



**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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Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

[support.forschung@kssg.ch](mailto:support.forschung@kssg.ch)