

Additional file 2

Table S2 Results of regression analyses of *OXTR* coding variants predicting gestational age (GA) in Danish case and control mothers¹

Variant	GA in weeks ²		GA as a binary trait ³	
	<i>n</i> ⁴	<i>P</i>	<i>n</i> (case, control) ⁴	<i>P</i>
rs237913 ⁵	1591	0.08	691, 900	0.11
rs237911	1598	0.22	696, 902	0.44
A11T	1602	0.53	697, 905	0.98
V45L	1602	0.28	696, 906	0.97
V211L	1590	0.39	698, 892	0.86
rs4686302 (A218T)	1592	0.57	701, 891	0.54
rs237902 (N230N)	1586	0.26	697, 889	0.35
rs61740241 (A238T)	1588	0.46	698, 890	0.26
A247A	1587	0.79	696, 891	0.44
rs151141371 (G252A)	1588	0.35	698, 890	0.25
rs144814761 (V281M)	1588	0.38	697, 891	0.98

¹The analysis was performed for all common and rare variants identified by sequencing.

²GA was considered as a continuous variable.

³GA was considered as a categorical variable (preterm, ≤ 36 weeks; and term, 40 weeks).

⁴Number of individuals successfully genotyped and included in the analysis.

⁵The best genetic effect model.