Supplementary Results

Cross-sectional correlations.

Table 1 reports cross-sectional correlations between age, testosterone, and PDS for females (highlighted in pink) and males (highlighted in blue) for each time point.

Table 1.

Timepoint 1									
	PDS	Testosterone	Age						
PDS		0.45**	0.44**						
Testosterone	0.40*		.43**						
Age	0.25	0.51**							
Timepoint 2									
	PDS	Testosterone	Age						
PDS		.39*	.60**						
Testosterone	.55***		.42**						
Age	.39*	.41*							
Timepoint 3									
	PDS	Testosterone	Age						
PDS		0.28	.56***						
Testosterone	0.61***		0.25						
Age	0.69***	.52***							

* (females=pink, males=blue)

Inclusion of additional covariates in longitudinal analyses

Supplementary analyses confirmed that all significant findings remained unchanged when we included additional covariates related to motion, "not sure" responses, and total number of trials. Initial values at T1 and change since T1 in each of these variables was added to the final models selected for hippocampus (Table 2) and DLPFC (Table 3).

Motion. Motion was measured as the number of repaired volumes based on ArtRepair (>1mm motion or >2% signal change) for each participant. Children's average number repaired volumes was M(SD)=14.22(11.92) at T1, M(SD)=10.72(10.03) at T2, and M(SD)=7.87(8.47) at T3 corresponding to 4 to 7% of volumes. The number of repaired volumes was negatively related to age at T1 (*r*=-.25, *p*=.02) and T2 (*r*=-.27, *p*=.01), and marginally at T3 (*r*=-.20, *p*=.08). To account for these developmental effects, we added the number of repaired volumes at T1 and change in repaired volumes since T1 in our final models of hippocampal and DLPFC activation. Repaired volumes at T1 was not a significant predictor of activation (*ps*>.29), but change in number of repaired volumes was significant when added to the final hippocampal models (PDS model: b=-.01, *p*=.003, Testosterone model: b=-.01, *p*=.003), indicating that children whose number of repaired volumes was significant for the DLPFC models (*ps*>.50). Importantly, all our main findings still held for the both the hippocampal (PDS model: main effect of region b=.02, *p*=.04; time² b=1.32 *p*<.001; T1 age x Change in PDS b=.15, *p*<.001; Testosterone model: T1 age x Gender x Change in Testosterone, b=-.01, *p*=.03).

Not Sure Rates. "Not sure" rates were measured as the number of "not sure" responses divided by the total number of hits. Children's "not sure" response rates were M(SD)=.26(.19) at T1, M(SD)=.17(.13) at T2, and M(SD)=.14(.12) at T3. "Not sure" rates negatively associated with age at T1 (*r*=-.33, *p*=.001) and T2 (*r*=-.25, *p*=.02), but not T3 (*r*=-.15, *p*=.19). To account for these developmental effects, we added "not sure" rates at T1 and change in "not sure" rates since T1 in our final models. Neither "not sure" rate at T1 or change in "not sure" rates were significant predictors when added to the final hippocampal models *p*>.49; when added to the final DLPFC model, the change in "not sure" rates marginally predicted DLPFC activation (PDS model: b=-.41, *p*=.06, Testosterone model: b=-.37, *p*=.07). However, all our main findings remain significant for the both the hippocampal (PDS model: main effect of region b=.02, *p*=.04; time² b=1.27 *p*<.001; T1 age x Change in PDS b=.14, *p*<.001; Testosterone model: T1 age x Gender x Change in Testosterone, b=-.01, *p*<.001) and DLPFC model: time² b=1.26 *p*<.001; Testosterone model: T1 age x Gender x Change in Testosterone, b=-.01, *p*<.001)

Total Number of Trials. The total number of trials was calculated as the sum of correct and incorrect itemcontext trials that were included for fMRI activation estimates. Total trial number was M(SD)=102(24) at T1, M(SD)=111(25) at T2, and M(SD)=119(18) at T3. Total trial number was marginally positively associated with age at T1 (r=.19, p=.07), significantly at T2 (r=.27, p=.02), and not significantly at T3 (r=.12, p=.30). To account for these developmental effects, we added total trial number at T1 and change in total trial number since T1 in our final models. Total trial number at T1 was not a significant predictor of activation (ps>.27), but change in number of total trials was significant when added to the final hippocampal models (PDS model: b=.003, p<.001, Testosterone model: b=.004, p<.001), indicating that children whose number of total trials increased over time exhibited higher hippocampal activation. Initial total trial number was not significant for the DLPFC models (ps>.20) but change in number of total trials was marginal for the model including PDS (b=.002, p=.08) and significant for the model including testosterone (b=.004, p<.001). Importantly, our main results remain significant for the hippocampal model (PDS model: main effect of region b=.02, p=.04; time² b=1.34 p<.001; T1 age x Change in PDS b=.11, p=.006; Testosterone model: T1 age x Gender x Change in Testosterone, b=-.01, p < .001). Our main results also remain largely the same for the DLPFC models (PDS Model: time² b=2.12 p<.001; Testosterone model: T1 age x Gender x Change in Testosterone, b=-.01, p=.058), although we note the 3-way interaction was marginal at p=.058.

