

# David T Pride

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1514770/publications.pdf>

Version: 2024-02-01

49  
papers

3,796  
citations

159585  
30  
h-index

214800  
47  
g-index

49  
all docs

49  
docs citations

49  
times ranked

4425  
citing authors

#	ARTICLE	IF	CITATIONS
1	Evidence of a robust resident bacteriophage population revealed through analysis of the human salivary virome. ISME Journal, 2012, 6, 915-926.	9.8	295
2	Evolutionary Implications of Microbial Genome Tetranucleotide Frequency Biases. Genome Research, 2003, 13, 145-158.	5.5	232
3	Electronic cigarette inhalation alters innate immunity and airway cytokines while increasing the virulence of colonizing bacteria. Journal of Molecular Medicine, 2016, 94, 667-679.	3.9	204
4	Library preparation methodology can influence genomic and functional predictions in human microbiome research. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 14024-14029.	7.1	179
5	The human urine virome in association with urinary tract infections. Frontiers in Microbiology, 2015, 6, 14.	3.5	173
6	Altered Oral Viral Ecology in Association with Periodontal Disease. MBio, 2014, 5, e01133-14.	4.1	171
7	Human oral viruses are personal, persistent and gender-consistent. ISME Journal, 2014, 8, 1753-1767.	9.8	159
8	Considerations for the Use of Phage Therapy in Clinical Practice. Antimicrobial Agents and Chemotherapy, 2022, 66, AAC0207121.	3.2	151
9	Microbial diversity in individuals and their household contacts following typical antibiotic courses. Microbiome, 2016, 4, 39.	11.1	135
10	Molecular Analysis of Sarcoidosis Tissues for <i>Mycobacterium</i> Species DNA. Emerging Infectious Diseases, 2002, 8, 1334-1341.	4.3	128
11	Metagenomic detection of phage-encoded platelet-binding factors in the human oral cavity. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 4547-4553.	7.1	123
12	Effects of Long Term Antibiotic Therapy on Human Oral and Fecal Viromes. PLoS ONE, 2015, 10, e0134941.	2.5	119
13	East Asian genotypes of <i>Helicobacter pylori</i> strains in Amerindians provide evidence for its ancient human carriage. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 15107-15111.	7.1	115
14	Molecular Bases and Role of Viruses in the Human Microbiome. Journal of Molecular Biology, 2014, 426, 3892-3906.	4.2	113
15	Evidence of host-virus co-evolution in tetranucleotide usage patterns of bacteriophages and eukaryotic viruses. BMC Genomics, 2006, 7, 8.	2.8	112
16	Bacteriophage and their potential roles in the human oral cavity. Journal of Oral Microbiology, 2015, 7, 27423.	2.7	109
17	Association between living environment and human oral viral ecology. ISME Journal, 2013, 7, 1710-1724.	9.8	108
18	Allelic Variation within <i>Helicobacter pylori</i> babA and babB. Infection and Immunity, 2001, 69, 1160-1171.	2.2	105

