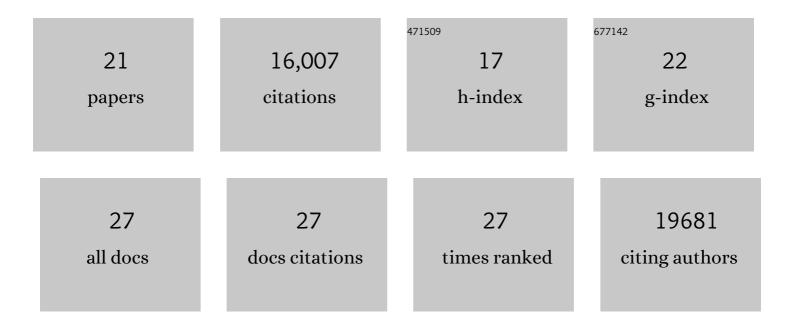
## Lingjing Jiang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1775938/publications.pdf Version: 2024-02-01



LINCUNG JUNG

#	Article	IF	CITATIONS
1	A semiparametric model for betweenâ€subject attributes: Applications to betaâ€diversity of microbiome data. Biometrics, 2022, 78, 950-962.	1.4	5
2	Utilizing stability criteria in choosing feature selection methods yields reproducible results in microbiome data. Biometrics, 2022, 78, 1155-1167.	1.4	4
3	Multiomic Analyses of Nascent Preterm Infant Microbiomes Differentiation Suggest Opportunities for Targeted Intervention. Advanced Biology, 2022, 6, .	2.5	4
4	Context-aware dimensionality reduction deconvolutes gut microbial community dynamics. Nature Biotechnology, 2021, 39, 165-168.	17.5	61
5	SARS-CoV-2 detection status associates with bacterial community composition in patients and the hospital environment. Microbiome, 2021, 9, 132.	11.1	37
6	Naturalization of the microbiota developmental trajectory of Cesarean-born neonates after vaginal seeding. Med, 2021, 2, 951-964.e5.	4.4	37
7	Effects of the microalgae Chlamydomonas on gastrointestinal health. Journal of Functional Foods, 2020, 65, 103738.	3.4	66
8	Nutrition and the Gut Microbiota in 10- to 18-Month-Old Children Living in Urban Slums of Mumbai, India. MSphere, 2020, 5, .	2.9	20
9	Vitamin D metabolites and the gut microbiome in older men. Nature Communications, 2020, 11, 5997.	12.8	88
10	Host variables confound gut microbiota studies of human disease. Nature, 2020, 587, 448-454.	27.8	324
11	High-Resolution Longitudinal Dynamics of the Cystic Fibrosis Sputum Microbiome and Metabolome through Antibiotic Therapy. MSystems, 2020, 5, .	3.8	47
12	Human Skin, Oral, and Gut Microbiomes Predict Chronological Age. MSystems, 2020, 5, .	3.8	80
13	QIIME 2 Enables Comprehensive Endâ€ŧoâ€End Analysis of Diverse Microbiome Data and Comparative Studies with Publicly Available Data. Current Protocols in Bioinformatics, 2020, 70, e100.	25.8	212
14	Reproducible, interactive, scalable and extensible microbiome data science using QIIME 2. Nature Biotechnology, 2019, 37, 852-857.	17.5	11,167
15	Calour: an Interactive, Microbe-Centric Analysis Tool. MSystems, 2019, 4, .	3.8	28
16	Evaluating Metagenomic Prediction of the Metaproteome in a 4.5-Year Study of a Patient with Crohn's Disease. MSystems, 2019, 4, .	3.8	40
17	Phylogenetic Placement of Exact Amplicon Sequences Improves Associations with Clinical Information. MSystems, 2018, 3, .	3.8	376
18	American Gut: an Open Platform for Citizen Science Microbiome Research. MSystems, 2018, 3, .	3.8	604

LINGJING JIANG

#	Article	IF	CITATIONS
19	An Elegan(t) Screen for Drug-Microbe Interactions. Cell Host and Microbe, 2017, 21, 555-556.	11.0	2
20	A communal catalogue reveals Earth's multiscale microbial diversity. Nature, 2017, 551, 457-463.	27.8	1,942
21	Discrete False-Discovery Rate Improves Identification of Differentially Abundant Microbes. MSystems, 2017, 2, .	3.8	73