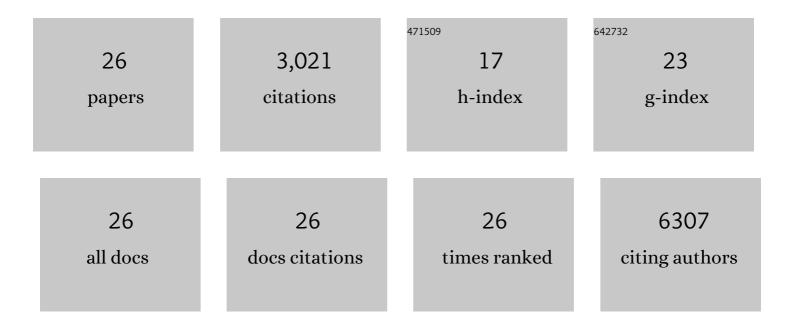
Aubrey G Bailey

List of Publications by Year in descending order

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ALIRDEN C. RALLEN

#	Article	IF	CITATIONS
1	Association Between Breast Milk Bacterial Communities and Establishment and Development of the Infant Gut Microbiome. JAMA Pediatrics, 2017, 171, 647.	6.2	749
2	Inflammation, Antibiotics, and Diet as Environmental Stressors of the Gut Microbiome in Pediatric Crohn's Disease. Cell Host and Microbe, 2015, 18, 489-500.	11.0	646
3	Comparison of placenta samples with contamination controls does not provide evidence for a distinct placenta microbiota. Microbiome, 2016, 4, 29.	11.1	447
4	Virus-helminth coinfection reveals a microbiota-independent mechanism of immunomodulation. Science, 2014, 345, 578-582.	12.6	238
5	A role for bacterial urease in gut dysbiosis and Crohn's disease. Science Translational Medicine, 2017, 9, .	12.4	171
6	Human and rat gut microbiome composition is maintained following sleep restriction. Proceedings of the United States of America, 2017, 114, E1564-E1571.	7.1	106
7	Peptidoglycan from the gut microbiota governs the lifespan of circulating phagocytes at homeostasis. Blood, 2016, 127, 2460-2471.	1.4	88
8	INSPIIRED: A Pipeline for Quantitative Analysis of Sites of New DNA Integration in Cellular Genomes. Molecular Therapy - Methods and Clinical Development, 2017, 4, 39-49.	4.1	81
9	Characterization of Resistance Gene Analogues (RGAs) in Apple (Malus × domestica Borkh.) and Their Evolutionary History of the Rosaceae Family. PLoS ONE, 2014, 9, e83844.	2.5	71
10	A Pilot Characterization of the Human Chronobiome. Scientific Reports, 2017, 7, 17141.	3.3	70
11	Population structure of human gut bacteria in a diverse cohort from rural Tanzania and Botswana. Genome Biology, 2019, 20, 16.	8.8	66
12	INSPIIRED: Quantification and Visualization Tools for Analyzing Integration Site Distributions. Molecular Therapy - Methods and Clinical Development, 2017, 4, 17-26.	4.1	60
13	Associations of the vaginal microbiota with HIV infection, bacterial vaginosis, and demographic factors. Aids, 2017, 31, 895-904.	2.2	44
14	Extracellular vesicles from Kaposi Sarcoma-associated herpesvirus lymphoma induce long-term endothelial cell reprogramming. PLoS Pathogens, 2019, 15, e1007536.	4.7	40
15	High-Density Amplicon Sequencing Identifies Community Spread and Ongoing Evolution of SARS-CoV-2 in the Southern United States. Cell Reports, 2020, 33, 108352.	6.4	38
16	Allometry and Ecology of the Bilaterian Gut Microbiome. MBio, 2018, 9, .	4.1	29
17	Quantitation of HIV DNA integration: Effects of differential integration site distributions on Alu-PCR assays. Journal of Virological Methods, 2013, 189, 53-57.	2.1	21
18	Dietary Regulation of the Gut Microbiota Engineered by a Minimal Defined Bacterial Consortium. PLoS ONE. 2016. 11. e0155620.	2.5	16

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AUBREY G BAILEY

#	Article	IF	CITATIONS
19	Bringing it all together. Aids, 2013, 27, 835-838.	2.2	14
20	Novel modulators of p53-signaling encoded by unknown genes of emerging viruses. PLoS Pathogens, 2021, 17, e1009033.	4.7	12
21	Burden of respiratory viral infection in persons with human immunodeficiency virus. Influenza and Other Respiratory Viruses, 2020, 14, 465-469.	3.4	9
22	Increased Microbial Diversity Found in Successful Versus Unsuccessful Recipients of a Next-Generation FMT for Recurrent Clostridium difficile Infection. Open Forum Infectious Diseases, 2015, 2, .	0.9	4
23	The Subgingival Oral Microbiome in Pediatric Patients With Crohn's Disease. Inflammatory Bowel Diseases, 2012, 18, S61.	1.9	1
24	Sa1803 Longitudinal Characterization of the Subgingival Microbiota in Pediatric Crohn's Disease. Gastroenterology, 2013, 144, S-309-S-310.	1.3	0
25	Su2068 16S Gene Sequencing Reveals a Unique Bacterial Signature in the Gut Microbiota of Pediatric Patients With Cystic Fibrosis and Pancreatic Insufficiency. Gastroenterology, 2015, 148, S-589.	1.3	0
26	Breastfeeding Influences Establishment and Development of the Infant Microbiome. Open Forum Infectious Diseases, 2015, 2, .	0.9	0