



**L Philip Schumm**

**Kontakt**

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## Publikationen (4)

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, Ananthkrishnan 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, Zaghiyan 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.

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Lees C, van der Meulen-de Jong A, Theatre E, Schreiber S, Satsangi J, Lee J, Lawrance I, Kupcinkas L, Kennedy N, Weersma R, Wilson D, Barrett J, McGovern D, Radford-Smith G, Silverberg M, Mansfield J, Rioux J, Vermeire S, Parkes M, Huang H, Hov J, Halfvarson J, Annese V, Andrews J, Andersen V, Ahmad T, Zeissig S, Schumm L, Jostins L, Boucher G, Brand S, Brant S, Cho J, Hakonarson H, Goyette P, Geary R, Franke A, Ferguson L, Duerr R, Dubinsky M, Daly M, Cleyne I. Inherited determinants of Crohn's disease and ulcerative colitis phenotypes: a genetic association study. *Lancet* 2015; 387:156-67.

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, Ananthkrishnan 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.

## Projekte (0)

Keine Resultate gefunden.

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