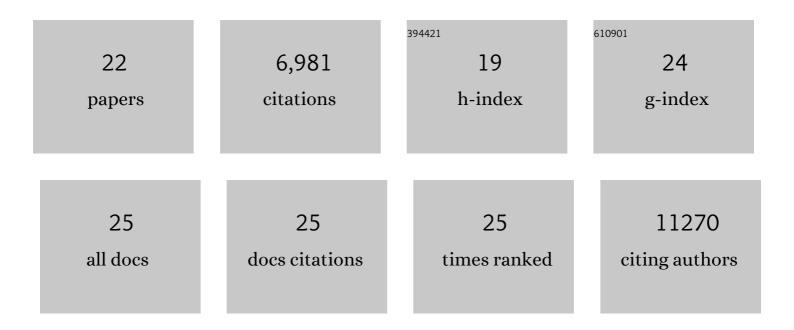
## Floris Imhann

List of Publications by Year in descending order

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



FLODIS IMHANN

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Whole exome sequencing analyses reveal gene–microbiota interactions in the context of IBD. Gut, 2021, 70, gutjnl-2019-319706.   | 12.1 | 26        |
| 2  | Habitual dietary intake of IBD patients differs from population controls: a case–control study.<br>European Journal of Nutrition, 2021, 60, 345-356.  | 3.9  | 22        |
| 3  | Patient attitudes towards faecal sampling for gut microbiome studies and clinical care reveal positive engagement and room for improvement. PLoS ONE, 2021, 16, e0249405.   | 2.5  | 4         |
| 4  | 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       |
| 5  | Impact of commonly used drugs on the composition and metabolic function of the gut microbiota.<br>Nature Communications, 2020, 11, 362.   | 12.8 | 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. PLoS ONE, 2019, 14, e0211328.   | 2.5  | 10        |
| 7  | Anti-inflammatory Gut Microbial Pathways Are Decreased During Crohn's Disease Exacerbations.<br>Journal of Crohn's and Colitis, 2019, 13, 1439-1449.  | 1.3  | 39        |
| 8  | MOLGENIS research: advanced bioinformatics data software for non-bioinformaticians.<br>Bioinformatics, 2019, 35, 1076-1078.   | 4.1  | 58        |
| 9  | The 1000IBD project: multi-omics data of 1000 inflammatory bowel disease patients; data release 1. BMC Gastroenterology, 2019, 19, 5.   | 2.0  | 68        |
| 10 | Analysis of 1135 gut metagenomes identifies sex-specific resistome profiles. Gut Microbes, 2019, 10, 358-366.   | 9.8  | 118       |
| 11 | Gut microbiome structure and metabolic activity in inflammatory bowel disease. Nature Microbiology, 2019, 4, 293-305.   | 13.3 | 1,094     |
| 12 | Genomic and Expression Analyses Identify a Disease-Modifying Variant for Fibrostenotic Crohn's<br>Disease. Journal of Crohn's and Colitis, 2018, 12, 582-588.   | 1.3  | 16        |
| 13 | Interplay of host genetics and gut microbiota underlying the onset and clinical presentation of inflammatory bowel disease. Gut, 2018, 67, 108-119.   | 12.1 | 590       |
| 14 | Gut microbiota composition and functional changes in inflammatory bowel disease and irritable bowel syndrome. Science Translational Medicine, 2018, 10, .   | 12.4 | 351       |
| 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.   | 1.9  | 33        |
| 16 | HSPA6 is an ulcerative colitis susceptibility factor that is induced by cigarette smoke and protects<br>intestinal epithelial cells by stabilizing anti-apoptotic Bcl-XL. Biochimica Et Biophysica Acta -<br>Molecular Basis of Disease, 2016, 1862, 788-796. | 3.8  | 16        |
| 17 | Population-based metagenomics analysis reveals markers for gut microbiome composition and diversity. Science, 2016, 352, 565-569.   | 12.6 | 1,398     |
| 18 | The effect of host genetics on the gut microbiome. Nature Genetics, 2016, 48, 1407-1412.  | 21.4 | 672       |

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|----|--|------|-----------|
| 19 | Proton pump inhibitors affect the gut microbiome. Gut, 2016, 65, 740-748.  | 12.1 | 885       |
| 20 | Pooled Resequencing of 122 Ulcerative Colitis Genes in a Large Dutch Cohort Suggests<br>Population-Specific Associations of Rare Variants in MUC2. PLoS ONE, 2016, 11, e0159609. | 2.5  | 21        |
| 21 | The Gut Microbiome Contributes to a Substantial Proportion of the Variation in Blood Lipids.<br>Circulation Research, 2015, 117, 817-824.  | 4.5  | 534       |
| 22 | Complex host genetics influence the microbiome in inflammatory bowel disease. Genome Medicine, 2014, 6, 107.   | 8.2  | 322       |