



International IBD Genetics Consortium

Kontakt

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Publikationen (2)

Akhlaghpour M, Haritunians T, More S, Thomas L, Stamps D, Dube S, Li D, Yang S, Landers C, Mengesha E, Hamade H, Murali R, Potdar A, Wolf A, Botwin G, Khrom M, International IBD Genetics Consortium, Ananthakrishnan A, Faubion W, Jabri B, Lira S, Newberry R, Sandler R, Sartor R, Xavier R, Brant S, Cho J, Duerr R, Lazarev M, Rioux J, Schumm L, Silverberg M, Zoghiyan K, Fleshner P, Melmed G, Vasiliauskas E, Ha C, Rabizadeh S, Syal G, Bonthala N, Ziring D, Targan S, Long M, McGovern D, Michelsen K, Brand S. Genetic coding variant in complement factor B (CFB) is associated with increased risk for perianal Crohn's disease and leads to impaired CFB cleavage and phagocytosis. *Gut* 2023; 72:2068-2080.

Momozawa Y, Hoentjen F, Löwenberg M, Oldenburg B, Pierik M, Vander Meulen-de Jong A, Janneke van der Woude C, Visschedijk M, International IBD Genetics Consortium, Lathrop M, Hugot J, Weersma R, De Vos M, Franchimont D, Vermeire S, Kubo M, Louis E, Bouma G, Amininejad L, Dmitrieva J, Theatre E, Deffontaine V, Rahmouni S, Charlotheaux B, Crins F, Docampo E, Elansary M, Gori A, Lecut C, Mariman R, Mni M, Oury C, Altukhov I, Alexeev D, Aulchenko Y, Georges M. IBD risk loci are enriched in multigenic regulatory modules encompassing putative causative genes. *Nat Commun* 2018; 9:2427.

Projekte (0)

Keine Resultate gefunden.

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