



**Christiane Wolf**

**Kontakt**

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## Publikationen (6)

Schnitzler F, Friedrich M, Angelberger M, Diegelmann J, Stallhofer J, Wolf C, Dütschler J, Truniger S, Olszak T, Beigel F, Tillack C, Lohse P, Brand S. Development of a uniform, very aggressive disease phenotype in all homozygous carriers of the NOD2 mutation p.Leu1007fsX1008 with Crohn's disease and active smoking status resulting in ileal stenosis requiring surgery. PloS one 2020; 15:e0236421.

Schnitzler F, Grüner N, Rust C, Guba M, Denk G, Zachoval R, Göke B, Tillack C, Beigel F, Olszak T, Angelberger M, Wolf C, Karbalai N, Habicht A, Fischereeder M, Schönermarck U, Stallhofer J, Friedrich M, Brand S. Solid Organ Transplantation in Patients with Inflammatory Bowel Diseases (IBD): Analysis of Transplantation Outcome and IBD Activity in a Large Single Center Cohort. PloS one 2015; 10:e0135807.

Schnitzler F, Lohse P, Glas J, Göke B, Beigel F, Tillack C, Olszak T, Diegelmann J, Angelberger M, Stallhofer J, Wolf C, Friedrich M, Brand S. The NOD2 Single Nucleotide Polymorphism rs72796353 (IVS4+10 A>C) Is a Predictor for Perianal Fistulas in Patients with Crohn's Disease in the Absence of Other NOD2 Mutations. PloS one 2015; 10:e0116044.

Schnitzler F, Lohse P, Glas J, Göke B, Stallhofer J, Tillack C, Beigel F, Olszak T, Diegelmann J, Angelberger M, Wolf C, Friedrich M, Brand S. The NOD2 p.Leu1007fsX1008 mutation (rs2066847) is a stronger predictor of the clinical course of Crohn's disease than the FOXO3A intron variant rs12212067. PloS one 2014; 9:e108503.

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- $\kappa$ 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.

## Projekte (0)

Keine Resultate gefunden.

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