



Supplementary Materials

Table S1. Genotype effects on brain curvature.

Gene	SNPs	Age-adjusted/FDR									
		Left Hemisphere					Right Hemisphere				
		Lateral Orbitofrontal Gyrus	Medial Orbitofrontal Gyrus	Parahippocampal Gyrus	Posterior Cingulate Gyrus	Superior Temporal Gyrus	Lateral Orbitofrontal Gyrus	Medial Orbitofrontal Gyrus	Parahippocampal Gyrus	Posterior Cingulate Gyrus	Superior Temporal Gyrus
<i>OXTR</i>	rs1042778	0.850/0.982	0.033/0.414	0.899/0.981	0.680/0.765	0.080/0.240	0.098/0.528	<0.001/<0.001	0.394/0.974	0.273/0.734	0.005/0.043
<i>OXTR</i>	rs2268493	0.399/0.857	0.287/0.575	0.749/0.981	0.013/0.034	0.045/0.230	0.100/0.528	0.035/0.079	0.400/0.974	0.260/0.734	0.017/0.057
<i>OXTR</i>	rs53576	0.564/0.890	0.224/0.575	0.547/0.923	0.015/0.034	0.027/0.230	0.175/0.528	0.016/0.075	0.711/0.990	0.260/0.734	0.002/0.041
<i>FADS2</i>	rs174570	0.508/0.857	0.329/0.575	0.819/0.981	0.011/0.034	0.051/0.230	0.193/0.528	0.021/0.075	0.799/0.974	0.386/0.734	0.008/0.043
<i>FADS2</i>	rs174576	0.743/0.912	0.256/0.575	0.911/0.981	<0.001/<0.001	0.023/0.230	0.187/0.528	0.023/0.075	0.736/0.990	0.473/0.749	0.007/0.043
<i>FADS2</i>	rs498793	0.668/0.890	0.751/0.819	0.429/0.827	0.015/0.034	0.041/0.230	0.036/0.528	0.054/0.104	0.933/0.990	0.181/0.734	0.003/0.041
<i>COMT</i>	rs740603	0.488/0.857	0.588/0.756	0.703/0.981	0.012/0.034	0.143/0.322	0.349/0.628	0.002/0.027	0.905/0.990	0.384/0.734	0.010/0.045
<i>COMT</i>	rs165774	0.983/0.983	0.121/0.575	0.895/0.981	<0.001/<0.001	0.063/0.240	0.094/0.528	0.024/0.075	0.760/0.990	0.173/0.734	0.015/0.057
<i>COMT</i>	rs174696	0.429/0.857	0.555/0.749	0.722/0.981	0.047/0.098	0.101/0.273	0.349/0.628	0.024/0.075	0.136/0.974	0.324/0.734	0.037/0.111

SNP, single-nucleotide polymorphism; *OXTR*, oxytocin receptor; *FADS2*, fatty acid desaturase 2; *COMT*, catechol-o-methyltransferase; FDR, false discovery rate.

Table S2. Genotype effects on brain volume.

Gene	SNPs	Age-adjusted/FDR				
		Total Gray Matter Volume	Left Hemisphere Cortical Gray Matter Volume	Right Hemisphere Cortical Gray Matter Volume	Left Hemisphere Cerebral White Matter Volume	Right Hemisphere Cerebral White Matter Volume
		<i>OXTR</i>	rs1042778	0.748/0.805	0.721/0.972	0.768/0.839
<i>OXTR</i>	rs2268493	0.730/0.805	0.972/0.972	0.548/0.748	0.901/0.992	0.816/0.995
<i>OXTR</i>	rs53576	0.805/0.805	0.821/0.972	0.839/0.839	0.992/0.992	0.995/0.995
<i>FADS2</i>	rs174570	0.698/0.805	0.876/0.972	0.582/0.748	0.682/0.992	0.825/0.995
<i>FADS2</i>	rs174576	0.613/0.805	0.750/0.972	0.513/0.748	0.954/0.992	0.675/0.995
<i>FADS2</i>	rs498793	0.196/0.805	0.428/0.972	0.267/0.748	0.321/0.992	0.125/0.995
<i>COMT</i>	rs740603	0.377/0.805	0.369/0.972	0.388/0.748	0.760/0.992	0.809/0.995
<i>COMT</i>	rs165774	0.583/0.805	0.772/0.972	0.492/0.748	0.413/0.992	0.488/0.995
<i>COMT</i>	rs174696	0.314/0.805	0.265/0.972	0.164/0.748	0.689/0.992	0.452/0.995

