

S2 Table. Polygenic Transmission of nsCL/P genetic risk variants in independent European and Asian trios

P-value Inclusion Threshold	European Trios (N = 2209)			Asian Trios (N = 2593)		
	Number of SNPs	Beta (95% C.I.) ¹	P Value	Number of SNPs	Beta (95% C.I.) ¹	P Value
0.000001	4	0.30 (0.23, 0.37)	1.8x 10 ⁻¹⁶	10	0.12 (0.05, 0.19)	8.0x 10 ⁻⁴
0.000005	13	0.30 (0.23, 0.37)	1.1x 10 ⁻¹⁶	18	0.16 (0.09, 0.23)	7.0x 10 ⁻⁶
0.00001	17	0.32 (0.25, 0.39)	3.5x 10⁻¹⁸	23	0.19 (0.12, 0.25)	1.7x 10⁻⁷
0.00005	44	0.25 (0.18, 0.33)	1.0x 10 ⁻¹¹	53	0.16 (0.09, 0.13)	1.1x 10 ⁻⁵
0.0001	69	0.24 (0.17, 0.31)	2.1x 10 ⁻¹¹	81	0.12 (0.05, 0.19)	0.001
0.0005	222	0.17 (0.10, 0.23)	2.7x 10 ⁻⁶	244	0.06 (0.00, 0.13)	0.056
0.001	405	0.15 (0.08, 0.22)	4.6x 10 ⁻⁵	437	0.08 (0.01, 0.14)	0.020
0.005	1,626	0.19 (0.12, 0.26)	5.3x 10 ⁻⁸	1793	0.12 (0.05, 0.18)	3.5x10 ⁻⁴
0.01	3,002	0.16 (0.09, 0.23)	9.2x 10 ⁻⁶	3,334	0.10 (0.04, 0.17)	0.002
0.05	11,400	0.16 (0.10, 0.23)	1.7x 10 ⁻⁶	13,222	0.09 (0.02, 0.15)	0.009
0.1	20,133	0.16 (0.10, 0.23)	2.2x 10 ⁻⁶	23,421	0.06 (-0.01, 0.12)	0.096
0.5	64,727	0.17 (0.10, 0.24)	1.1x 10 ⁻⁶	74,832	0.03 (-0.03, 0.10)	0.31
1	92,527	0.16 (0.09, 0.23)	2.6x 10 ⁻⁶	107,809	0.03 (-0.04, 0.10)	0.38

¹ Standardised difference in genetic score between parents and offspring