

Addendum

Pubertal development mediates the association between family environment and brain structure and function in childhood – ADDENDUM

Sandra Thijssen, Paul F. Collins and Monica Luciana

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In November 2019, the Adolescent Brain and Cognitive Development consortium communicated that previously released functional MRI data from Philips scanners has been processed incorrectly and should not be analyzed. The resting-state fMRI analyses reported in Thijssen et al. (2019) include data from Philips scanners. We have reanalyzed our resting-state fMRI data excluding participants scanned on a Philips scanner (n = 256). Excluding the Philips data did not significantly affect our results. For the new results, please see below. The conclusions described in the manuscript remain unchanged.

Resting-state fMRI

In the total sample excluding those scanned with Phillips scanners, the total, direct, and indirect effects of Family Environment on cingulo-opercular network–left amygdala functional connectivity were $\beta = 0.068$, p = .003, $\beta = 0.059$, p = .010, $\beta = 0.009$, p = .071, respectively. For cingulo-opercular network-right amygdala functional connectivity, the total, direct, and indirect effects were $\beta = 0.044$, p = .055, $\beta = 0.036$, p = .122, $\beta = 0.008$, p = .106, respectively. Thus, Family Environment was positively

associated with cingulo-opercular network–amygdala functional connectivity. For the left amygdala–cingulo-opercular network functional connectivity, the indirect effect of family environment on functional connectivity via pubertal stage indicated a trend in the expected direction. For right amygdala–cingulo-opercular network functional connectivity, the indirect effect no longer indicates a trend (p > .1). As the effect size of the indirect effect increased from $\beta = 0.007$ to $\beta = 0.008$ when excluding the Philips data, this difference is solely explained by decreased power.

The exploratory analyses stratified by sex suggest that the total and direct effects of Family Environment on cingulo-opercular network–left amygdala functional connectivity were significant for girls, whereas a trend was found for the indirect effect (β = 0.090, p = .005, β = 0.078, p = .017, β = 0.012, p = .093, respectively). For boys, no significant effects were found (β = 0.049, p = .112, β = 0.044, p = .157, β = 0.005, p = .459, respectively). For cingulo-opercular network–right amygdala functional connectivity, no significant effects were found for girls nor boys (girls: β = 0.061, p = .071, β = 0.053, p = .132, β = 0.008, p = .226 for total, direct, and indirect effects, respectively; boys β = 0.030, p = .322, β = 0.023, p = .459, β = 0.007, p = .289, for total, direct, and indirect effects, respectively).

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Table 6. Mediation model parameters--Cinculo-opercular network-amygdala connectivity

		CON-left amygdala				CON-right amygdala			
	β	S.E.	β/S.E.	р	β	S.E.	β /S.E.	p	
Family Environment [∞]	0.059	0.023	2.582	.010	0.036	0.023	1.548	.122	
Pubertal stage ⁺	-0.064	0.033	-1.923	.054	-0.055	0.033	-1.692	.091	
Age	0.002	0.023	0.106	.915	0.021	0.024	0.878	.380	
Sex	-0.031	0.027	-1.160	.246	-0.055	0.028	-1.984	.047	
Race	-0.040	0.023	-1.768	.077	-0.032	0.021	-1.531	.126	
		Outcome: P	ubertal stage		Outcome: Pubertal stage				
Family Environment ^f	-0.136	0.022	-6.189	<.001	-0.136	0.022	-6.189	<.001	
Age	0.231	0.021	11.159	<.001	0.231	0.021	11.159	<.001	
Sex	-0.504	0.016	-30.950	<.001	-0.504	0.016	-30.950	<.001	
Race	0.136	0.020	6.686	<.001	0.136	0.020	6.686	<.001	

Note: CON = cingulo-opercular network; $^{\infty}$ = direct effect; $^{+}$ = indirect effect Step 2; f = indirect effect Step 1.

