

Chenhong Zhang

List of Publications by Citations

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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.

46
papers

4,521
citations

22
h-index

55
g-index

55
ext. papers

6,208
ext. citations

8.7
avg, IF

5.27
L-index

#	Paper	IF	Citations
46	Gut bacteria selectively promoted by dietary fibers alleviate type 2 diabetes. <i>Science</i> , 2018 , 359, 1151-1156	11.9	904
45	Interactions between gut microbiota, host genetics and diet relevant to development of metabolic syndromes in mice. <i>ISME Journal</i> , 2010 , 4, 232-41	11.9	633
44	Modulation of gut microbiota during probiotic-mediated attenuation of metabolic syndrome in high fat diet-fed mice. <i>ISME Journal</i> , 2015 , 9, 1-15	11.9	536
43	Structural modulation of gut microbiota in life-long calorie-restricted mice. <i>Nature Communications</i> , 2013 , 4, 2163	17.4	305
42	Structural resilience of the gut microbiota in adult mice under high-fat dietary perturbations. <i>ISME Journal</i> , 2012 , 6, 1848-57	11.9	294
41	Structural modulation of gut microbiota during alleviation of type 2 diabetes with a Chinese herbal formula. <i>ISME Journal</i> , 2015 , 9, 552-62	11.9	267
40	Fiber-utilizing capacity varies in Prevotella- versus Bacteroides-dominated gut microbiota. <i>Scientific Reports</i> , 2017 , 7, 2594	4.9	216
39	Dietary Modulation of Gut Microbiota Contributes to Alleviation of Both Genetic and Simple Obesity in Children. <i>EBioMedicine</i> , 2015 , 2, 968-84	8.8	198
38	Structural Alteration of Gut Microbiota during the Amelioration of Human Type 2 Diabetes with Hyperlipidemia by Metformin and a Traditional Chinese Herbal Formula: a Multicenter, Randomized, Open Label Clinical Trial. <i>MBio</i> , 2018 , 9,	7.8	139
37	Dysbiosis of Gut Microbiota Associated with Clinical Parameters in Polycystic Ovary Syndrome. <i>Frontiers in Microbiology</i> , 2017 , 8, 324	5.7	121
36	Strain-Specific Anti-inflammatory Properties of Two Strains on Chronic Colitis in Mice. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019 , 9, 239	5.9	118
35	Accelerated dysbiosis of gut microbiota during aggravation of DSS-induced colitis by a butyrate-producing bacterium. <i>Scientific Reports</i> , 2016 , 6, 27572	4.9	99
34	Remodelling of the gut microbiota by hyperactive NLRP3 induces regulatory T cells to maintain homeostasis. <i>Nature Communications</i> , 2017 , 8, 1896	17.4	96
33	Alterations in the gut microbiome and metabolism with coronary artery disease severity. <i>Microbiome</i> , 2019 , 7, 68	16.6	93
32	Predominant gut Lactobacillus murinus strain mediates anti-inflammaging effects in calorie-restricted mice. <i>Microbiome</i> , 2018 , 6, 54	16.6	65
31	Strain-level dissection of the contribution of the gut microbiome to human metabolic disease. <i>Genome Medicine</i> , 2016 , 8, 41	14.4	59
30	Initial gut microbiota structure affects sensitivity to DSS-induced colitis in a mouse model. <i>Science China Life Sciences</i> , 2018 , 61, 762-769	8.5	49

