

Type	Proband	Sibling
Silent	214	124
Missense1	158	114
Missense2	78	55
Missense3	319	147
Nonsense	51	16
Splice	18	3
Total SNV	842	461
Frameshift	54	15

Table 2: The statistics of the *de novo* mutations in autism probands and unaffected siblings. The missense labels are based on predictions from PolyPhen2. Missense1-3 correspond to “benign”, “possibly damaging” and “probably damaging” mutations, respectively. The last row is the counts of frameshift indels.