Huanzi Zhong

List of Publications by Year in descending order

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Version: 2024-02-01

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	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
2	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
3	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
4	An Expanded Gene Catalog of Mouse Gut Metagenomes. MSphere, 2021, 6, .	2.9	13
5	A genome-wide association study for gut metagenome in Chinese adults illuminates complex diseases. Cell Discovery, 2021, 7, 9.	6.7	49
6	Characterization of respiratory microbial dysbiosis in hospitalized COVID-19 patients. Cell Discovery, 2021, 7, 23.	6.7	34
7	A transomic cohort as a reference point for promoting a healthy human gut microbiome. Medicine in Microecology, 2021, 8, 100039.	1.6	24
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 fecal DNA extraction protocols for metagenomic studies. GigaScience, 2020, 9, .	6.4	35
10	A catalog of microbial genes from the bovine rumen unveils a specialized and diverse biomass-degrading environment. GigaScience, 2020, 9, .	6.4	35
11	Effect of Caloric Restriction on BMI, Gut Microbiota, and Blood Amino Acid Levels in Non-Obese Adults. Nutrients, 2020, 12, 631.	4.1	36
12	Multiple approaches for massively parallel sequencing of SARS-CoV-2 genomes directly from clinical samples. Genome Medicine, 2020, 12, 57.	8.2	104
13	Distinct gut metagenomics and metaproteomics signatures in prediabetics and treatment-naÃ-ve type 2 diabetics. EBioMedicine, 2019, 47, 373-383.	6.1	101
14	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
15	Assessment of the cPAS-based BGISEQ-500 platform for metagenomic sequencing. GigaScience, 2018, 7, 1-8.	6.4	168
16	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
17	Gut microbiome and serum metabolome alterations in obesity and after weight-loss intervention. Nature Medicine, 2017, 23, 859-868.	30.7	1,074
18	The gut microbiome in atherosclerotic cardiovascular disease. Nature Communications, 2017, 8, 845.	12.8	1,029

#	Article	IF	CITATIONS
19	The microbiota continuum along the female reproductive tract and its relation to uterine-related diseases. Nature Communications, 2017, 8, 875.	12.8	572
20	Analyses of gut microbiota and plasma bile acids enable stratification of patients for antidiabetic treatment. Nature Communications, 2017, 8, 1785.	12.8	312
21	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
22	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
23	An integrated catalog of reference genes in the human gut microbiome. Nature Biotechnology, 2014, 32, 834-841.	17.5	1,664
24	M-GWAS for the Gut Microbiome in Chinese Adults Illuminates on Complex Diseases. SSRN Electronic Journal, $0, , .$	0.4	4