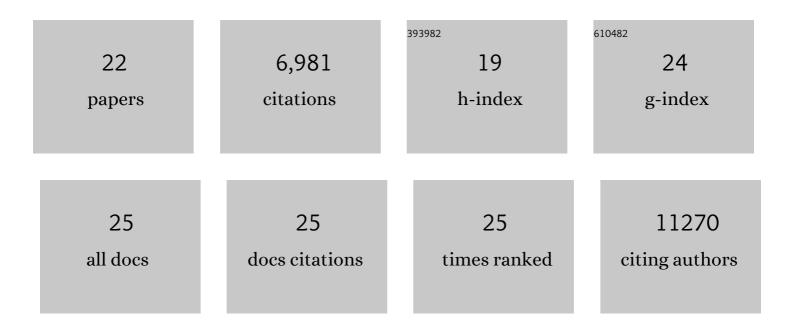
## Floris Imhann

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/133506/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Population-based metagenomics analysis reveals markers for gut microbiome composition and diversity. Science, 2016, 352, 565-569.	6.0	1,398
2	Gut microbiome structure and metabolic activity in inflammatory bowel disease. Nature Microbiology, 2019, 4, 293-305.	5.9	1,094
3	Proton pump inhibitors affect the gut microbiome. Gut, 2016, 65, 740-748.	6.1	885
4	The effect of host genetics on the gut microbiome. Nature Genetics, 2016, 48, 1407-1412.	9.4	672
5	Interplay of host genetics and gut microbiota underlying the onset and clinical presentation of inflammatory bowel disease. Gut, 2018, 67, 108-119.	6.1	590
6	The Gut Microbiome Contributes to a Substantial Proportion of the Variation in Blood Lipids. Circulation Research, 2015, 117, 817-824.	2.0	534
7	Impact of commonly used drugs on the composition and metabolic function of the gut microbiota. Nature Communications, 2020, 11, 362.	5.8	416
8	Gut microbiota composition and functional changes in inflammatory bowel disease and irritable bowel syndrome. Science Translational Medicine, 2018, 10, .	5.8	351
9	Complex host genetics influence the microbiome in inflammatory bowel disease. Genome Medicine, 2014, 6, 107.	3.6	322
10	Long-term dietary patterns are associated with pro-inflammatory and anti-inflammatory features of the gut microbiome. Gut, 2021, 70, 1287-1298.	6.1	246
11	Analysis of 1135 gut metagenomes identifies sex-specific resistome profiles. Gut Microbes, 2019, 10, 358-366.	4.3	118
12	The 1000IBD project: multi-omics data of 1000 inflammatory bowel disease patients; data release 1. BMC Gastroenterology, 2019, 19, 5.	0.8	68
13	MOLGENIS research: advanced bioinformatics data software for non-bioinformaticians. Bioinformatics, 2019, 35, 1076-1078.	1.8	58
14	Anti-inflammatory Gut Microbial Pathways Are Decreased During Crohn's Disease Exacerbations. Journal of Crohn's and Colitis, 2019, 13, 1439-1449.	0.6	39
15	Cohort profile: design and first results of the Dutch IBD Biobank: a prospective, nationwide biobank of patients with inflammatory bowel disease. BMJ Open, 2017, 7, e016695.	0.8	33
16	Whole exome sequencing analyses reveal gene–microbiota interactions in the context of IBD. Gut, 2021, 70, gutjnl-2019-319706.	6.1	26
17	Habitual dietary intake of IBD patients differs from population controls: a case–control study. European Journal of Nutrition, 2021, 60, 345-356.	1.8	22
18	Pooled Resequencing of 122 Ulcerative Colitis Genes in a Large Dutch Cohort Suggests Population-Specific Associations of Rare Variants in MUC2, PLoS ONF, 2016, 11, e0159609	1.1	21

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#	Article	IF	CITATIONS
19	HSPA6 is an ulcerative colitis susceptibility factor that is induced by cigarette smoke and protects intestinal epithelial cells by stabilizing anti-apoptotic Bcl-XL. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2016, 1862, 788-796.	1.8	16
20	Genomic and Expression Analyses Identify a Disease-Modifying Variant for Fibrostenotic Crohn's Disease. Journal of Crohn's and Colitis, 2018, 12, 582-588.	0.6	16
21	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.	1.1	10
22	Patient attitudes towards faecal sampling for gut microbiome studies and clinical care reveal positive engagement and room for improvement. PLoS ONE, 2021, 16, e0249405.	1.1	4