

# Floris Imhann

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/133506/publications.pdf>

Version: 2024-02-01

22  
papers

6,981  
citations

393982

19  
h-index

610482

24  
g-index

25  
all docs

25  
docs citations

25  
times ranked

11270  
citing authors

#	ARTICLE	IF	CITATIONS
1	Whole exome sequencing analyses reveal gene-microbiota interactions in the context of IBD. <i>Gut</i> , 2021, 70, gutjnl-2019-319706.	6.1	26
2	Habitual dietary intake of IBD patients differs from population controls: a case-control study. <i>European Journal of Nutrition</i> , 2021, 60, 345-356.	1.8	22
3	Patient attitudes towards faecal sampling for gut microbiome studies and clinical care reveal positive engagement and room for improvement. <i>PLoS ONE</i> , 2021, 16, e0249405.	1.1	4
4	Long-term dietary patterns are associated with pro-inflammatory and anti-inflammatory features of the gut microbiome. <i>Gut</i> , 2021, 70, 1287-1298.	6.1	246
5	Impact of commonly used drugs on the composition and metabolic function of the gut microbiota. <i>Nature Communications</i> , 2020, 11, 362.	5.8	416
6	SLC39A8 missense variant is associated with Crohn's disease but does not have a major impact on gut microbiome composition in healthy subjects. <i>PLoS ONE</i> , 2019, 14, e0211328.	1.1	10
7	Anti-inflammatory Gut Microbial Pathways Are Decreased During Crohn's Disease Exacerbations. <i>Journal of Crohn's and Colitis</i> , 2019, 13, 1439-1449.	0.6	39
8	MOLGENIS research: advanced bioinformatics data software for non-bioinformaticians. <i>Bioinformatics</i> , 2019, 35, 1076-1078.	1.8	58
9	The 1000IBD project: multi-omics data of 1000 inflammatory bowel disease patients; data release 1. <i>BMC Gastroenterology</i> , 2019, 19, 5.	0.8	68
10	Analysis of 1135 gut metagenomes identifies sex-specific resistome profiles. <i>Gut Microbes</i> , 2019, 10, 358-366.	4.3	118
11	Gut microbiome structure and metabolic activity in inflammatory bowel disease. <i>Nature Microbiology</i> , 2019, 4, 293-305.	5.9	1,094
12	Genomic and Expression Analyses Identify a Disease-Modifying Variant for Fibrostenotic Crohn's Disease. <i>Journal of Crohn's and Colitis</i> , 2018, 12, 582-588.	0.6	16
13	Interplay of host genetics and gut microbiota underlying the onset and clinical presentation of inflammatory bowel disease. <i>Gut</i> , 2018, 67, 108-119.	6.1	590
14	Gut microbiota composition and functional changes in inflammatory bowel disease and irritable bowel syndrome. <i>Science Translational Medicine</i> , 2018, 10, .	5.8	351
15	Cohort profile: design and first results of the Dutch IBD Biobank: a prospective, nationwide biobank of patients with inflammatory bowel disease. <i>BMJ Open</i> , 2017, 7, e016695.	0.8	33
16	HSPA6 is an ulcerative colitis susceptibility factor that is induced by cigarette smoke and protects intestinal epithelial cells by stabilizing anti-apoptotic Bcl-XL. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2016, 1862, 788-796.	1.8	16
17	Population-based metagenomics analysis reveals markers for gut microbiome composition and diversity. <i>Science</i> , 2016, 352, 565-569.	6.0	1,398
18	The effect of host genetics on the gut microbiome. <i>Nature Genetics</i> , 2016, 48, 1407-1412.	9.4	672

#	ARTICLE	IF	CITATIONS
19	Proton pump inhibitors affect the gut microbiome. <i>Gut</i> , 2016, 65, 740-748.	6.1	885
20	Pooled Resequencing of 122 Ulcerative Colitis Genes in a Large Dutch Cohort Suggests Population-Specific Associations of Rare Variants in MUC2. <i>PLoS ONE</i> , 2016, 11, e0159609.	1.1	21
21	The Gut Microbiome Contributes to a Substantial Proportion of the Variation in Blood Lipids. <i>Circulation Research</i> , 2015, 117, 817-824.	2.0	534
22	Complex host genetics influence the microbiome in inflammatory bowel disease. <i>Genome Medicine</i> , 2014, 6, 107.	3.6	322