Hippocampus									
								. 95%	6 CI
	SD	b	SE	df	t	р		Lower	Upper
PDS Bandom Effect									
	0.284							0 247	0 327
Time	0.204							0.247	0.327
Fixed Effect	0.240							0.201	0.200
Intercept		-0.033	0.039	1291	-0.84	0.40		-0.111	0.044
Initial Age		-0.047	0.025	117	-1.93	0.06		-0.096	0.001
region:Body		0.024	0.012	1291	2.02	0.04	*	0.001	0.046
region:Head		0.018	0.012	1291	1.58	0.12		-0.005	0.041
Time		-4.176	1.245	1291	-3.35	<.001	***	-6.619	-1.733
Time ²		1.362	0.346	1291	3.94	<.001	***	0.683	2.040
Gender		0.001	0.051	117	0.02	0.99		-0.100	0.102
Hemisphere		0.001	0.010	1291	0.14	0.89		-0.017	0.020
Initial PDS		0.010	0.057	117	0.17	0.86		-0.103	0.122
Change in PDS		0.131	0.043	1291	3.04	0.002	**	0.046	0.216
Initial Repaired Scans		-0.002	0.002	117	-0.86	0.39		-0.006	0.002
Change in Repaired Scans		-0.003	0.002	1291	-1.50	0.13		-0.007	0.001
Initial Not Sure Rate		-0.008	0.130	117	-0.06	0.95		-0.266	0.251
Change in Not Sure Rate		0.025	0.134	1291	0.19	0.85		-0.238	0.289
Initial Trial Count		0.000	0.001	117	-0.12	0.90		-0.002	0.002
Change in Trial Count		0.002	0.001	1291	3.28	0.001	**	0.001	0.004
Initial Age*Change in PDS		0.119	0.041	1291	2.94	0.003	**	0.039	0.198
Testosterone									
Random Effect									
Intercept	0.291							0.253	0.335
Time	0.287							0.239	0.345
Fixed Effect									
Intercept		0.013	0.039	1187	0.35	0.73		-0.063	0.090
Initial Age		0.012	0.036	111	0.33	0.74		-0.059	0.083
region:Body		0.017	0.011	1187	1.57	0.12		-0.004	0.039
region:Head		0.014	0.011	1187	1.25	0.21		-0.008	0.036
Time		-2.461	1.257	1187	-1.96	0.05	*	-4.928	0.006
Lime		0.921	0.331	1187	2.79	0.01	**	0.272	1.570
Gender		-0.037	0.050	111	-0.74	0.46		-0.136	0.062
Hemisphere		0.004	0.009	118/	0.47	0.64		-0.014	0.022
		0.000	0.002	111	-0.16	0.87	*	-0.004	0.003
Unange in Testosterone		-0.003	0.002	1187	-2.00	0.05		-0.006	0.000
Change in Repaired Scalls		-0.003	0.002	1107	-1.24	0.22		-0.000	0.002
Unarige in Repaired Scans		-0.002	0.002	110/	-1.32 1.02	0.19		-0.000	0.001
Initial Not Sure Rate Change in Not Sure Pete		0.142	0.130	1187	1.03	0.31		-0.132	0.410
Initial Trial Count		-0.001	0.129	111	-1 11	0.13		-0.000	0.440
Change in Trial Count		0.001	0.001	1187	6 15	< 0.21	***	0.003	0.001
Gender * Change in Testosterone		0.004	0.001	1187	1 78	0.08		0.000	0.000
Initial Age*Change in Testosterone		0.004	0.002	1187	5.58	< 001	***	0.005	0.000
Initial Age * Gender		-0.025	0.046	111	-0.54	0.59		-0.115	0.066
Initial Age* Gender * Change in Testosterone		-0.007	0.002	1187	-3.29	<.001	***	-0.012	-0.003

