

Erratum

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Integrating genetic and gene expression evidence into genome-wide association analysis of gene sets

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The Acknowledgments section in this paper was incomplete and should instead read as follows:

The results published here are in whole or part based upon data generated by The Cancer Genome Atlas pilot project established by the NCI and NHGRI. Information about TCGA and the investigators and institutions who constitute The Cancer Genome Atlas (TCGA) Research Network can be found at <http://cancergenome.nih.gov/>. The TCGA data analyzed here are available through dbGAP, accession phs000489.v3.p7 (http://www.ncbi.nlm.nih.gov/projects/gap/cgi-bin/study.cgi?study_id=phs000489.v3.p7).

The authors apologize for the incomplete Acknowledgments.



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