

CORRIGENDUM

AH11, a pivotal neurodevelopmental gene, and *C6orf217* are associated with susceptibility to schizophrenia

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In the version of the article initially published, three minor alleles (MA) and the resulting over-transmitted alleles (OTA) were switched in Table 1 as well as in the online Supplementary Table S1. For SNP no. 32 (rs6902485), the MA is G and the OTA is G. For SNP no. 35 (rs9494332), the MA is G and the OTA is A. For SNP no. 37 (rs911507), the MA is G and the OTA is A. Further in Table S1 SNP no. 160 (rs225630), the MA is G and the OTA is G. In Supplementary Table S2, the resulting haplotypes comprised of the above

SNPs need to be corrected accordingly. The correct alleles of the haplotypes are: for haplotype no. 83 GGA, haplotype no. 84 AAA, haplotype no. 85 AAG, haplotype no. 86 GAG, haplotype no. 87 GAA, haplotype no. 88 AAA, haplotype no. 89 AGA, haplotype no. 90 AGC, haplotype no. 91 AAT, haplotype no. 92 GAT, haplotype no. 93 GCA, haplotype no. 94 ATG, haplotype no. 95 CAA, haplotype no. 96 TGC, haplotype no. 97 AAA, haplotype no. 98 AAG, haplotype no. 99 ACG, haplotype no. 100 GCG, haplotype no. 101 AAT, haplotype no. 102 AGC, haplotype no. 103 CGC, haplotype no. 104 ATG, haplotype no. 105 GCA.

These changes do not alter the significance of the results.