SNP, single-nucleotide polymorphism; *OXTR*, oxytocin receptor; *FADS2*, fatty acid desaturase 2; *COMT*, catechol-o-methyltransferase; FDR, false discovery rate.

Table S3. Genotype effects on brain volume in the preterm and control groups controlling for age at scan.

Gene	SNPs	Age-adjusted/P _{FDR}				
		Total Gray Matter Volume	Left Hemisphere Cortical Gray Matter Volume	Right Hemisphere Cortical Gray Matter Volume	Left Hemisphere Cerebral White Matter Volume	Right Hemisphere Cerebral White Matter Volume
		<i>OXTR</i>	rs1042778	0.113/0.848	0.494/0.943	0.271/0.933
<i>OXTR</i>	rs2268493	0.272/0.848	0.835/0.943	0.575/0.933	0.083/0.330	0.079/0.304
<i>OXTR</i>	rs53576	0.151/0.848	0.705/0.943	0.284/0.933	0.072/0.330	0.064/0.304
<i>FADS2</i>	rs174570	0.188/0.848	0.751/0.943	0.349/0.943	0.110/0.330	0.072/0.304
<i>FADS2</i>	rs174576	0.189/0.848	0.632/0.943	0.364/0.943	0.109/0.330	0.068/0.304
<i>FADS2</i>	rs498793	0.201/0.848	0.697/0.943	0.336/0.933	0.083/0.330	0.079/0.304
<i>COMT</i>	rs740603	0.332/0.848	0.908/0.943	0.476/0.933	0.158/0.388	0.140/0.378
<i>COMT</i>	rs165774	0.073/0.848	0.362/0.943	0.135/0.933	0.078/0.330	0.068/0.304
<i>COMT</i>	rs174696	0.323/0.848	0.898/0.943	0.466/0.933	0.089/0.330	0.090/0.304

SNP, single-nucleotide polymorphism; *OXTR*, oxytocin receptor; *FADS2*, fatty acid desaturase 2; *COMT*, catechol-o-methyltransferase; FDR, false discovery rate.

Table S4. Genotype effects on brain thickness.

Gene	SNPs	Age-adjusted/FDR									
		Left Hemisphere					Right Hemisphere				
		Lateral Orbitofrontal	Medial Orbitofrontal	Parahippocampal	Posterior Cingulate	Superior Temporal	Lateral Orbitofrontal	Medial Orbitofrontal	Parahippocampal	Posterior Cingulate	Superior Temporal
<i>OXTR</i>	rs1042778	0.210/0.962	0.031/0.155	0.296/0.920	0.371/0.669	0.113/0.917	0.253/0.885	0.869/0.869	0.497/0.688	0.682/0.682	0.044/0.440
<i>OXTR</i>	rs2268493	0.855/0.962	0.989/0.989	0.368/0.920	0.468/0.669	0.751/0.917	0.826/0.885	0.222/0.740	0.529/0.688	0.299/0.682	0.315/0.525
<i>OXTR</i>	rs53576	0.996/0.996	0.512/0.853	0.332/0.920	0.343/0.669	0.569/0.917	0.885/0.885	0.456/0.869	0.799/0.888	0.302/0.682	0.957/0.957
<i>FADS2</i>	rs174570	0.851/0.962	0.839/0.989	0.789/0.957	0.920/0.920	0.558/0.917	0.749/0.885	0.832/0.869	0.095/0.605	0.575/0.682	0.296/0.525
<i>FADS2</i>	rs174576	0.610/0.962	0.471/0.853	0.697/0.957	0.764/0.849	0.287/0.917	0.787/0.885	0.768/0.869	0.121/0.605	0.540/0.682	0.215/0.525
<i>FADS2</i>	rs498793	0.491/0.962	0.932/0.989	0.663/0.957	0.310/0.669	0.780/0.917	0.411/0.885	0.681/0.869	0.499/0.688	0.469/0.682	0.248/0.525
<i>AP0E</i>	rs405509	0.833/0.962	0.958/0.989	0.171/0.920	0.055/0.550	0.536/0.917	0.441/0.885	0.704/0.869	0.550/0.688	0.063/0.630	0.857/0.952
<i>COMT</i>	rs740603	0.341/0.962	0.380/0.853	0.979/0.979	0.609/0.761	0.903/0.917	0.482/0.885	0.106/0.555	0.266/0.688	0.269/0.682	0.416/0.594
<i>COMT</i>	rs165774	0.682/0.962	0.463/0.853	0.861/0.957	0.444/0.669	0.917/0.917	0.831/0.885	0.427/0.869	0.371/0.688	0.675/0.682	0.787/0.952
<i>COMT</i>	rs174696	0.055/0.962	0.031/0.155	0.723/0.957	0.117/0.585	0.521/0.917	0.041/0.410	0.111/0.555	0.940/0.940	0.420/0.682	0.146/0.525

