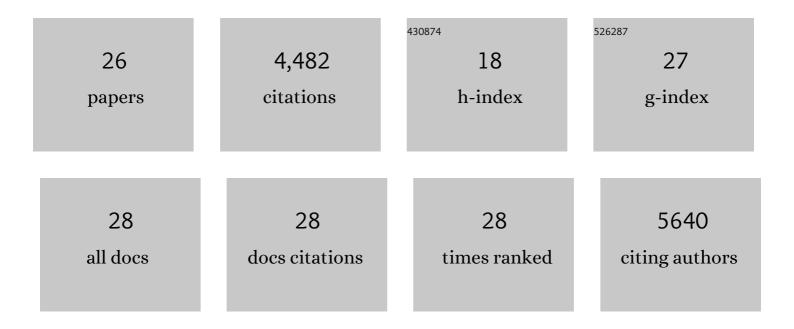
Andrew Maltez Thomas

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/377616/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Integrating taxonomic, functional, and strain-level profiling of diverse microbial communities with bioBakery 3. ELife, 2021, 10, .	6.0	808
2	Meta-analysis of fecal metagenomes reveals global microbial signatures that are specific for colorectal cancer. Nature Medicine, 2019, 25, 679-689.	30.7	734
3	Metagenomic analysis of colorectal cancer datasets identifies cross-cohort microbial diagnostic signatures and a link with choline degradation. Nature Medicine, 2019, 25, 667-678.	30.7	602
4	Microbiome connections with host metabolism and habitual diet from 1,098 deeply phenotyped individuals. Nature Medicine, 2021, 27, 321-332.	30.7	477
5	Precise phylogenetic analysis of microbial isolates and genomes from metagenomes using PhyloPhlAn 3.0. Nature Communications, 2020, 11, 2500.	12.8	368
6	Intestinal Akkermansia muciniphila predicts clinical response to PD-1 blockade in patients with advanced non-small-cell lung cancer. Nature Medicine, 2022, 28, 315-324.	30.7	225
7	Butyrate Protects Mice from Clostridium difficile-Induced Colitis through an HIF-1-Dependent Mechanism. Cell Reports, 2019, 27, 750-761.e7.	6.4	212
8	Microbial community structure and dynamics in thermophilic composting viewed through metagenomics and metatranscriptomics. Scientific Reports, 2016, 6, 38915.	3.3	183
9	Cross-cohort gut microbiome associations with immune checkpoint inhibitor response in advanced melanoma. Nature Medicine, 2022, 28, 535-544.	30.7	158
10	Tissue-Associated Bacterial Alterations in Rectal Carcinoma Patients Revealed by 16S rRNA Community Profiling. Frontiers in Cellular and Infection Microbiology, 2016, 6, 179.	3.9	125
11	Multiple levels of the unknown in microbiome research. BMC Biology, 2019, 17, 48.	3.8	101
12	Faecal microbiota transplantation for the treatment of diarrhoea induced by tyrosine-kinase inhibitors in patients with metastatic renal cell carcinoma. Nature Communications, 2020, 11, 4333.	12.8	82
13	Intestinal microbiota influences clinical outcome and side effects of early breast cancer treatment. Cell Death and Differentiation, 2021, 28, 2778-2796.	11.2	72
14	Alcohol and tobacco consumption affects bacterial richness in oral cavity mucosa biofilms. BMC Microbiology, 2014, 14, 250.	3.3	71
15	Altered Fecal Small RNA Profiles in Colorectal Cancer Reflect Gut Microbiome Composition in Stool Samples. MSystems, 2019, 4, .	3.8	59
16	Gut Microbiota Profile of Obese Diabetic Women Submitted to Roux-en-Y Gastric Bypass and Its Association with Food Intake and Postoperative Diabetes Remission. Nutrients, 2020, 12, 278.	4.1	47
17	Prophylactic Supplementation of Bifidobacterium longum 51A Protects Mice from Ovariectomy-Induced Exacerbated Allergic Airway Inflammation and Airway Hyperresponsiveness. Frontiers in Microbiology, 2017, 8, 1732.	3.5	27
18	Tributyrin Attenuates Metabolic and Inflammatory Changes Associated with Obesity through a GPR109A-Dependent Mechanism. Cells, 2020, 9, 2007.	4.1	25

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#	Article	IF	CITATIONS
19	Deletion and low expression of NFKBIA are associated with poor prognosis in lower-grade glioma patients. Scientific Reports, 2016, 6, 24160.	3.3	17
20	A probiotic has differential effects on allergic airway inflammation in A/J and C57BL/6 mice and is correlated with the gut microbiome. Microbiome, 2021, 9, 134.	11.1	14
21	Complete genome sequence and analysis of Alcaligenes faecalis strain Mc250, a new potential plant bioinoculant. PLoS ONE, 2020, 15, e0241546.	2.5	9
22	Genes Encoding Microbial Acyl Coenzyme A Binding Protein/Diazepam-Binding Inhibitor Orthologs Are Rare in the Human Gut Microbiome and Show No Links to Obesity. Applied and Environmental Microbiology, 2021, 87, e0047121.	3.1	7
23	Genomics and epidemiology for gastric adenocarcinomas (GE4GAC): a Brazilian initiative to study gastric cancer. Applied Cancer Research, 2019, 39, .	1.0	5
24	Maternal supplementation with a synbiotic has distinct outcomes on offspring gut microbiota formation in A/J and C57BL/6 mice, differentially affecting airway inflammatory cell infiltration and mucus production. Journal of Functional Foods, 2019, 61, 103496.	3.4	4
25	Comparative Metagenomics. Methods in Molecular Biology, 2018, 1704, 243-260.	0.9	2
26	Microbiomes of Field-Grown Maize and Soybean in Southeastern and Central Brazil Inferred by High-Throughput 16S and Internal Transcribed Spacer Amplicon Sequencing. Microbiology Resource Announcements, 2021, 10, e0052821.	0.6	1