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Author Correction: Identifying Genetic Factors That Contribute to the Increased Risk of Congenital Heart Defects in Infants With Down Syndrome

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OPEN

Author Correction: Identifying genetic factors that contribute to the increased risk of congenital heart defects in infants with Down syndrome

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Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-020-74650-4>, published online 22 October 2020

Ivan P. Moskowitz was omitted from the author list in the original version of this Article.

The Author Contributions section now reads:

“C.E.T., A.M.H., M.E.Z., S.L.S., M.P.E., D.J.C., and J.G.M. were all involved in the conception and design of the work presented in this manuscript. C.E.T. and A.M.H. jointly drafted the manuscript, with all authors reviewing and providing feedback on the manuscript. Ascertainment of the samples and phenotype assessment of individuals with DS were accomplished through the efforts of S.L.S., C.L.M., T.C.R., K.J.D., G.T.C., R.H.R., P.J.G., J.E.O., D.C.B., L.W., C.L.C., I.P.M., and E.G. (sponsor of PGC-approved ancillary project). H.R.J., D.J.C., and B.L.R. performed mapping and variant calling for the WES and WGS datasets, and together with C.E.T. performed QC of these data. C.E.T. performed the gene- and pathway-level (SKAT) analyses. H.C. and J.O. were involved in the initial phase of these datasets and helped to set the framework for these analyses. H.J.C., B.D.K., A.J.A., and E.G. generated the GWAS summary results used as training data for the PRS analyses. A.M.H. performed imputation of the array genotype data, and QC and analyses for the PRS application.”

The Acknowledgements section now reads:

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The original Article and accompanying Supplementary Information file have been corrected.



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