



Bertram Müller-Myhsok

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Publications (14)

Glas J, Duerr R, Franke A, Kamboh M, Achkar J, Balschun T, Müller-Myhsok B, Wolf C, Olszak T, Wetzke M, Diegelmann J, Pasciuto G, Czamara D, Seiderer J, Brand S. PTGER4 expression-modulating polymorphisms in the 5p13.1 region predispose to Crohn's disease and affect NF- κ B and XBP1 binding sites. *PLoS one* 2012; 7:e52873.

Glas J, Czamara D, Diegelmann J, Müller-Myhsok B, Lohse P, Wolf C, Ochsenkühn T, Göke B, Lass U, Olszak T, Beigel F, Weidinger M, Pfennig S, Tillack C, Fries C, Seiderer J, Brand S. CEACAM6 gene variants in inflammatory bowel disease. *PLoS one* 2011; 6:e19319.

Glas J, Czamara D, Diegelmann J, Lohse P, Ochsenkühn T, Göke B, Müller-Myhsok B, Weidinger M, Laubender R, Olszak T, Jürgens M, Beigel F, Pfennig S, Tillack C, Seiderer J, Brand S. The NOD2 single nucleotide polymorphisms rs2066843 and rs2076756 are novel and common Crohn's disease susceptibility gene variants. *PLoS one* 2010; 5:e14466.

Glas J, Czamara D, Diegelmann J, Müller-Myhsok B, Folwaczny M, Ochsenkühn T, Göke B, Weidinger M, Olszak T, Beigel F, Wetzke M, Pfennig S, Tengler B, Fischer D, Seiderer J, Brand S. Pregnane X receptor (PXR/NR112) gene haplotypes modulate susceptibility to inflammatory bowel disease. *Inflamm Bowel Dis* 2010; 17:1917-24.

Glas J, Roeske D, Diegelmann J, Müller-Myhsok B, Ochsenkühn T, Göke B, Paschos E, Wetzke M, Pfennig S, Markus C, Seiderer J, Brand S. Role of PPARG gene variants in inflammatory bowel disease. *Inflamm Bowel Dis* 2010; 17:1057-8.

Glas J, Roeske D, Müller-Myhsok B, Diegelmann J, Ochsenkühn T, Göke B, Folwaczny M, Lohse P, Epplen J, Klein W, Pfennig S, Weidinger M, Beigel F, Fries C, Nagy M, Seiderer J, Brand S. Evidence for STAT4 as a common autoimmune gene: rs7574865 is associated with colonic Crohn's disease and early disease onset. *PLoS one* 2010; 5:e10373.

Török H, Müller-Myhsok B, Folwaczny C, Göke B, Folwaczny M, Ochsenkühn T, Lohse P, Klein W, Wetzke M, Teshome M, Tonenchi L, Endres I, Glas J, Brand S. Epistasis between Toll-like receptor-9 polymorphisms and variants in NOD2 and IL23R modulates susceptibility to Crohn's disease. *Am J Gastroenterol* 2009; 104:1723-33.

Glas J, Müller-Myhsok B, Ochsenkühn T, Göke B, Lohse P, Folwaczny M, Koletzko S, Lacher M, Schiemann U, Griga T, Epplen J, Klein W, Pfennig S, Wetzke M, Ripke S, Stallhofer J, Brand S. Novel genetic risk markers for ulcerative colitis in the IL2/IL21 region are in epistasis with IL23R and suggest a common genetic background for ulcerative colitis and celiac disease. *Am J Gastroenterol* 2009; 104:1737-44.

Glas J, Klein W, Epplen J, Schiemann U, Mussack T, Lohse P, Göke B, Ochsenkühn T, Folwaczny M, Müller-Myhsok B, Griga T, Jürgens M, Stallhofer J, Seiderer J, Pasciuto G, Tillack C, Diegelmann J, Pfennig S, Konrad A, Schmechel S, Wetzke M, Török H, Brand S. rs224136 on chromosome 10q21.1 and variants in PHOX2B, NCF4, and FAM92B are not major genetic risk factors for susceptibility to Crohn's disease in the German population. *Am J Gastroenterol* 2009; 104:665-72.

Seiderer J, Lohse P, Müller-Myhsok B, Ochsenkühn T, Göke B, Konrad A, Schmechel S, Jürgens M, Pfennig S, Tillack C, Stallhofer J, Glas J, Diegelmann J, Elben I, Brand S. Role of the novel Th17 cytokine IL-17F in inflammatory bowel disease (IBD): upregulated colonic IL-17F expression in active Crohn's disease and analysis of the IL17F p.His161Arg polymorphism in IBD. *Inflamm Bowel Dis* 2008; 14:437-45.

Seiderer J, Reinecker H, Lohse P, Ochsenkühn T, Göke B, Müller-Myhsok B, Jürgens M, Pfennig S, Niess J, Glas J, Tillack C, Leistner D, Dambacher J, Brand S. Genotype-phenotype analysis of the CXCL16 p.Ala181Val polymorphism in inflammatory bowel disease. *Clin Immunol* 2008; 127:49-55.

Glas J, Klein W, Epplen J, Folwaczny C, Lohse P, Göke B, Ochsenkühn T, Mussack T, Folwaczny M, Müller-Myhsok B, Griga T, Haller D, Pfennig S, Konrad A, Schmechel S, Dambacher J, Seiderer J, Schroff F, Wetzke M, Roeske D, Török H, Tonenchi L, Brand S. The ATG16L1 gene variants rs2241879 and rs2241880 (T300A) are strongly associated with susceptibility to Crohn's disease in the German population. *Am J Gastroenterol* 2007; 103:682-91.

Glas J, Epplen J, Schiemann U, Folwaczny C, Lohse P, Göke B, Ochsenkühn T, Müller-Myhsok B, Folwaczny M, Mussack T, Klein W, Griga T, Maier K, Seiderer J, Wetzke M, Konrad A, Török H, Schmechel S, Tonenchi L, Grassl C, Dambacher J, Pfennig S, Brand S. rs1004819 is the main disease-associated IL23R variant in German Crohn's disease patients: combined analysis of IL23R, CARD15, and OCTN1/2 variants. *PloS one* 2007; 2:e819.

Glas J, Folwaczny M, Ochsenkühn T, Brand S, Schnitzler F, Seiderer J, Lohse P, Schiemann U, Griga T, Epplen J, Klein W, Wetzke M, Mussack T, Müller-Myhsok B, Tonenchi L, Török H, Folwaczny C. Role of the NFKB1 -94ins/delATTG promoter polymorphism in IBD and potential interactions with polymorphisms in the CARD15/NOD2, IKBL, and IL-1RN genes. *Inflamm Bowel Dis* 2006; 12:606-11.

Projects (0)

No results found.

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