Supplementary Material, Table S1 Study samples and the number of markers analyzed in the follow-up and fine-map stages per locus. Abbreviations: Ext ped=extended pedigree, AUT=autism study sample, AS=Asperger syndrome study sample.

| Locus | Follow-up microsatellites | Study samples | Fine-map SNPs | Study samples |
|-------|---------------------------|---------------|------------------|----------------------|
| 1q23 | 23 ^a | Ext ped | 31 | Ext ped, AUT, AS |
| 15q12 | 7 ^b | Ext ped | 41 | Ext ped, AUT, AS |
| 19p13 | 7 ^c | Ext ped | 80 | Ext ped ^d |
| 14q11 | 1 | Ext ped | X | X |
| 13q12 | 2 | Ext ped | X | X |
| 8q23 | 1 | Ext ped | X | X |
| 6р | 1 | Ext ped | X | X |
| 5q | 2 | Ext ped | X | X |

^afrom D1S1631 to D1S1589; region 55.2 cM ^bfrom D1SS97 to D15S165; region 10.4 cM

^cfrom D19S883 to D19S873; region 24.1 cM

^dmarkers for *TLE6*, *TLE2* and *AES* genes were analyzed also in the AUT and AS study samples