

Yueqiong Ni

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.

14
papers

583
citations

9
h-index

15
g-index

15
ext. papers

862
ext. citations

9.7
avg, IF

3.91
L-index

#	Paper	IF	Citations
14	Probiotics modulated gut microbiota suppresses hepatocellular carcinoma growth in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E1306-15	11.5	288
13	Gut Microbiome Fermentation Determines the Efficacy of Exercise for Diabetes Prevention. <i>Cell Metabolism</i> , 2020 , 31, 77-91.e5	24.6	95
12	Predictable modulation of cancer treatment outcomes by the gut microbiota. <i>Microbiome</i> , 2020 , 8, 28	16.6	44
11	The Environmental Exposures and Inner- and Intercity Traffic Flows of the Metro System May Contribute to the Skin Microbiome and Resistome. <i>Cell Reports</i> , 2018 , 24, 1190-1202.e5	10.6	32
10	A Molecular-Level Landscape of Diet-Gut Microbiome Interactions: Toward Dietary Interventions Targeting Bacterial Genes. <i>MBio</i> , 2015 , 6, e01263-15	7.8	29
9	COMAN: a web server for comprehensive metatranscriptomics analysis. <i>BMC Genomics</i> , 2016 , 17, 622	4.5	26
8	Developing a molecular roadmap of drug-food interactions. <i>PLoS Computational Biology</i> , 2015 , 11, e1004048	5.048	22
7	NutriChem 2.0: exploring the effect of plant-based foods on human health and drug efficacy. <i>Database: the Journal of Biological Databases and Curation</i> , 2017 , 2017,	5	11
6	Designing Dietary Recommendations Using System Level Interactomics Analysis and Network-Based Inference. <i>Frontiers in Physiology</i> , 2017 , 8, 753	4.6	9
5	A metagenome-wide association study of gut microbiome and visceral fat accumulation. <i>Computational and Structural Biotechnology Journal</i> , 2020 , 18, 2596-2609	6.8	8
4	Distinct composition and metabolic functions of human gut microbiota are associated with cachexia in lung cancer patients. <i>ISME Journal</i> , 2021 , 15, 3207-3220	11.9	8
3	Metabolic modeling predicts specific gut bacteria as key determinants for <i>Candida albicans</i> colonization levels. <i>ISME Journal</i> , 2021 , 15, 1257-1270	11.9	8
2	An integrative understanding of the large metabolic shifts induced by antibiotics in critical illness. <i>Gut Microbes</i> , 2021 , 13, 1993598	8.8	2
1	Gut microbiome functionality might be associated with exercise tolerance and recurrence of resected early-stage lung cancer patients. <i>PLoS ONE</i> , 2021 , 16, e0259898	3.7	1