



Richard K Russell

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Publications (2)

Anderson C, Barrett J, Wilson R, Ahmed A, Karaminejadranjbar M, Fulga T, Baeumler T, Danesh J, Roberts D, Ouwehand W, Sambrook J, Elkadri A, Griffiths A, COLORS in IBD group investigators, Uhlig H, Wilson D, Muise A, Shah N, Snapper S, NIDDK IBD Genetics Consortium, UK IBD Genetics Consortium, Swiss IBD Cohort Investigators, INTERVAL Study, Oxford IBD cohort study investigators, Moore C, Sullivan P, Wedrychowicz A, Russell R, Rodrigues A, Posovszky C, Croft N, Kammermeier J, Pandey S, Fachal L, Cavounidis A, Moutsianas L, Schwerd T, Barakat F, Auth M, Heuschkel R, Gilmour K, Matte J, Ferry H, Thapar N, Parkes M, Satsangi J, Travis S, Braegger C, Fyderek K, Zilbauer M, Serra E. Somatic mosaicism and common genetic variation contribute to the risk of very-early-onset inflammatory bowel disease. *Nat Commun* 2020; 11:995.

Kupcinkas L, Potocnik U, Prescott N, Regueiro M, Rotter J, Russell R, Sanderson J, Sans M, Satsangi J, Schreiber S, Simms L, Sventoraityte J, Ponsioen C, Palmieri O, Kugathasan S, Latiano A, Laukens D, Lawrance I, Lees C, Louis E, Mahy G, Mansfield J, Morgan A, Mowat C, Newman W, Targan S, Taylor K, Tremelling M, Hakonarson H, Brant S, Radford-Smith G, Mathew C, Rioux J, Schadt E, Daly M, Franke A, Parkes M, Vermeire S, Barrett J, Annese V, Silverberg M, Verspaget H, De Vos M, Wijmenga C, Wilson D, Winkelmann J, Xavier R, Zeissig S, Zhang B, Zhang C, Zhao H, Cho J, Karlsen T, Jostins L, Theatre E, Spain S, Raychaudhuri S, Goyette P, Wei Z, Abraham C, Achkar J, Ahmad T, Amininejad L, Ananthakrishnan A, Andersen V, Cleyne I, Ning K, Ripke S, Weersma R, Duerr R, McGovern D, Hui K, Lee J, Schumm L, Sharma Y, Anderson C, Essers J, Mitrovic M, Andrews J, Baidoo L, Balschun T, Ferguson L, Franchimont D, Fransen K, Geary R, Georges M, Gieger C, Glas J, Haritunians T, Hart A, Hawkey C, Hedl M, Ellinghaus D, Edwards C, Bampton P, Bitton A, Boucher G, Brand S, Büning C, Cohain A, Cichon S, D'Amato M, De Jong D, Devaney K, Dubinsky M, Hu X. Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. *Nature* 2012; 491:119–24.

Projects (0)

No results found.

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