TableS9. Mediation model parameters--Cinculo-opercular network-amygdala connectivity in girls

		CON-left amygdala				CON-right amygdala			
	β	S.E.	β/S.E.	р	β	S.E.	β/S.E.	р	
Family Environment [∞]	0.078	0.033	2.391	.027	0.053	0.035	1.505	.132	
Pubertal stage ⁺	-0.071	0.039	-1.816	.069	-0.049	0.039	-1.261	.207	
Age	-0.048	0.034	-1.404	.160	0.014	0.031	0.416	.677	
Race	-0.044	0.036	-1.209	.227	-0.050	0.033	-1.637	.102	
		Outcome: Pu	ubertal stage		Outcome: Pubertal stage				
Family Environment ^f	-0.171	0.034	-5.058	<.001	-0.171	0.034	-5.058	<.001	
Age	0.300	0.030	9.991	.001	0.300	0.030	9.991	.001	
Race	0.111	0.034	3.295	<.001	0.111	0.034	3.295	<.001	

Note: $^{\infty}$ = direct effect; $^{+}$ = indirect effect Step 2; f = indirect effect Step 1; CON = cingulo-opercular network.

TableS12. Mediation model parameters—Cinculo-opercular network—amygdala connectivity in boys

		CONeft amygdala				CON-right amygdala			
	β	S.E.	β/S.E.	р	β	S.E.	β/S.E.	р	
Family Environment [∞]	0.044	0.031	1.415	.157	0.023	0.031	0.740	.459	
Pubertal stage ⁺	-0.034	0.043	-0.794	.427	-0.048	0.042	-1.146	.252	
Age	0.042	0.032	1.298	.194	0.027	0.030	0.900	.368	
Race	-0.038	0.029	-1.286	.198	-0.016	0.029	-0.554	.579	
		Outcome: Pu	ubertal stage		Outcome: Pubertal stage				
Family Environment ^f	-0.142	0.036	-3.960	<.001	-0.142	0.036	-3.960	<.001	
Age	0.223	0.035	6.289	<.001	0.223	0.035	6.289	<.001	
Race	0.212	0.031	6.846	<.001	0.212	0.031	6.846	<.001	

Note: $^{\infty}$ = direct effect; $^{+}$ = indirect effect Step 2; f = indirect effect Step 1; CON = cingulo-opercular network.

Somato-motor mouth network-amygdala functional connectivity

For the resting-state model with motor processing measures, onlythe total and direct effects of Family Environment on

SOMM-left amygdala FC were significant (β = 0.060, p = .005, β = 0.061, p = .006, respectively), but not the indirect effect (β = -0.001, p = .847). No associations between Family Environment and SOMM-right amygdala were found (β = -0.013,

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p = .583, $\beta = -0.015$, p = .542, $\beta = 0.002$, p = .693, for total, direct, and indirect effects, respectively).

For the resting-state model with motor processing measures, in girls the total and direct effects, and in boys only the total effect of Family Environment on SOMM–left amygdala FC were significant (girls: $\beta = 0.056$, p = .048, $\beta = 0.064$, p = .036, $\beta = -0.008$, p = .302 for total, direct, and indirect effects, respectively; boys

β = 0.063, p = .041, β = 0.056, p = .074, β = 0.006, p = .357. for total, direct, and indirect effects, respectively). No significant associations were found between Family Environment and SOMM-right amygdala (girls: β = 0.027, p = .421, β = 0.026, p = .452, β = 0.001, p = .886 for total, direct, and indirect effects, respectively; boys β = -0.043, p = .157, β = -0.046, p = .141, β = 0.003, p = .606 for total, direct, and indirect effects, respectively).

TableS6. Mediation model parameters -- Somatomotor-mouth network-amygdala connectivity

		SOMM-left amygdala				SOMM-right amygdala			
	β	S.E.	β/S.E.	р	β	S.E.	β /S.E.	р	
Family Environment [∞]	0.061	0.022	2.767	.006	-0.015	0.024	-0.610	.542	
Pubertal stage ⁺	0.007	0.035	0.199	.843	-0.013	0.033	-0.404	.686	
Age	-0.021	0.023	-0.925	.355	-0.011	0.025	-0.437	.662	
Sex	-0.009	0.029	-0.309	.757	0.007	0.026	0.279	.780	
Race	-0.015	0.022	-0.678	.498	-0.026	0.022	-1.182	.237	
		Outcome: P	ubertal stage		Outcome: Pubertal stage				
Family Environment ^f	-0.136	0.022	-6.189	<.001	-0.136	0.022	-6.189	<.001	
Age	0.231	0.021	11.159	<.001	0.231	0.021	11.159	<.001	
Sex	-0.504	0.016	-30.950	<.001	-0.504	0.016	-30.950	<.001	
Race	0.136	0.020	6.686	<.001	0.136	0.020	6.686	<.001	

