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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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