#	ARTICLE	IF	CITATIONS
19	Analysis of streptococcal CRISPRs from human saliva reveals substantial sequence diversity within and between subjects over time. <i>Genome Research</i> , 2011, 21, 126-136.	5.5	104
20	Shared and Distinct Features of Human Milk and Infant Stool Viromes. <i>Frontiers in Microbiology</i> , 2018, 9, 1162.	3.5	100
21	Characterization of bacteriophage communities and CRISPR profiles from dental plaque. <i>BMC Microbiology</i> , 2014, 14, 175.	3.3	83
22	Concerted evolution between duplicated genetic elements in <i>Helicobacter pylori</i> . <i>Journal of Molecular Biology</i> , 2002, 316, 629-642.	4.2	78
23	Transmission of viruses via our microbiomes. <i>Microbiome</i> , 2016, 4, 64.	11.1	70
24	Comparisons of clustered regularly interspaced short palindromic repeats and viromes in human saliva reveal bacterial adaptations to salivary viruses. <i>Environmental Microbiology</i> , 2012, 14, 2564-2576.	3.8	57
25	Chemostat culture systems support diverse bacteriophage communities from human feces. <i>Microbiome</i> , 2015, 3, 58.	11.1	50
26	Transcriptome analysis of bacteriophage communities in periodontal health and disease. <i>BMC Genomics</i> , 2015, 16, 549.	2.8	48
27	The Virome of Cerebrospinal Fluid: Viruses Where We Once Thought There Were None. <i>Frontiers in Microbiology</i> , 2019, 10, 2061.	3.5	48
28	Conservation of streptococcal CRISPRs on human skin and saliva. <i>BMC Microbiology</i> , 2014, 14, 146.	3.3	44
29	Immunogenicity and protective efficacy of recombinant <i>Clostridium difficile</i> flagellar protein FliC. <i>Emerging Microbes and Infections</i> , 2016, 5, 1-10.	6.5	44
30	Superoxide dismutase A antigens derived from molecular analysis of sarcoidosis granulomas elicit systemic Th-1 immune responses. <i>Respiratory Research</i> , 2008, 9, 36.	3.6	40
31	Benchmarking urine storage and collection conditions for evaluating the female urinary microbiome. <i>Scientific Reports</i> , 2019, 9, 13409.	3.3	33
32	Fecal Viral Community Responses to High-Fat Diet in Mice. <i>MSphere</i> , 2020, 5, .	2.9	33
33	Cervicovaginal Microbiome Composition Is Associated with Metabolic Profiles in Healthy Pregnancy. <i>MBio</i> , 2020, 11, .	4.1	30
34	Genome signature analysis of thermal virus metagenomes reveals Archaea and thermophilic signatures. <i>BMC Genomics</i> , 2008, 9, 420.	2.8	28
35	Comparison of Multiplex Gastrointestinal Pathogen Panel and Conventional Stool Testing for Evaluation of Diarrhea in Patients with Inflammatory Bowel Diseases. <i>Digestive Diseases and Sciences</i> , 2019, 64, 382-390.	2.3	22
36	Comparison of Three Nucleic Acid Amplification Tests and Culture for Detection of Group B <i>Streptococcus</i> from Enrichment Broth. <i>Journal of Clinical Microbiology</i> , 2019, 57, .	3.9	21

#	ARTICLE	IF	CITATIONS
37	Identification of Horizontally Acquired Genetic Elements in <i>Helicobacter pylori</i> and Other Prokaryotes Using Oligonucleotide Difference Analysis. <i>Journal of Genome Science and Technology</i> , 2002, 1, 2-15.	0.5	19
38	Compositional Differences in the Oral Microbiome of E-cigarette Users. <i>Frontiers in Microbiology</i> , 2021, 12, 599664.	3.5	18
39	A Rapid and Low-Cost Pathogen Detection Platform by Using a Molecular Agglutination Assay. <i>ACS Central Science</i> , 2018, 4, 1485-1494.	11.3	15
40	Global transcription of CRISPR loci in the human oral cavity. <i>BMC Genomics</i> , 2015, 16, 401.	2.8	14
41	Temporal variations in bacterial community diversity and composition throughout intensive care unit renovations. <i>Microbiome</i> , 2020, 8, 86.	11.1	14
42	Animal Models of Phage Therapy. <i>Frontiers in Microbiology</i> , 2021, 12, 631794.	3.5	13
43	Aseptic Barriers Allow a Clean Contact for Contaminated Stethoscope Diaphragms. <i>Mayo Clinic Proceedings Innovations, Quality &amp; Outcomes</i> , 2020, 4, 21-30.	2.4	12
44	Comparison of two nucleic acid amplification tests (NAATs) and two antigen tests for detection of SARS-CoV-2 from upper respiratory specimens. <i>Journal of Clinical Virology Plus</i> , 2021, 1, 100011.	1.0	9
45	Phage Cocktails Constrain the Growth of <i>Enterococcus</i> . <i>MSystems</i> , 2022, 7, .	3.8	9
46	Identification of staphylococcal phage with reduced transcription in human blood through transcriptome sequencing. <i>Frontiers in Microbiology</i> , 2015, 6, 216.	3.5	4
47	Increased Innate Immune Susceptibility in Hyperpigmented Bacteriophage-Resistant Mutants of <i>Pseudomonas aeruginosa</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 0, , .	3.2	3
48	Draft Genome Sequence of an <i>Enterococcus faecalis</i> ATCC 19433 Siphovirus Isolated from Raw Domestic Sewage. <i>Genome Announcements</i> , 2017, 5, .	0.8	2
49	Molecular Mirror Technology Facilitates High-Throughput, Accurate SARS-CoV-2 Testing. <i>Microbiology Spectrum</i> , 2021, 9, e0039221.	3.0	0