Note: $^{\infty}$ = direct effect; $^{+}$ = indirect effect Step 2; f = indirect effect Step 1; SOMM = somato-motor mouth network.

TableS17. Mediation model parameters—Somatomotor-mouth network-amygdala connectivity in girls

		SOMM-left amygdala			SOMM-right amygdala			
	β	S.E.	β/S.E.	р	β	S.E.	β /S.E.	p
Family Environment [∞]	0.064	0.031	2.099	.036	0.026	0.035	0.0753	.452
Pubertal stage ⁺	0.045	0.041	1.094	.274	-0. 005	0.037	-0.146	.884
Age	-0.004	0.031	-0.113	.910	0.010	0.035	0.281	.779
Race	-0.071	0.032	-2.208	.027	-0.052	0.032	-1.592	.111
		Outcome: Pi	ubertal stage		Outcome: Pubertal stage			
Family Environment ^f	-0.171	0.034	-5.058	<.001	-0.171	0.034	-5.058	<.001
Age	0.300	0.030	9.991	.000	0.300	0.030	9.991	.000
Race	0.111	0.034	3.295	<.001	0.111	0.034	3.295	<.001

Note: [∞] = direct effect; ⁺ = indirect effect Step 2; ^f = indirect effect Step 1; SOMM = somato-motor mouth network.

TableS20. Mediation model parameters -- Somatomotor-mouth network-amygdala connectivity in boys

		SOMM-left amygdala				SOMM-right amygdala			
	β	S.E.	β/S.E.	р	β	S.E.	β/S.E.	р	
Family Environment [∞]	0.056	0.032	1.787	.074	-0.046	0.031	-1.474	.141	
Pubertal stage ⁺	-0.043	0.045	-0.972	.331	-0.023	0.044	-0.528	.597	
Age	-0.031	0.031	-1.003	.316	-0.024	0.034	-0.698	.485	
Race	0.034	0.029	1.160	.246	-0.002	0.031	-0.060	.952	
		Outcome: Pubertal stage				Outcome: Pubertal stage			
Family Environment ^f	-0.142	0.036	-3.960	<.001	-0.142	0.036	-3.960	<.001	
Age	0.223	0.035	6.289	<.001	0.223	0.035	6.289	<.001	
Race	0.212	0.031	6.846	<.001	0.212	0.031	6.846	<.001	

Note: [∞] = direct effect; ⁺ = indirect effect Step 2; ^f = indirect effect Step 1.SOMM = somato-motor mouth network.

Table3. Correlation between MRI measures

	ACC CA	ACC FA	Amygdala SV	CON-l amygdala FC	CON-r amygdala FC
ACC CT	082	072	.067	.003	.036
ACC CA		257	017	.033	.013
ACC FA			049	012	038
Amygdala SV				058	033
CON amygdala FC					.586

Note: All measures are residualized for data collection site. Gray matter measures were further residualized for total brain volume. ACC = anterior cingulate cortex; CT = cortical thickness; CA = cortical area; FA = fractional anisotropy; SC = subcortical volume; CON = cingulo-opercular network; I = left; r = right; FC = functional connectivity.

TableS1. Correlations among brain measures of motor processing

	Precentral CA	Precentral FA	SOMM-L Amygdala FC	SOMMR Amygdala FC
Precentral CT	423	.204	.070	.023
Precentral CA		114	046	011
Precentral FA			031	.033
SOMM-L Amygdala FC				197

Note: SOMM = somatomotor-mouth network; FC = functional connectivity; CT = cortical thickness; CA = cortical area; FA = fractional anisotropy.

Reference

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