			DLPFC							
									95%	6 CI
		SD	b	SE	df	t	р		Lower	Upper
PDS										
Random Effect										
li li	ntercept	0.339							0.291	0.396
	Ime	0.226							0.180	0.284
Fixed Effect			0.040	0.050	224	0.00	0.74		0 4 0 0	0.004
 			-0.018	0.056	334	-0.33	0.74		-0.128	0.091
111			-0.004	0.033	117	-0.11	0.92		-0.009	0.002
	Time ²		-1.00Z	0.090	224	-1.17	0.24	***	-2.010	0.711
	Condor		2.123	0.301	117	2.00	<.001 0.04	*	0.295	2.000
Hem	isphore		-0.147	0.070	334	-2.11	0.04		-0.205	-0.009
Init			-0.003	0.019	117	-0.23	0.00		-0.043	0.033
Change	in PDS		0 106	0.077	334	1 54	0.14		-0.200	0.033
Initial Repaired	d Scans		-0.003	0.003	117	-0.92	0.36		-0.009	0.003
Change in Repaired	d Scans		0.000	0.003	334	1 41	0.00		-0.002	0.000
Initial Not Su	ire Rate		-0 127	0.000	117	-0.71	0.48		-0.480	0.226
Change in Not Su	ire Rate		-0.460	0.216	334	-2 13	0.03	*	-0.885	-0.034
Initial Tria	al Count		-0.002	0.001	117	-1.64	0.10		-0.005	0.000
Change in Tria	al Count		0.002	0.001	334	1.95	0.05	*	0.000	0.004
Initial Age*Change	in PDS		0.058	0.055	334	1.04	0.30		-0.051	0.166
Testosterone										
Random Effect										
lı lı	ntercept	0.349							0.300	0.407
	Time	0.218							0.170	0.278
Fixed Effect										
lı lı	ntercept		0.012	0.050	304	0.24	0.81		-0.086	0.110
Ini	itial Age		-0.001	0.047	111	-0.02	0.98		-0.095	0.093
	Time		-0.611	0.703	304	-0.87	0.39		-1.994	0.772
	l ime ⁻		1.643	0.347	304	4.73	<.001	***	0.960	2.326
	Gender		-0.152	0.067	111	-2.29	0.02	*	-0.284	-0.020
Hem Initial Tests	lisphere		0.001	0.019	304	0.06	0.95		-0.036	0.038
Initial Lesto	sterone		-0.003	0.003	111	-1.30	0.20		-0.008	0.002
	sterone		-0.004	0.003	304	-1.47	0.14		-0.009	0.001
			-0.001	0.003	111	-0.37	0.72		-0.007	0.005
Change in Repaired	u Scans		0.005	0.003	304	1.09	0.09		-0.001	0.010
Initial NOL SU Chango in Not Su			-0.022	0.100	111 204	-0.1Z	0.90		-0.309	0.340
Change in NOL SU			-0.320	0.207	111	0.71	0.12		-0.732	0.001
Initial Irial Count Change in Trial Count			0.001	0.001	304	-0.71	0.40 < 001	***	-0.004 0.002	0.002
Conder * Change in Tratectorer			0.004	0.001	304	0.14 0 11	001	*	0.002	0.000
Gender " Change in Testosterone			11111		. 11/4	2.14	0.00		0.001	0.014
	sterone		0.006	0.000	304	2.23	0.03	*	0.001	0.011
Initial Age Change in Testo Initial Age *	sterone		0.006	0.003	304 111	2.23	0.03	*	0.001	0.011

Table 3. Longitudinal Model Results for DLPFC Item-Context Association Activation Including Additional Covariates

Supplementary Figures



Figure 1. Longitudinal results for hippocampal item-context association activation. A.) depicts the change in puberty score by initial age interaction, and B.) depicts the change in testosterone by initial age by gender interaction. Individual values are plotted by points and individual participants are connected by horizontal lines. Participants were separated into three age groups based on initial age (younger <8.7 yrs, middle >8.7 and <10.2, older >10.2 yrs). The plotted regression lines were estimated from the full longitudinal models reported in the manuscript for three specific initial ages (8.7 years, 10.2 years, and 11.7 years).



Figure 2. Longitudinal results for DLPFC item-context association activation. A.) depicts the change in puberty score by initial age interaction, and B.) depicts the change in testosterone by initial age by gender interaction. Individual values are plotted by points and individual participants are connected by horizontal lines. Participants were separated into three age groups based on initial age (younger <8.7 yrs, middle >8.7 and <10.2, older >10.2 yrs). The plotted regression lines were estimated from the full longitudinal models reported in the manuscript for three specific initial ages (8.7 years, 10.2 years, and 11.7 years).