



J Kenneth Baillie

Kontakt

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Publikationen (1)

Kousathanas A, Pairo-Castineira E, Rawlik K, Stuckey A, Odhams C, Walker S, Russell C, Malinauskas T, Wu Y, Millar J, Shen X, Elliott K, Griffiths F, Oosthuyzen W, Morrice K, Keating S, Wang B, Rhodes D, Klaric L, Zechner M, Parkinson N, Siddiq A, Goddard P, Donovan S, Maslove D, Nichol A, Semple M, Zainy T, Maleady-Crowe F, Todd L, Salehi S, Knight J, Elgar G, Chan G, Arumugam P, Patch C, Rendon A, Bentley D, Kingsley C, Kosmicki J, Horowitz J, Baras A, Abecasis G, Ferreira M, Justice A, Mirshahi T, Oetjens M, Rader D, Ritchie M, Verma A, Fowler T, Shankar-Hari M, Summers C, Hinds C, Horby P, Ling L, McAuley D, Montgomery H, Openshaw P, Elliott P, Walsh T, Tenesa A, Fawkes A, Murphy L, Rowan K, Ponting C, Vitart V, Wilson J, Yang J, Bretherick A, Scott R, Hendry S, Moutsianas L, Law A, Caulfield M, Baillie J, Albrich W. Whole-genome sequencing reveals host factors underlying critical COVID-19. *Nature* 2022; 607:97-103.

Projekte (0)

Keine Resultate gefunden.

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