

## Author Correction

GenOMICC Investigators; SCOURGE Consortium; ISARICC Investigators; The 23andMe COVID-19 Team; Pairo-Castineira, Erola; Rawlik, Konrad; Bretherick, Andrew D.; Qi, Ting; Wu, Yang; Nassiri, Isar; McConkey, Glenn A.; Zechner, Marie; Klaric, Lucija; Griffiths, Fiona; Oosthuyzen, Wilna; Kousathanas, Athanasios; Richmond, Anne; Millar, Jonathan; Russell, Clark D.; Malinauskas, Tomas

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# Author Correction: GWAS and meta-analysis identifies 49 genetic variants underlying critical COVID-19

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Erola Pairo-Castineira, Konrad Rawlik, Andrew D. Bretherick, Ting Qi, Yang Wu, Isar Nassiri, Glenn A. McConkey, Marie Zechner, Lucija Klaric, Fiona Griffiths, Wilna Oosthuyzen, Athanasios Kousathanas, Anne Richmond, Jonathan Millar, Clark D. Russell, Tomas Malinauskas, Ryan Thwaites, Kirstie Morrice, Sean Keating, David Maslove, Alistair Nichol, Malcolm G. Semple, Julian Knight, Manu Shankar-Hari, Charlotte Summers, Charles Hinds, Peter Horby, Lowell Ling, Danny McAuley, Hugh Montgomery, Peter J. M. Openshaw, Colin Begg, Timothy Walsh, Albert Tenesa, Carlos Flores, José A. Riancho, Augusto Rojas-Martinez, Pablo Lapunzina, GenOMICC Investigators, SCOURGE Consortium, ISARICC Investigators, The 23andMe COVID-19 Team, Jian Yang, Chris P. Ponting, James F. Wilson, Veronique Vitart, Malak Abedalthagafi, Andre D. Luchessi, Esteban J. Parra, Raquel Cruz, Angel Carracedo, Angie Fawkes, Lee Murphy, Kathy Rowan, Alexandre C. Pereira, Andy Law, Benjamin Fairfax, Sara Clohisey Hendry & J. Kenneth Baillie

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In the version of this article initially published, the name of Ana Margarita Baldi3n-Elorza, of the SCOURGE Consortium, appeared incorrectly (as Ana Mar3a Baldion) and has now been amended in the HTML and PDF versions of the article.



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