Kantonsspital St.Gallen

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Publikationen (5)

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.

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.

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.

Projekte (0)

Keine Resultate gefunden.

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