

Tonya Ward

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

1,592
citations

759055

12
h-index

1125617

13
g-index

14
all docs

14
docs citations

14
times ranked

3091
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of shared and disease-specific host gene-microbiome associations across human diseases using multi-omic integration. <i>Nature Microbiology</i> , 2022, 7, 780-795.	5.9	57
2	Multi-Omics Analyses Show Disease, Diet, and Transcriptome Interactions With the Virome. <i>Gastroenterology</i> , 2021, 161, 1194-1207.e8.	0.6	28
3	Longitudinal Multi-omics Reveals Subset-Specific Mechanisms Underlying Irritable Bowel Syndrome. <i>Cell</i> , 2020, 182, 1460-1473.e17.	13.5	217
4	Fecal microbiota transplantation reverses antibiotic and chemotherapy-induced gut dysbiosis in mice. <i>Scientific Reports</i> , 2018, 8, 6219.	1.6	99
5	Vaccination Against <i>Lawsonia intracellularis</i> Decreases Shedding of <i>Salmonella enterica</i> serovar Typhimurium in Co-Infected Pigs and Alters the Gut Microbiome. <i>Scientific Reports</i> , 2018, 8, 2857.	1.6	24
6	Associations Between Nutrition, Gut Microbiome, and Health in A Novel Nonhuman Primate Model. <i>Scientific Reports</i> , 2018, 8, 11159.	1.6	60
7	Intergenerational transfer of antibiotic-perturbed microbiota enhances colitis in susceptible mice. <i>Nature Microbiology</i> , 2018, 3, 234-242.	5.9	118
8	Advances in the use of genetically modified plant biomass for biodiesel generation. <i>Biofuels, Bioproducts and Biorefining</i> , 2017, 11, 749-764.	1.9	11
9	Diverse bacterial communities exist on canine skin and are impacted by cohabitation and time. <i>PeerJ</i> , 2017, 5, e3075.	0.9	30
10	Captivity humanizes the primate microbiome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 10376-10381.	3.3	369
11	Pretreatment gut microbiome predicts chemotherapy-related bloodstream infection. <i>Genome Medicine</i> , 2016, 8, 49.	3.6	136
12	Antibiotics, Pediatric Dysbiosis, and Disease. <i>Cell Host and Microbe</i> , 2015, 17, 553-564.	5.1	428