



**Jeremy D Sanderson**

**Kontakt**

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## Publikationen (2)

Franke A, Büning C, Duerr R, Nöthen M, Wang J, Vatn M, Mathew C, Sanderson J, Rutgeerts P, Vermeire S, McArdle W, Strachan D, Wijmenga C, Sans M, Brand S, Glas J, Parkes M, Schreiber S, Rosenstiel P, Subramani S, Karlsen T, Nothnagel M, Daly M, D'Amato M, Halfvarson J, Annese V, Latiano A, Illig T, Winkelmann J, Ponsioen C, Weersma R, Nikolaus S, Liu X, Doncheva N, Skieceviciene J, Rivas M, Keller A, Ellinghaus E, Bromberg Y, Stade B, Jiang T, Till A, Lipinski S, Zeissig S, Zhang H, Liu Q, Jiang F, Krawczak M, Kayser M, Kupcinkas L, Vogel U, Andersen V, Lee J, Berzuini C, Goodall J, Boehm B, Häsler R, Albrecht M, Mayr G, Forster M, Ellinghaus D. Association between variants of PRDM1 and NDP52 and Crohn's disease, based on exome sequencing and functional studies. *Gastroenterology* 2013; 145:339-47.

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, Gearry 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.

## Projekte (0)

Keine Resultate gefunden.

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Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

[support.forschung@kssg.ch](mailto:support.forschung@kssg.ch)