



Julia Seiderer

Kontakt

Julia Seiderer

Publikationen (37)

Beigel F, Brand S, Seiderer J, Göke B, Laubender R, Van Steen K, John J, Breitenreicher S, Tillack C, Schnitzler F, Steinborn A, Ochsenkühn T. Risk of malignancies in patients with inflammatory bowel disease treated with thiopurines or anti-TNF alpha antibodies. *Pharmacoepidemiol Drug Saf* 2014; 23:735-44.

Glas J, Czamara D, Diegelmann J, Göke B, Friedrich M, Steib C, Beigel F, Wetzke M, Tsekeri E, Olszak T, Fries C, Stallhofer J, Bues S, Seiderer J, Brand S. IRGM variants and susceptibility to inflammatory bowel disease in the German population. *PLoS one* 2013; 8:e54338.

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, Ochsenkühn T, Göke B, Wetzke M, Steib C, Stallhofer J, Beigel F, Friedrich M, Tillack C, Fries C, Olszak T, Wagner J, Seiderer J, Brand S. Analysis of IL12B gene variants in inflammatory bowel disease. *PLoS one* 2012; 7:e34349.

Glas J, Czamara D, Diegelmann J, Karbalai N, Ochsenkühn T, Göke B, Steib C, Friedrich M, Stallhofer J, Tillack C, Beigel F, Wetzke M, Olszak T, Seiderer J, Wagner J, Brand S. PTPN2 gene variants are associated with susceptibility to both Crohn's disease and ulcerative colitis supporting a common genetic disease background. *PLoS one* 2012; 7:e33682.

Glas J, Czamara D, Diegelmann J, Friedrich M, Steib C, Beigel F, Olszak T, Tillack C, Fries C, Wetzke M, Bayrle C, Seiderer J, Brand S. The role of osteopontin (OPN/SPP1) haplotypes in the susceptibility to Crohn's disease. *PLoS one* 2011; 6:e29309.

Beigel F, Ochsenkühn T, Seiderer J, Göke B, Weidinger M, Breitenreicher S, Schnitzler F, Tillack C, Laubender R, Löhr B, Brand S. Iron status and analysis of efficacy and safety of ferric carboxymaltose treatment in patients with inflammatory bowel disease. *Digestion* 2011; 85:47-54.

Schnitzler F, Seiderer J, Stallhofer J, Brand S. Dominant disease-causing effect of NOD2 mutations in a family with all family members affected by Crohn's disease. *Inflamm Bowel Dis* 2011; 18:395-6.

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.

Beigel F, Schnitzler F, Paul Laubender R, Pfennig S, Weidinger M, Göke B, Seiderer J, Ochsenkühn T, Brand S. Formation of antinuclear and double-strand DNA antibodies and frequency of lupus-like syndrome in anti-TNF- α antibody-treated patients with inflammatory bowel disease. *Inflamm Bowel Dis* 2011; 17:91-8.

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, Seiderer J, Tillack C, Paschos E, Wetzke M, Diegelmann J, Czamara D, Brand S. Functional SFTPD gene variants are not associated with susceptibility to inflammatory bowel disease in the German population. *Inflamm Bowel Dis* 2010; 17:1439–40.

Diegelmann J, Seiderer J, Niess J, Haller D, Göke B, Reinecker H, Brand S. Expression and regulation of the chemokine CXCL16 in Crohn's disease and models of intestinal inflammation. *Inflamm Bowel Dis* 2010; 16:1871–81.

Jürgens M, Ochsenkühn T, Göke B, Tillack C, Beigel F, Hasbargen U, Hübener C, Filik L, Brand S, Seiderer J. Safety of adalimumab in Crohn's disease during pregnancy: case report and review of the literature. *Inflamm Bowel Dis* 2010; 16:1634–6.

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 PPAR γ 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.

Jürgens M, Herrmann K, Lohse P, Göke B, Kreis M, Schnitzler F, Weidinger M, Beigel F, Tillack C, Pfennig S, Wagner J, Wetzke M, Glas J, Seiderer J, Laubender R, Brand S, Ochsenkühn T. The presence of fistulas and NOD2 homozygosity strongly predict intestinal stenosis in Crohn's disease independent of the IL23R genotype. *J Gastroenterol* 2010; 45:721–31.

Jürgens M, Ochsenkühn T, Glas J, Göke B, Lohse P, Tillack C, Schnitzler F, Stallhofer J, Pfennig S, Beigel F, Wetzke M, Wagner J, Seiderer J, Weidinger M, Hartl F, Laubender R, Brand S. Disease activity, ANCA, and IL23R genotype status determine early response to infliximab in patients with ulcerative colitis. *Am J Gastroenterol* 2010; 105:1811–9.

