Rinse K Weersma

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/1402993/publications.pdf

Version: 2024-02-01

24 papers	1,789 citations	623734 14 h-index	642732 23 g-index
24 all docs	24 docs citations	24 times ranked	2418 citing authors

#	Article	IF	Citations
1	Role of the gut microbiome in mediating lactose intolerance symptoms. Gut, 2022, 71, 215-217.	12.1	18
2	Cross-cohort gut microbiome associations with immune checkpoint inhibitor response in advanced melanoma. Nature Medicine, 2022, 28, 535-544.	30.7	158
3	Environmental factors shaping the gut microbiome in a Dutch population. Nature, 2022, 604, 732-739.	27.8	239
4	Whole exome sequencing analyses reveal geneâ€“microbiota interactions in the context of IBD. Gut, 2021, 70, gutjnl-2019-319706.	12.1	26
5	Genetic Risk Scores Identify Genetic Aetiology of Inflammatory Bowel Disease Phenotypes. Journal of Crohn's and Colitis, 2021, 15, 930-937.	1.3	8
6	Environmental factors associated with biological use and surgery in inflammatory bowel disease. Journal of Gastroenterology and Hepatology (Australia), 2021, 36, 1022-1034.	2.8	2
7	Donor genetic variants as risk factors for thrombosis after liver transplantation: A genome-wide association study. American Journal of Transplantation, 2021, 21, 3133-3147.	4.7	4
8	Long-term dietary patterns are associated with pro-inflammatory and anti-inflammatory features of the gut microbiome. Gut, 2021, 70, 1287-1298.	12.1	246
9	Large-scale genetic analyses in an understudied disease: haemorrhoidal disease. Gut, 2021, 70, 1429-1430.	12.1	2
10	A combination of fecal calprotectin and human beta-defensin 2 facilitates diagnosis and monitoring of inflammatory bowel disease. Gut Microbes, 2021, 13, 1943288.	9.8	4
11	Inulin-grown Faecalibacterium prausnitzii cross-feeds fructose to the human intestinal epithelium. Gut Microbes, 2021, 13, 1993582.	9.8	12
12	Predicted efficacy of a pharmacogenetic passport for inflammatory bowel disease. Alimentary Pharmacology and Therapeutics, 2020, 51, 1105-1115.	3.7	17
13	Interaction between drugs and the gut microbiome. Gut, 2020, 69, 1510-1519.	12.1	451
14	Latent cytomegalovirus infection does not influence long-term disease outcomes in inflammatory bowel disease, but is associated with later onset of disease. Scandinavian Journal of Gastroenterology, 2020, 55, 891-896.	1.5	2
15	Development and validation of a web-based questionnaire to identify environmental risk factors for inflammatory bowel disease: the Groningen IBD Environmental Questionnaire (GIEQ). Journal of Gastroenterology, 2019, 54, 238-248.	5.1	16
16	SLC39A8 missense variant is associated with Crohn's disease but does not have a major impact on gut microbiome composition in healthy subjects. PLoS ONE, 2019, 14, e0211328.	2.5	10
17	Anti-inflammatory Gut Microbial Pathways Are Decreased During Crohnâ€™s Disease Exacerbations. Journal of Crohn's and Colitis, 2019, 13, 1439-1449.	1.3	39
18	Analysis of 1135 gut metagenomes identifies sex-specific resistome profiles. Gut Microbes, 2019, 10, 358-366.	9.8	118

#	Article	IF	Citations
19	The influence of proton pump inhibitors and other commonly used medication on the gut microbiota. Gut Microbes, 2017, 8, 351-358.	9.8	136
20	A large variety of clinical features and concomitant disorders in celiac disease â€“ A cohort study in the Netherlands. Digestive and Liver Disease, 2016, 48, 499-505.	0.9	51
21	Down the line from genome-wide association studies in inflammatory bowel disease: the resulting clinical benefits and the outlook for the future. Expert Review of Clinical Immunology, 2015, 11, 33-44.	3.0	13
22	How will insights from genetics translate to clinical practice in inflammatory bowel disease?. Bailliere's Best Practice and Research in Clinical Gastroenterology, 2014, 28, 387-397.	2.4	15
23	Three ulcerative colitis susceptibility loci are associated with primary sclerosing cholangitis and indicate a role for IL2, REL , and CARD9. Hepatology, 2011, 53, 1977-1985.	7.3	110
24	Analysis of SNPs with an effect on gene expression identifies UBE2L3 and BCL3 as potential new risk genes for Crohn's disease. Human Molecular Genetics, 2010, 19, 3482-3488.	2.9	92