



David Maslove

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.

Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

support.forschung@kssg.ch