

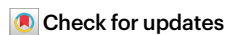


Author Correction: Rare variant analyses in 51,256 type 2 diabetes cases and 370,487 controls reveal the pathogenicity spectrum of monogenic diabetes genes

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Alicia Huerta-Chagoya, Philip Schroeder, Ravi Mandla, Jiang Li, Lowri Morris, Maheak Vora, Ahmed Alkanaq, Dorka Nagy, Lukasz Szczerbinski, Jesper G. S. Madsen , Silvia Bonàs-Guarch , Fanny Mollandin , Joanne B. Cole , Bianca Porneala, Kenneth Westerman, Josephine H. Li , Toni I. Pollin, Jose C. Florez , Anna L. Gloyn , David J. Carey, Inês Cebola , Uyenlinh L. Mirshahi , Alisa K. Manning , Aaron Leong, Miriam Udler & Josep M. Mercader

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