Beigel F, Jürgens M, Filik L, Bader L, Lück C, Göke B, Ochsenkühn T, Brand S, Seiderer J. Severe Legionella pneumophila pneumonia following infliximab therapy in a patient with Crohn's disease. *Inflamm Bowel Dis* 2009; 15:1240–4.

Seiderer J, Brand S. IL-22: a two-headed cytokine in IBD?. *Inflamm Bowel Dis* 2009; 15:473–4.

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.

Tillack C, Seiderer J, Brand S, Göke B, Reiser M, Schaefer C, Diepolder H, Ochsenkühn T, Herrmann K. Correlation of magnetic resonance enteroclysis (MRE) and wireless capsule endoscopy (CE) in the diagnosis of small bowel lesions in Crohn's disease. *Inflamm Bowel Dis* 2008; 14:1219–28.

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.

Dambacher J, Beigel F, Seiderer J, Haller D, Göke B, Auernhammer C, Brand S. Interleukin 31 mediates MAP kinase and STAT1/3 activation in intestinal epithelial cells and its expression is upregulated in inflammatory bowel disease. *Gut* 2007; 56:1257-65.

Dambacher J, Lohse P, Ochsenkühn T, Göke B, Diebold J, Otte J, Tillack C, Konrad A, Hofbauer K, Pfennig S, Schnitzler F, Susic Z, Seiderer J, Staudinger T, Brand S. Macrophage migration inhibitory factor (MIF) -173G/C promoter polymorphism influences upper gastrointestinal tract involvement and disease activity in patients with Crohn's disease. *Inflamm Bowel Dis* 2007; 13:71-82.

Seiderer J, Lohse P, Göke B, Sackmann M, Tillack C, Dambacher J, Hofbauer K, Herrmann K, Pfennig S, Staudinger T, Brand S, Schnitzler F, Ochsenkühn T. Homozygosity for the CARD15 frameshift mutation 1007fs is predictive of early onset of Crohn's disease with ileal stenosis, entero-enteral fistulas, and frequent need for surgical intervention with high risk of re-stenosis. *Scand J Gastroenterol* 2006; 41:1421-32.

Seiderer J, Lohse P, Göke B, Schoenberg S, Pfennig S, Crispin A, Hatz R, Schnitzler F, Herrmann K, Brand S, Ochsenkühn T. Predictive value of the CARD15 variant 1007fs for the diagnosis of intestinal stenoses and the need for surgery in Crohn's disease in clinical practice: results of a prospective study. *Inflamm Bowel Dis* 2006; 12:1114-21.

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.

Seiderer J, Zech C, Diebold J, Schoenberg S, Brand S, Tillack C, Göke B, Ochsenkühn T. Nodular regenerative hyperplasia: a reversible entity associated with azathioprine therapy. *Eur J Gastroenterol Hepatol* 2006; 18:553-5.

Brand S, Auernhammer C, Göke B, Ochsenkühn T, Seiderer J, Herrmann K, Leclair S, Popp A, Jagla W, Marquardt A, Diepolder H, Otte J, Eichhorst S, Zitzmann K, Olszak T, Beigel F, Dambacher J. IL-22 is increased in active Crohn's disease and promotes proinflammatory gene expression and intestinal epithelial cell migration. *Am J Physiol Gastrointest Liver Physiol* 2006; 290:G827-38.

Schnitzler F, Ochsenkühn T, Göke B, Tillack C, Seiderer J, Hofbauer K, Pfennig S, Staudinger T, Brand S, Lohse P. Eight novel CARD15 variants detected by DNA sequence analysis of the CARD15 gene in 111 patients with inflammatory bowel disease. *Immunogenetics* 2006; 58:99-106.

Brand S, Ochsenkühn T, Göke B, Konrad A, Tillack C, Seiderer J, Pfennig S, Staudinger T, Schnitzler F, Dambacher J, Hofbauer K, Lohse P. Increased expression of the chemokine fractalkine in Crohn's disease and association of the fractalkine receptor T280M polymorphism with a fibrostenosing disease phenotype. *Am J Gastroenterol* 2006; 101:99-106.

Brand S, Lohse P, Göke B, Crispin A, Konrad A, Tillack C, Seiderer J, Dambacher J, Hofbauer K, Pfennig S, Schnitzler F, Staudinger T, Ochsenkühn T. The role of Toll-like receptor 4 Asp299Gly and Thr399Ile polymorphisms and CARD15/NOD2 mutations in the susceptibility and phenotype of Crohn's disease. *Inflamm Bowel Dis* 2005; 11:645-52.

Projekte (0)

Keine Resultate gefunden.

Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

support.forschung@kssg.ch