Pajau Vangay

List of Publications by Year
in descending order

24 papers	4,323 citations	430874 18 h-index	610901 24 g-index
26 all docs	26 docs citations	26 times ranked	7947 citing authors

#	Article	IF	Citations
1	Microbiome Metadata Standards: Report of the National Microbiome Data Collaborativeâ€™s Workshop and Follow-On Activities. MSystems, 2021, 6, .	3.8	28
2	Participatory Microbiome Research With Hmong and Karen Communities: Lessons Learned. Journal of Participatory Research Methods, 2021, 2, .	0.9	1
3	US Immigration Is Associated With Rapid and Persistent Acquisition of Antibiotic Resistance Genes in the Gut. Clinical Infectious Diseases, 2020, 71, 419-421.	5.8	10
4	Wild primate microbiomes prevent weight gain in germ-free mice. Animal Microbiome, 2020, 2, 16.	3.8	7
5	The National Microbiome Data Collaborative: enabling microbiome science. Nature Reviews Microbiology, 2020, 18, 313-314.	28.6	42
6	Dietary Patterns Correspond with Microbiome Composition (FS07-02-19). Current Developments in Nutrition, 2019, 3, nzz040.FS07-02-19.	0.3	0
7	Daily Sampling Reveals Personalized Diet-Microbiome Associations in Humans. Cell Host and Microbe, 2019, 25, 789-802.e5.	11.0	441
8	Microbiome Learning Repo (ML Repo): A public repository of microbiome regression and classification tasks. GigaScience, 2019, 8, .	6.4	54
9	US Immigration Westernizes the Human Gut Microbiome. Cell, 2018, 175, 962-972.e10.	28.9	511
10	Associations Between Nutrition, Gut Microbiome, and Health in A Novel Nonhuman Primate Model. Scientific Reports, 2018, 8, 11159.	3.3	60
11	Substituting whole grains for refined grains in a 6-wk randomized trial favorably affects energy-balance metrics in healthy men and postmenopausal women. American Journal of Clinical Nutrition, 2017, 105, 589-599.	4.7	74
12	Substituting whole grains for refined grains in a 6-wk randomized trial has a modest effect on gut microbiota and immune and inflammatory markers of healthy adults. American Journal of Clinical Nutrition, 2017, 105, 635-650.	4.7	203
13	Fecal concentrations of bacterially derived vitamin K forms are associated with gut microbiota composition but not plasma or fecal cytokine concentrations in healthy adults. American Journal of Clinical Nutrition, 2017, 106, 1052-1061.	4.7	71
14	Stable Engraftment of Bifidobacterium longum AH1206 in the Human Gut Depends on Individualized Features of the Resident Microbiome. Cell Host and Microbe, 2016, 20, 515-526.	11.0	337
15	Captivity humanizes the primate microbiome. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 10376-10381.	7.1	369
16	Systematic improvement of amplicon marker gene methods for increased accuracy in microbiome studies. Nature Biotechnology, 2016, 34, 942-949.	17.5	623
17	Antibiotic-mediated gut microbiome perturbation accelerates development of type 1 diabetes in mice. Nature Microbiology, 2016, 1, 16140.	13.3	275
18	Chemotherapyâ€driven dysbiosis in the intestinal microbiome. Alimentary Pharmacology and Therapeutics, 2015, 42, 515-528.	3.7	334

#	Article	IF	Citations
19	Antibiotics, Pediatric Dysbiosis, and Disease. Cell Host and Microbe, 2015, 17, 553-564.	11.0	428
20	The guts of obesity: progress and challenges in linking gut microbes to obesity. Discovery Medicine, 2015, 19, 81-8.	0.5	8
21	Complex host genetics influence the microbiome in inflammatory bowel disease. Genome Medicine, 2014, 6, 107.	8.2	322
22	Multi-omics analysis of inflammatory bowel disease. Immunology Letters, 2014, 162, 62-68.	2.5	42
23	Classification of Listeria monocytogenes Persistence in Retail Delicatessen Environments Using Expert Elicitation and Machine Learning. Risk Analysis, 2014, 34, 1830-1845.	2.7	7
24	Food Microbe Tracker: A Web-Based Tool for Storage and Comparison of Food-Associated Microbes. Journal of Food Protection, 2013, 76, 283-294.	1.7	70