

Supplementary table 1. CNVs of participants' children

CNV	N
11q14.1 deletion	2
13q12.2 duplication	1
15q11.2 deletion	22
15q11.2 duplication	1
15q13.2 duplication	2
15q13.3 deletion	11
15q13.3 duplication	10
16p11.2 deletion (and distal deletion)	37
16p11.2 duplication	12
16p12.2 deletion	2
16p13.11 deletion	4
16p13.11 duplication	2
1q21.1 deletion	13
1q21.1 duplication	17
22q11.2 deletion	80
22q11.2 duplication	19
22q12.3 duplication	1
22q13.3 deletion	2
2p16.3 deletion	1
3q29 deletion	4
4q28.3	1
7p22.1 deletion	1
7q11.23 duplication	1
8q12.3 deletion	1
9q34.3 deletion	10
17q23 duplication	1
SHANK3 deletion	1

Xp21.1	1
XXXY	1
NRXN1	12
We note here the primary CNV for which the family was recruited. 12% of our sample had additional CNVs.	