

CORRECTION

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Correction: GWAS on family history of Alzheimer's disease

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In the original Article, GWAS analysis of the UK Biobank data was based on standardised SNP dosages (standardised by their standard deviation) as opposed to raw SNP dosages. The standard error-weighted meta-analysis of the UK Biobank data with the IGAP findings was

therefore incorrect. The meta-analysis has been re-run as previously specified (the key findings remain although four new genome-wide significant loci have been identified). There have also been updates to the Supplementary Tables and Fig. 1.

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