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Publications (4)

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, Kupcinskas 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.

Thomsen I, Vatn M, Silverberg M, Duerr R, Padyukov L, Brand S, Sans M, Annese V, Achkar J, Melum E, Durie P, Sandford R, Mayr G, König I, Hveem K, Cleynen I, Gutierrez-Achury J, Ricaño-Ponce I, van Heel D, Björnsson E, Boberg K, Marschall H, Schreiber S, Manns M, Färkkilä M, Dale A, Chapman R, Lazaridis K, Franke A, Anderson C, Cho J, Bergquist A, Alexander G, Chazouillères O, Bowlus C, Wijmenga C, Schrupf E, Vermeire S, Albrecht M, Rioux J, Karlsen T, Næss S, Liu J, Gotthardt D, Pares A, Ellinghaus D, Shah T, Juran B, Milkiewicz P, Rust C, Schramm C, Hirschfield G, Invernizzi P, Eksteen B, Hov J, Folseraas T, Ellinghaus E, Rushbrook S, Doncheva N, Andreassen O, Weersma R, Weismüller T, Müller T, Srivastava B, Saarela J, Leppä V, Dorfman R, Alvaro D, Floreani A, Onengut-Gumuscu S, Rich S, Thompson W, Mason A, Teufel A, Sterneck M, Dalekos G, Nöthen M, Herms S, Winkelmann J, Mitrovic M, Braun F, Ponsioen C, Croucher P, Schork A. Dense genotyping of immune-related disease regions identifies nine new risk loci for primary sclerosing cholangitis. *Nat Genet* 2013; 45:670-5.

Kupcinskas 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, Cleynen 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.

Phillips A, van den Berg L, Vatn M, Verspaget H, Walters T, Wijmenga C, Wilson D, Westra H, Xavier R, Zhao Z, Ponsioen C, Andersen V, Torkvist L, Targan S, Steinhart A, Prescott N, Proctor D, Roberts R, Russell R, Rutgeerts P, Sanderson J, Sans M, Schumm P, Seibold F, Sharma Y, Simms L, Seielstad M, Gazouli M, Anagnou N, Satsangi J, Cho J, Schreiber S, Daly M, Barrett J, Parkes M, Annese V, Hakonarson H, Radford-Smith G, Duerr R, Vermeire S, Weersma R, Chamailard M, Brant S, Karlsen T, Kupcinskas L, Sventoraityte J, Mansfield J, Kugathasan S, Silverberg M, Halfvarson J, Rotter J, Mathew C, Griffiths A, Geary R, Ahmad T, Rioux J, Panés J, Anderson C, Baldassano R, Barclay M, Bayless T, Brand S, Büning C, Colombel J, Denson L, De Vos M, Dubinsky M, Edwards C, Ellinghaus D, Fehrmann R, Baidoo L, Bumpstead S, Boucher G, Lees C, Franke A, D'Amato M, Taylor K, Lee J, Goyette P, Imielinski M, Latiano A, Lagacé C, Scott R, Amininejad L, Floyd J, Florin T, Libioulle C, Louis E, McGovern D, Milla M, Montgomery G, Morley K, Mowat C, Ng A, Newman W, Ophoff R, Papi L, Palmieri O, Levine A, Lemann M, Franchimont D, Franke L, Georges M, Glas J, Glazer N, Guthery S, Haritunians T, Hayward N, Hugot J, Jobin G, Laukens D, Lawrance I, Peyrin-Biroulet L. Meta-analysis identifies 29 additional ulcerative colitis risk loci, increasing the number of confirmed associations to 47. *Nat Genet* 2011; 43:246-52.

Projects (0)

No results found.

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