29	The Association between Cardiorespiratory Fitness and Gut Microbiota Composition in Premenopausal Women. <i>Nutrients</i> , 2017 , 9,	6.7	38
28	Genomic Microdiversity of Underlying Differential Strain-Level Responses to Dietary Carbohydrate Intervention. <i>MBio</i> , 2017 , 8,	7.8	31
27	A More Robust Gut Microbiota in Calorie-Restricted Mice Is Associated with Attenuated Intestinal Injury Caused by the Chemotherapy Drug Cyclophosphamide. <i>MBio</i> , 2019 , 10,	7.8	25
26	Genetically Obese Human Gut Microbiota Induces Liver Steatosis in Germ-Free Mice Fed on Normal Diet. <i>Frontiers in Microbiology</i> , 2018 , 9, 1602	5.7	25
25	Fecal menaquinone profiles of overweight adults are associated with gut microbiota composition during a gut microbiota-targeted dietary intervention. <i>American Journal of Clinical Nutrition</i> , 2015 , 102, 84-93	7	24
24	Diminution of the gut resistome after a gut microbiota-targeted dietary intervention in obese children. <i>Scientific Reports</i> , 2016 , 6, 24030	4.9	22
23	Guild-based analysis for understanding gut microbiome in human health and diseases. <i>Genome Medicine</i> , 2021 , 13, 22	14.4	22
22	The structural alteration of gut microbiota in low-birth-weight mice undergoing accelerated postnatal growth. <i>Scientific Reports</i> , 2016 , 6, 27780	4.9	18
21	Causality in dietary interventions-building a case for gut microbiota. <i>Genome Medicine</i> , 2018 , 10, 62	14.4	17
20	Timing of Calorie Restriction in Mice Impacts Host Metabolic Phenotype with Correlative Changes in Gut Microbiota. <i>MSystems</i> , 2019 , 4,	7.6	14
19	Strain Promoted by a High-Fiber Diet in Genetic Obese Child Alleviates Lipid Metabolism and Modifies Gut Microbiota in Mice on a Western Diet. <i>Microorganisms</i> , 2020 , 8,	4.9	11
18	Fungal and bacterial microbiome dysbiosis and imbalance of trans-kingdom network in asthma. <i>Clinical and Translational Allergy</i> , 2020 , 10, 42	5.2	10
17	DNA Phosphorothioate Modifications Are Widely Distributed in the Human Microbiome. <i>Biomolecules</i> , 2020 , 10,	5.9	8
16	Association Between Gut Microbiota and Symptomatic Hand Osteoarthritis: Data From the Xiangya Osteoarthritis Study. <i>Arthritis and Rheumatology</i> , 2021 , 73, 1656-1662	9.5	8
15	Regulated Inflammation and Lipid Metabolism in Colon mRNA Expressions of Obese Germfree Mice Responding to B29 Combined with the High Fat Diet. <i>Frontiers in Microbiology</i> , 2016 , 7, 1786	5.7	7
14	miRNA-Gene Regulatory Network in Gnotobiotic Mice Stimulated by Dysbiotic Gut Microbiota Transplanted From a Genetically Obese Child. <i>Frontiers in Microbiology</i> , 2019 , 10, 1517	5.7	5
13	Minimizing spurious features in 16S rRNA gene amplicon sequencing		5
12	Clinical characteristics associated with the properties of gut microbiota in peritoneal dialysis patients. <i>Peritoneal Dialysis International</i> , 2021 , 41, 298-306	2.8	5

11	Characteristics of the Gut Microbiota and Metabolism in Patients With Latent Autoimmune Diabetes in Adults: A Case-Control Study. <i>Diabetes Care</i> , 2021 , 44, 2738-2746	14.6	4
10	Ketogenic Diets Induced Glucose Intolerance and Lipid Accumulation in Mice with Alterations in Gut Microbiota and Metabolites. <i>MBio</i> , 2021 , 12,	7.8	4
9	Ginsenoside Rb1 Improves Metabolic Disorder in High-Fat Diet-Induced Obese Mice Associated With Modulation of Gut Microbiota.. <i>Frontiers in Microbiology</i> , 2022 , 13, 826487	5.7	4
8	Insights into gut microbiome and its functional pathways in asthma patients through high-throughput sequencing. <i>Future Microbiology</i> , 2021 , 16, 421-438	2.9	3
7	Active phase prebiotic feeding alters gut microbiota, induces weight-independent alleviation of hepatic steatosis and serum cholesterol in high-fat diet-fed mice. <i>Computational and Structural Biotechnology Journal</i> , 2021 , 19, 448-458	6.8	3
6	The effect of calorie intake, fasting, and dietary composition on metabolic health and gut microbiota in mice. <i>BMC Biology</i> , 2021 , 19, 51	7.3	3
5	High-Fiber Diet or Combined With Acarbose Alleviates Heterogeneous Phenotypes of Polycystic Ovary Syndrome by Regulating Gut Microbiota.. <i>Frontiers in Endocrinology</i> , 2021 , 12, 806331	5.7	2
4	Gut Bacteria Shared by Children and Their Mothers Associate with Developmental Level and Social Deficits in Autism Spectrum Disorder. <i>MSphere</i> , 2020 , 5,	5	2
3	Quantification of Human Oral and Fecal by Use of Quantitative Real-Time PCR Targeting the Gene. <i>Frontiers in Microbiology</i> , 2019 , 10, 2910	5.7	2
2	A randomized controlled trial for response of microbiome network to exercise and diet intervention in patients with nonalcoholic fatty liver disease.. <i>Nature Communications</i> , 2022 , 13, 2555	17.4	2
1	A Comprehensive Analysis of Genomics and Metagenomics in a Heterozygote Familial Hypercholesterolemia Family. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021 , 11, 605954	5.9	1