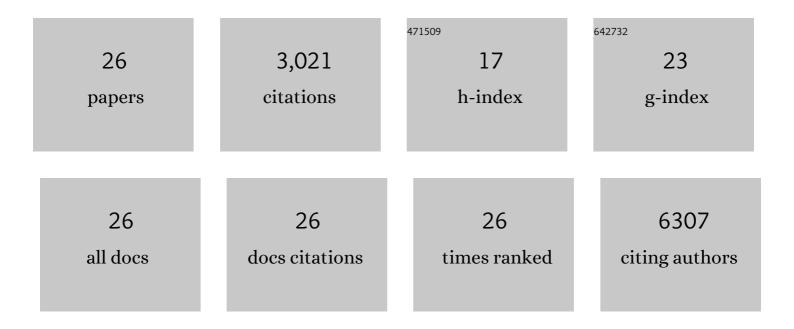
Aubrey G Bailey

List of Publications by Year in descending order

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ALIRDEV C. RALLEV

#	Article	IF	CITATIONS
1	Novel modulators of p53-signaling encoded by unknown genes of emerging viruses. PLoS Pathogens, 2021, 17, e1009033.	4.7	12
2	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
3	Burden of respiratory viral infection in persons with human immunodeficiency virus. Influenza and Other Respiratory Viruses, 2020, 14, 465-469.	3.4	9
4	Population structure of human gut bacteria in a diverse cohort from rural Tanzania and Botswana. Genome Biology, 2019, 20, 16.	8.8	66
5	Extracellular vesicles from Kaposi Sarcoma-associated herpesvirus lymphoma induce long-term endothelial cell reprogramming. PLoS Pathogens, 2019, 15, e1007536.	4.7	40
6	Allometry and Ecology of the Bilaterian Gut Microbiome. MBio, 2018, 9, .	4.1	29
7	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
8	INSPIIRED: Quantification and Visualization Tools for Analyzing Integration Site Distributions. Molecular Therapy - Methods and Clinical Development, 2017, 4, 17-26.	4.1	60
9	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
10	Association Between Breast Milk Bacterial Communities and Establishment and Development of the Infant Gut Microbiome. JAMA Pediatrics, 2017, 171, 647.	6.2	749
11	Associations of the vaginal microbiota with HIV infection, bacterial vaginosis, and demographic factors. Aids, 2017, 31, 895-904.	2.2	44
12	A Pilot Characterization of the Human Chronobiome. Scientific Reports, 2017, 7, 17141.	3.3	70
13	A role for bacterial urease in gut dysbiosis and Crohn's disease. Science Translational Medicine, 2017, 9, .	12.4	171
14	Peptidoglycan from the gut microbiota governs the lifespan of circulating phagocytes at homeostasis. Blood, 2016, 127, 2460-2471.	1.4	88
15	Comparison of placenta samples with contamination controls does not provide evidence for a distinct placenta microbiota. Microbiome, 2016, 4, 29.	11.1	447
16	Dietary Regulation of the Gut Microbiota Engineered by a Minimal Defined Bacterial Consortium. PLoS ONE, 2016, 11, e0155620.	2.5	16
17	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
18	Breastfeeding Influences Establishment and Development of the Infant Microbiome. Open Forum Infectious Diseases, 2015, 2, .	0.9	0

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#	Article	lF	CITATIONS
19	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
20	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
21	Virus-helminth coinfection reveals a microbiota-independent mechanism of immunomodulation. Science, 2014, 345, 578-582.	12.6	238
22	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
23	Sa1803 Longitudinal Characterization of the Subgingival Microbiota in Pediatric Crohn's Disease. Gastroenterology, 2013, 144, S-309-S-310.	1.3	0
24	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
25	Bringing it all together. Aids, 2013, 27, 835-838.	2.2	14
26	The Subgingival Oral Microbiome in Pediatric Patients With Crohn's Disease. Inflammatory Bowel Diseases, 2012, 18, S61.	1.9	1