.02 (-0.16,0.19)	0.25 (0.18,0.31)	95	5
Apnea-Anxiety	0.24 (-0.02, 0.51)	0.20 (0.02, 0.37)	0.22 (0.15,0.29)	47	53
Apnea-Externalizing	0.42 (0.13,0.72)	0.20 (0.02, 0.38)	0.29 (0.22,0.35)	57	43
Depression-Anxiety	0.80 (0.69,0.89)	0.49 (0.40,0.57)	0.62 (0.59,0.65)	54	46
Externalizing-Depression	0.58 (0.41,0.72)	0.41 (0.30, 0.50)	0.47 (0.43,0.50)	48	52
Externalizing-Anxiety	0.39 (0.22,0.55)	0.24 (0.13,0.35)	0.30 (0.25,0.35)	55	45

Note. CI: confidence interval; rE: environmental correlation; rG: genetic correlation rPh = phenotypic correlation