Huanzi Zhong

List of Publications by Year in descending order

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394421 642732 9,355 24 19 23 citations h-index g-index papers 33 33 33 14306 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Dynamics and Stabilization of the Human Gut Microbiome during the First Year of Life. Cell Host and Microbe, 2015, 17, 690-703.	11.0	2,276
2	An integrated catalog of reference genes in the human gut microbiome. Nature Biotechnology, 2014, 32, 834-841.	17.5	1,664
3	The oral and gut microbiomes are perturbed in rheumatoid arthritis and partly normalized after treatment. Nature Medicine, 2015, 21, 895-905.	30.7	1,306
4	Gut microbiome and serum metabolome alterations in obesity and after weight-loss intervention. Nature Medicine, 2017, 23, 859-868.	30.7	1,074
5	The gut microbiome in atherosclerotic cardiovascular disease. Nature Communications, 2017, 8, 845.	12.8	1,029
6	The microbiota continuum along the female reproductive tract and its relation to uterine-related diseases. Nature Communications, 2017, 8, 875.	12.8	572
7	Analyses of gut microbiota and plasma bile acids enable stratification of patients for antidiabetic treatment. Nature Communications, 2017, 8, 1785.	12.8	312
8	Gut microbiome-related effects of berberine and probiotics on type 2 diabetes (the PREMOTE study). Nature Communications, 2020, 11, 5015.	12.8	184
9	Assessment of the cPAS-based BGISEQ-500 platform for metagenomic sequencing. GigaScience, 2018, 7, 1-8.	6.4	168
10	Impact of early events and lifestyle on the gut microbiota and metabolic phenotypes in young school-age children. Microbiome, 2019, 7, 2.	11.1	135
11	Multiple approaches for massively parallel sequencing of SARS-CoV-2 genomes directly from clinical samples. Genome Medicine, 2020, 12, 57.	8.2	104
12	Distinct gut metagenomics and metaproteomics signatures in prediabetics and treatment-na \tilde{A} -ve type 2 diabetics. EBioMedicine, 2019, 47, 373-383.	6.1	101
13	Sex- and age-related trajectories of the adult human gut microbiota shared across populations of different ethnicities. Nature Aging, 2021, 1, 87-100.	11.6	86
14	Lipidomic profiling reveals distinct differences in plasma lipid composition in healthy, prediabetic, and type 2 diabetic individuals. GigaScience, 2017, 6, 1-12.	6.4	49
15	A genome-wide association study for gut metagenome in Chinese adults illuminates complex diseases. Cell Discovery, 2021, 7, 9.	6.7	49
16	Effect of Caloric Restriction on BMI, Gut Microbiota, and Blood Amino Acid Levels in Non-Obese Adults. Nutrients, 2020, 12, 631.	4.1	36
17	Assessment of fecal DNA extraction protocols for metagenomic studies. GigaScience, 2020, 9, .	6.4	35
18	A catalog of microbial genes from the bovine rumen unveils a specialized and diverse biomass-degrading environment. GigaScience, 2020, 9, .	6.4	35

#	Article	IF	CITATION
19	Characterization of respiratory microbial dysbiosis in hospitalized COVID-19 patients. Cell Discovery, 2021, 7, 23.	6.7	34
20	Combined berberine and probiotic treatment as an effective regimen for improving postprandial hyperlipidemia in type 2 diabetes patients: a double blinded placebo controlled randomized study. Gut Microbes, 2022, 14, 2003176.	9.8	27
21	A transomic cohort as a reference point for promoting a healthy human gut microbiome. Medicine in Microecology, 2021, 8, 100039.	1.6	24
22	An Expanded Gene Catalog of Mouse Gut Metagenomes. MSphere, 2021, 6, .	2.9	13
23	Distinct Functional Metagenomic Markers Predict the Responsiveness to Anti-PD-1 Therapy in Chinese Non-Small Cell Lung Cancer Patients. Frontiers in Oncology, 2022, 12, 837525.	2.8	6
24	M-GWAS for the Gut Microbiome in Chinese Adults Illuminates on Complex Diseases. SSRN Electronic Journal, $0, \dots$	0.4	4