Trine Nielsen

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

| 30 | 12,287 | 20 | 33 |
|-------------------|-----------------------|----------------------|-----------------|
| papers | citations | h-index | g-index |
| 33 ext. papers | 15,930 ext. citations | 21. 8 avg, IF | 5.03 L-index |

| # | Paper | IF | Citations |
|----|---|------------------|-----------|
| 30 | Impairment of gut microbial biotin metabolism and host biotin status in severe obesity: effect of biotin and prebiotic supplementation on improved metabolism <i>Gut</i> , 2022 , | 19.2 | 5 |
| 29 | Microbiome and metabolome features of the cardiometabolic disease spectrum <i>Nature Medicine</i> , 2022 , | 50.5 | 4 |
| 28 | Combinatorial, additive and dose-dependent drug-microbiome associations. <i>Nature</i> , 2021 , | 50.4 | 11 |
| 27 | A Previously Undescribed Highly Prevalent Phage Identified in a Danish Enteric Virome Catalog. <i>MSystems</i> , 2021 , 6, e0038221 | 7.6 | 0 |
| 26 | Human and preclinical studies of the host-gut microbiome co-metabolite hippurate as a marker and mediator of metabolic health. <i>Gut</i> , 2021 , 70, 2105-2114 | 19.2 | 13 |
| 25 | Conjugated C-6 hydroxylated bile acids in serum relate to human metabolic health and gut Clostridia species. <i>Scientific Reports</i> , 2021 , 11, 13252 | 4.9 | 0 |
| 24 | Statin therapy is associated with lower prevalence of gut microbiota dysbiosis. <i>Nature</i> , 2020 , 581, 310-3 | B 155 0.4 | 100 |
| 23 | Describing the fecal metabolome in cryogenically collected samples from healthy participants. <i>Scientific Reports</i> , 2020 , 10, 885 | 4.9 | 8 |
| 22 | Comparative Studies of the Gut Microbiota in the Offspring of Mothers With and Without Gestational Diabetes. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020 , 10, 536282 | 5.9 | 4 |
| 21 | Imidazole propionate is increased in diabetes and associated with dietary patterns and altered microbial ecology. <i>Nature Communications</i> , 2020 , 11, 5881 | 17.4 | 29 |
| 20 | Extracellular Vesicle Encapsulated MicroRNAs in Patients with Type 2 Diabetes Are Affected by Metformin Treatment. <i>Journal of Clinical Medicine</i> , 2019 , 8, | 5.1 | 26 |
| 19 | Metformin-induced changes of the gut microbiota in healthy young men: results of a non-blinded, one-armed intervention study. <i>Diabetologia</i> , 2019 , 62, 1024-1035 | 10.3 | 79 |
| 18 | Impact of a vegan diet on the human salivary microbiota. Scientific Reports, 2018, 8, 5847 | 4.9 | 50 |
| 17 | Aberrant intestinal microbiota in individuals with prediabetes. <i>Diabetologia</i> , 2018 , 61, 810-820 | 10.3 | 163 |
| 16 | Gestational diabetes is associated with change in the gut microbiota composition in third trimester of pregnancy and postpartum. <i>Microbiome</i> , 2018 , 6, 89 | 16.6 | 155 |
| 15 | A low-gluten diet induces changes in the intestinal microbiome of healthy Danish adults. <i>Nature Communications</i> , 2018 , 9, 4630 | 17.4 | 69 |
| 14 | Population-based studies of relationships between dietary acidity load, insulin resistance and incident diabetes in Danes. <i>Nutrition Journal</i> , 2018 , 17, 91 | 4.3 | 9 |

LIST OF PUBLICATIONS

| 1 | 3 | A computational framework to integrate high-throughput comicscdatasets for the identification of potential mechanistic links. <i>Nature Protocols</i> , 2018 , 13, 2781-2800 | 18.8 | 44 |
|---|---|--|------|--------------|
| 1 | 2 | Recovery of gut microbiota of healthy adults following antibiotic exposure. <i>Nature Microbiology</i> , 2018 , 3, 1255-1265 | 26.6 | 246 |
| 1 | 1 | Dietary Assessment in the MetaCardis Study: Development and Relative Validity of an Online Food Frequency Questionnaire. <i>Journal of the Academy of Nutrition and Dietetics</i> , 2017 , 117, 878-888 | 3.9 | 18 |
| 1 | Ο | Human gut microbes impact host serum metabolome and insulin sensitivity. <i>Nature</i> , 2016 , 535, 376-81 | 50.4 | 977 |
| 9 | ١ | Transcriptional interactions suggest niche segregation among microorganisms in the human gut. <i>Nature Microbiology</i> , 2016 , 1, 16152 | 26.6 | 38 |
| 8 | | Roux-en-Y gastric bypass surgery of morbidly obese patients induces swift and persistent changes of the individual gut microbiota. <i>Genome Medicine</i> , 2016 , 8, 67 | 14.4 | 187 |
| 7 | | Alterations in fecal microbiota composition by probiotic supplementation in healthy adults: a systematic review of randomized controlled trials. <i>Genome Medicine</i> , 2016 , 8, 52 | 14.4 | 290 |
| 6 | | Disentangling type 2 diabetes and metformin treatment signatures in the human gut microbiota. <i>Nature</i> , 2015 , 528, 262-266 | 50.4 | 1107 |
| 5 | | Mechanisms in endocrinology: Gut microbiota in patients with type 2 diabetes mellitus. <i>European Journal of Endocrinology</i> , 2015 , 172, R167-77 | 6.5 | 119 |
| 4 | | Identification and assembly of genomes and genetic elements in complex metagenomic samples without using reference genomes. <i>Nature Biotechnology</i> , 2014 , 32, 822-8 | 44.5 | 624 |
| 3 | | An integrated catalog of reference genes in the human gut microbiome. <i>Nature Biotechnology</i> , 2014 , 32, 834-41 | 44.5 | 1088 |
| 2 | | Richness of human gut microbiome correlates with metabolic markers. <i>Nature</i> , 2013 , 500, 541-6 | 50.4 | 2584 |
| 1 | | Enterotypes of the human gut microbiome. <i>Nature</i> , 2011 , 473, 174-80 | 50.4 | 424 0 |