

# Suisha Liang

## List of Publications by Year in descending order

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14  
papers

10,119  
citations

840776

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1058476

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14  
docs citations

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times ranked

14997  
citing authors

#	ARTICLE	IF	CITATIONS
1	Role of gut microbiota in travel-related acquisition of extended spectrum $\beta$ -lactamase-producing Enterobacteriaceae. <i>Journal of Travel Medicine</i> , 2021, 28, .	3.0	14
2	A transomic cohort as a reference point for promoting a healthy human gut microbiome. <i>Medicine in Microecology</i> , 2021, 8, 100039.	1.6	24
3	Metagenomic Survey Reveals More Diverse and Abundant Antibiotic Resistance Genes in Municipal Wastewater Than Hospital Wastewater. <i>Frontiers in Microbiology</i> , 2021, 12, 712843.	3.5	6
4	Dairy consumption and physical fitness tests associated with fecal microbiome in a Chinese cohort. <i>Medicine in Microecology</i> , 2021, 9, 100038.	1.6	6
5	Disease trends in a young Chinese cohort according to fecal metagenome and plasma metabolites. <i>Medicine in Microecology</i> , 2021, , 100037.	1.6	2
6	The Gut Microbiota Profile in Children with Prader-Willi Syndrome. <i>Genes</i> , 2020, 11, 904.	2.4	18
7	Impact of outdoor nature-related activities on gut microbiota, fecal serotonin, and perceived stress in preschool children: the Play&Grow randomized controlled trial. <i>Scientific Reports</i> , 2020, 10, 21993.	3.3	58
8	Establishment of a <i>Macaca fascicularis</i> gut microbiome gene catalog and comparison with the human, pig, and mouse gut microbiomes. <i>GigaScience</i> , 2018, 7, .	6.4	53
9	The gut microbiome in atherosclerotic cardiovascular disease. <i>Nature Communications</i> , 2017, 8, 845.	12.8	1,029
10	A reference gene catalogue of the pig gut microbiome. <i>Nature Microbiology</i> , 2016, 1, 16161.	13.3	416
11	Gut microbiome development along the colorectal adenoma-carcinoma sequence. <i>Nature Communications</i> , 2015, 6, 6528.	12.8	1,062
12	A catalog of the mouse gut metagenome. <i>Nature Biotechnology</i> , 2015, 33, 1103-1108.	17.5	422
13	An integrated catalog of reference genes in the human gut microbiome. <i>Nature Biotechnology</i> , 2014, 32, 834-841.	17.5	1,664
14	A metagenome-wide association study of gut microbiota in type 2 diabetes. <i>Nature</i> , 2012, 490, 55-60.	27.8	5,345