

# Aubrey G Bailey

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2735812/publications.pdf>

Version: 2024-02-01

26  
papers

3,021  
citations

471509

17  
h-index

642732

23  
g-index

26  
all docs

26  
docs citations

26  
times ranked

6307  
citing authors

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

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