Kantonsspital St.Gallen

Martin Lacher

Kontakt

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Publikationen (4)

Mingardo E, Beaman G, Grote P, Nordenskjöld A, Newman W, Woolf A, Eckstein M, Hilger A, Dworschak G, Rösch W, Ebert A, Stein R, Brusco A, Di Grazia M, Tamer A, Torres F, Hernandez J, Erben P, Maj C, Olmos J, Riancho J, Valero C, Hostettler I, Houlden H, Werring D, Schumacher J, Gehlen J, Giel A, Buerfent B, Arkani S, Åkesson E, Rotstein E, Ludwig M, Holmdahl G, Giorgio E, Berettini A, Keene D, Cervellione R, Younsi N, Ortlieb M, Oswald J, Haid B, Promm M, Neissner C, Hirsch K, Stehr M, Schäfer F, Schmiedeke E, Boemers T, van Rooij I, Feitz W, Marcelis C, Lacher M, Nelson J, Ure B, Fortmann C, Gale D, Chan M, Ludwig K, Nöthen M, Heilmann S, Zwink N, Jenetzky E, Odermatt B, Knapp M, Reutter H. A genome-wide association study with tissue transcriptomics identifies genetic drivers for classic bladder exstrophy. Commun Biol 2022; 5:1203.

Török H, Bellon V, Konrad A, Lacher M, Tonenchi L, Siebeck M, Brand S, De Toni E. Functional Toll-Like Receptor (TLR)2 polymorphisms in the susceptibility to inflammatory bowel disease. PloS one 2017; 12:e0175180.

Schroepf S, Koletzko S, von Schweinitz D, Berger M, Ballauff A, Helmbrecht J, Hoster E, Glas J, Lohse P, Prell C, Brand S, Kappler R, Lacher M. Strong overexpression of CXCR3 axis components in childhood inflammatory bowel disease. Inflamm Bowel Dis 2010; 16:1882-90.

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.

Projekte (0)

Keine Resultate gefunden.

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