Erratum: GWAS meta-analysis reveals novel loci and genetic correlates for general cognitive function: a report from the COGENT consortium

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CORRIGENDUM

GWAS meta-analysis reveals novel loci and genetic correlates for general cognitive function: a report from the COGENT consortium


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Data access for several cohorts used in this study was provided by the National Center for Biotechnology Information (NCBI) database of Genotypes and Phenotypes (dbGaP). dbGaP accession numbers for these cohorts were as follows:


Multi-Site Collaborative Study for Genotype-Phenotype Associations in Alzheimer’s Disease (GENADA): phs000219.v1.p1.

Genetics of Late Onset Alzheimer’s Disease Study (LOAD): phs000168.v1.p1.


The acknowledgment statements for these cohorts are provided below.

Framingham Heart Study: The Framingham Heart Study is conducted and supported by the National Heart, Lung, and Blood Institute (NHLBI) in collaboration with Boston University (contracts N01HC25195 and HHSN268201500001I). This article was not prepared in collaboration with investigators of the Framingham Heart Study and does not necessarily reflect the opinions or views of the Framingham Heart Study, Boston University, or the NHLBI. Funding for SHARe Affymetrix genotyping was provided by NHLBI contract N02HL64278. SHARe Illumina genotyping was provided under an agreement between Illumina and Boston University.

Cardiovascular Health Study (CHS): this research was supported by contracts HHSN268201200036C, HHSN268200800007C, N01HC85079, N01HC85080, N01HC85081, N01HC85082, N01HC85083, N01HC85084, N01HC85085, N01HC85086, N01HC35129, N01HC15103, N01HC55222, N01HC75150, N01HC45133 and N01HC85239; grants U01 HL080295 and U01 HL130014 from the NHLBI, and R01AG023629 from the National Institute on Aging, with additional contribution from the National Institute of Neurological Disorders and Stroke. A full list of principal CHS investigators and institutions can be found at https://chs-nhlbi.org/pi. This article was not prepared in collaboration with CHS investigators and does not necessarily reflect the opinions or views of CHS or the NHLBI.

Multi-Site Collaborative Study for Genotype-Phenotype Associations in Alzheimer’s Disease: the genotypic and associated phenotypic data used in the study were provided by the GlaxoSmithKline, R&D Limited. Details on data acquisition were published previously (Li H, Wetten S, Li L, St Jean PL, Upmanyu R, Surh L et al., Candidate single-nucleotide polymorphisms from a genome-wide association study of Alzheimer disease, Arch Neurol 2009; 66: 724–728). Funding for SHARe Affymetrix genotyping was provided by NHLBI contract N02HL64278. SHARe Illumina genotyping was provided under an agreement between Illumina and Boston University.

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Long Life Family Study: funding support for the Long Life Family Study was provided by the Division of Geriatrics and Clinical Gerontology, NIA. The study includes GWAS analyses for factors that contribute to long and healthy life. Assistance with phenotype harmonization and genotype cleaning as well as with general study coordination was provided by the Division of Geriatrics and Clinical Gerontology, NIA. Support for the collection of data sets and samples was provided via Multicenter Cooperative Agreement support by the Division of Geriatrics and Clinical Gerontology, NIA (U01AG023746, U01023755, U01023749, U01023744 and U01023712). Funding support for the genotyping, which was performed at the Johns Hopkins University Center for Inherited Disease Research, was provided by the NIA.

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