SNP, single-nucleotide polymorphism; *OXTR*, oxytocin receptor; *FADS2*, fatty acid desaturase 2; *COMT*, catechol-o-methyltransferase; FDR, false discovery rate.

Table S5. Genotype effects on brain curvature in the preterm and control groups controlling for age at scan.

Gene	SNPs	Age-adjusted/FDR									
		Left Hemisphere					Right Hemisphere				
		Lateral Orbitofrontal Gyrus	Medial Orbitofrontal Gyrus	Parahippocampal Gyrus	Posterior Cingulate Gyrus	Superior Temporal Gyrus	Lateral Orbitofrontal Gyrus	Medial Orbitofrontal Gyrus	Parahippocampal Gyrus	Posterior Cingulate Gyrus	Superior Temporal Gyrus
<i>OXTR</i>	rs1042778	0.259/0.957	0.044/0.531	0.276/0.827	0.387/0.992	0.112/0.833	0.317/0.914	0.980/0.981	0.644/0.820	0.788/0.851	0.070/0.545
<i>OXTR</i>	rs2268493	0.990/0.994	0.581/0.977	0.062/0.698	0.350/0.992	0.270/0.833	0.880/0.968	0.911/0.981	0.432/0.807	0.295/0.556	0.456/0.663
<i>OXTR</i>	rs53576	0.427/0.957	0.712/0.977	0.139/0.698	0.948/0.992	0.676/0.879	0.417/0.914	0.723/0.981	0.296/0.807	0.303/0.556	0.150/0.545
<i>FADS2</i>	rs174570	0.695/0.994	0.673/0.977	0.409/0.893	0.961/0.992	0.520/0.877	0.486/0.914	0.941/0.981	0.482/0.807	0.412/0.654	0.198/0.545
<i>FADS2</i>	rs174576	0.494/0.957	0.881/0.977	0.562/0.972	0.992/0.992	0.906/0.986	0.385/0.914	0.871/0.981	0.614/0.820	0.615/0.751	0.232/0.545
<i>FADS2</i>	rs498793	0.994/0.994	0.631/0.977	0.329/0.827	0.694/0.992	0.528/0.877	0.968/0.968	0.959/0.981	0.408/0.807	0.438/0.657	0.241/0.545
<i>COMT</i>	rs740603	0.906/0.994	0.453/0.977	0.302/0.827	0.888/0.992	0.395/0.833	0.491/0.914	0.789/0.981	0.368/0.807	0.223/0.556	0.263/0.545
<i>COMT</i>	rs165774	0.601/0.994	0.882/0.977	0.648/0.972	0.822/0.992	0.986/0.986	0.338/0.914	0.639/0.981	0.546/0.819	0.513/0.693	0.182/0.545
<i>COMT</i>	rs174696	0.897/0.994	0.327/0.977	0.273/0.827	0.724/0.992	0.313/0.833	0.876/0.968	0.877/0.981	0.458/0.807	0.103/0.556	0.220/0.545

SNP, single-nucleotide polymorphism; *OXTR*, oxytocin receptor; *FADS2*, fatty acid desaturase 2; *COMT*, catechol-o-methyltransferase; FDR, false discovery rate.

Table S6. Significance of minor allele frequencies of SNPs and GA.

		Early Preterm (n = 9)	Late Preterm (n = 11)	Control (n = 6)	Pearson χ^2
<i>OXTR</i>	rs1042778	0	1	1	0.409
<i>OXTR</i>	rs2268493	3	3	0	0.571
<i>OXTR</i>	rs53576	5	7	3	0.650
<i>FADS2</i>	rs174570	5	7	4	0.511
<i>FADS2</i>	rs174576	5	7	5	0.569
<i>FADS2</i>	rs498793	5	1	0	0.102
<i>COMT</i>	rs740603	6	8	3	0.729
<i>COMT</i>	rs165774	2	3	1	0.985
<i>COMT</i>	rs174696	6	9	3	0.165

SNP, single-nucleotide polymorphism; *OXTR*, oxytocin receptor; *FADS2*, fatty acid desaturase 2; *COMT*, catechol-o-methyltransferase; GA, gestational age.

Table S7. The sex differences of cerebral development.

Characteristics	Male (n = 15)	Female (n = 11)	p-Value
Volume			
Total gray matter volume	107825.85 ± 13431.40	106143.82 ± 16315.27	0.776
Left hemisphere cortical gray matter volume	36660.81 ± 4637.06	36482.69 ± 6379.82	0.935
Right hemisphere cortical gray matter volume	36231.77 ± 4603.25	34957.95 ± 5693.85	0.534
Left hemisphere cerebral white matter volume	73300.92 ± 10143.10	65353.17 ± 8864.55	0.048
Right hemisphere cerebral white matter volume	69638.25 ± 7685.66	63059.96 ± 8244.22	0.047
Thickness			
Left lateral orbitofrontal gyrus	1.08 ± 0.14	1.18 ± 0.23	0.209
Right lateral orbitofrontal gyrus	1.07 ± 0.16	1.09 ± 0.17	0.708
Left medial orbitofrontal gyrus	1.13 ± 0.12	1.19 ± 0.19	0.371
Right medial orbitofrontal gyrus	1.18 ± 0.11	1.20 ± 0.23	0.749
Left parahippocampal gyrus	1.05 ± 0.16	1.05 ± 0.17	0.979
Right parahippocampal gyrus	1.05 ± 0.14	1.01 ± 0.27	0.671
Left posterior cingulate gyrus	1.33 ± 0.13	1.33 ± 0.15	0.909
Right posterior cingulate gyrus	1.32 ± 0.13	1.37 ± 0.10	0.248
Left superior temporal gyrus	1.40 ± 0.11	1.41 ± 0.19	0.923
Right superior temporal gyrus	1.42 ± 0.11	1.40 ± 0.21	0.814
Curvature			
Left lateral orbitofrontal gyrus	1.95 ± 0.57	2.08 ± 0.68	0.595
Right lateral orbitofrontal gyrus	2.11 ± 0.59	2.10 ± 0.68	0.962
Left medial orbitofrontal gyrus	1.56 ± 0.41	1.69 ± 0.62	0.531
Right medial orbitofrontal gyrus	1.68 ± 0.56	1.49 ± 0.46	0.367
Left parahippocampal gyrus	0.50 ± 0.29	0.62 ± 0.20	0.267
Right parahippocampal gyrus	0.54 ± 0.44	0.35 ± 0.24	0.204
Left posterior cingulate gyrus	0.92 ± 0.44	0.74 ± 0.27	0.220
Right posterior cingulate gyrus	0.90 ± 0.28	0.86 ± 0.41	0.753
Left superior temporal gyrus	5.27 ± 0.99	4.87 ± 0.80	0.284
Right superior temporal gyrus	4.53 ± 0.99	4.38 ± 0.88	0.698