Jason J Gill

List of Publications by Year in descending order

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Version: 2024-02-01

97	3,097	24	53
papers	citations	h-index	g-index
103	103	103	3526
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Development and Use of Personalized Bacteriophage-Based Therapeutic Cocktails To Treat a Patient with a Disseminated Resistant Acinetobacter baumannii Infection. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	795
2	Phage Choice, Isolation, and Preparation for Phage Therapy. Current Pharmaceutical Biotechnology, 2010, 11, 2-14.	1.6	334
3	Efficacy and Pharmacokinetics of Bacteriophage Therapy in Treatment of Subclinical Staphylococcus aureus Mastitis in Lactating Dairy Cattle. Antimicrobial Agents and Chemotherapy, 2006, 50, 2912-2918.	3.2	145
4	Efficacy of Bacteriophage Therapy in a Model of <i>Burkholderia cenocepacia </i> Pulmonary Infection. Journal of Infectious Diseases, 2010, 201, 264-271.	4.0	134
5	Effects of zinc bacitracin, bird age and access to range on bacterial microbiota in the ileum and caeca of broiler chickens. Journal of Applied Microbiology, 2008, 104, 1372-1382.	3.1	120
6	Phage therapy redux—What is to be done?. Science, 2015, 350, 1163-1164.	12.6	113
7	Bacteriophages of Erwinia amylovora. Applied and Environmental Microbiology, 2003, 69, 2133-2138.	3.1	111
8	Galaxy and Apollo as a biologist-friendly interface for high-quality cooperative phage genome annotation. PLoS Computational Biology, 2020, 16, e1008214.	3.2	96
9	Bovine whey proteins inhibit the interaction of Staphylococcus aureus and bacteriophage K. Journal of Applied Microbiology, 2006, 101, 377-386.	3.1	94
10	Development and Validation of a Microtiter Plate-Based Assay for Determination of Bacteriophage Host Range and Virulence. Viruses, 2018, 10, 189.	3.3	90
11	The Caulobacter crescentus phage phiCbK: genomics of a canonical phage. BMC Genomics, 2012, 13, 542.	2.8	85
12	The habits of highly effective phages: population dynamics as a framework for identifying therapeutic phages. Frontiers in Microbiology, 2014, 5, 618.	3.5	66
13	Genomic and Biochemical Characterization of Acinetobacter Podophage Petty Reveals a Novel Lysis Mechanism and Tail-Associated Depolymerase Activity. Journal of Virology, 2018, 92, .	3.4	65
14	Genomic and Functional Analyses of <i>Rhodococcus equi</i> Phages ReqiPepy6, ReqiPoco6, ReqiPine5, and ReqiDocB7. Applied and Environmental Microbiology, 2011, 77, 669-683.	3.1	54
15	Molecular Typing and Distribution of Staphylococcus aureus Isolates in Eastern Canadian Dairy Herds. Journal of Clinical Microbiology, 2004, 42, 3449-3455.	3.9	53
16	Genomes and Characterization of Phages Bcep22 and BceplL02, Founders of a Novel Phage Type in Burkholderia cenocepacia. Journal of Bacteriology, 2011, 193, 5300-5313.	2.2	52
17	From farm management to bacteriophage therapy: strategies to reduce antibiotic use in animal agriculture. Annals of the New York Academy of Sciences, 2019, 1441, 31-39.	3.8	52
18	Role of phages in the pathogenesis of Burkholderia, or †Where are the toxin genes in Burkholderia phages?'. Current Opinion in Microbiology, 2007, 10, 410-417.	5.1	50

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19	Cultivation of lipid-producing bacteria with lignocellulosic biomass: Effects of inhibitory compounds of lignocellulosic hydrolysates. Bioresource Technology, 2014, 161, 162-170.	9.6	50
20	Bacteriophages of wastewater foaming-associated filamentous Gordonia reduce host levels in raw activated sludge. Scientific Reports, 2015, 5, 13754.	3.3	49
21	Characterization of bacterial populations recovered from the teat canals of lactating dairy and beef cattle by 16S rRNA gene sequence analysis. FEMS Microbiology Ecology, 2006, 56, 471-481.	2.7	44
22	Genomic and Biological Analysis of Phage Xfas53 and Related Prophages of <i>Xylella fastidiosa</i> Journal of Bacteriology, 2010, 192, 179-190.	2.2	39
23	The multicomponent antirestriction system of phage P1 is linked to capsid morphogenesis. Molecular Microbiology, 2017, 105, 399-412.	2.5	33
24	Phage-based extraction of polyhydroxybutyrate (PHB) produced from synthetic crude glycerol. Science of the Total Environment, 2016, 557-558, 317-321.	8.0	25
25	Bacteriophage application restores ethanol fermentation characteristics disrupted by Lactobacillus fermentum. Biotechnology for Biofuels, 2015, 8, 132.	6.2	24
26	Comparative genomics of Acinetobacter baumannii and therapeutic bacteriophages from a patient undergoing phage therapy. Nature Communications, 2022, 13, .	12.8	20
27	Evaluation of Commercial Prototype Bacteriophage Intervention Designed for Reducing O157 and Non-O157 Shiga-Toxigenic Escherichia coli (STEC) on Beef Cattle Hide. Foods, 2018, 7, 114.	4.3	19
28	Characterization of a Novel Tectivirus Phage Toil and Its Potential as an Agent for Biolipid Extraction. Scientific Reports, 2018, 8, 1062.	3.3	18
29	Revised Genome Sequence of Staphylococcus aureus Bacteriophage K. Genome Announcements, 2014, 2,	0.8	17
30	Prevalence and Characterization of Salmonella enterica and Salmonella Bacteriophages Recovered from Beef Cattle Feedlots in South Texas. Journal of Food Protection, 2016, 79, 1332-1340.	1.7	17
31	Practical and Theoretical Considerations for the Use of Bacteriophages in Food Systems. , 0, , 217-235.		15
32	Genome-wide screens reveal Escherichia coli genes required for growth of T1-like phage LL5 and V5-like phage LL12. Scientific Reports, 2020, 10, 8058.	3.3	15
33	Bacteriophages and phage-derived products as antibacterial therapeutics. Expert Opinion on Therapeutic Patents, 2007, 17, 1341-1350.	5.0	14
34	Comparative Genomics of Three Novel Jumbo Bacteriophages Infecting Staphylococcus aureus. Journal of Virology, 2021, 95, e0239120.	3.4	13
35	Sheep in wolves' clothing: Temperate T7-like bacteriophages and the origins of the Autographiviridae. Virology, 2022, 568, 86-100.	2.4	12
36	Modeling of bacteriophage therapy. , 2008, , 439-464.		10

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37	DNA Packaging and Genomics of the <i>>Salmonella</i> >9NA-Like Phages. Journal of Virology, 2019, 93, .	3.4	8
38	Comparison of 2 fixatives in the porcine colon for in situ microbiota studies. Journal of Animal Science, 2019, 97, 4803-4809.	0.5	8
39	Effect of chronic and acute enterotoxigenic E. coli challenge on growth performance, intestinal inflammation, microbiome, and metabolome of weaned piglets. Scientific Reports, 2022, 12, 5024.	3.3	8
40	Complete Genome Sequence of Stenotrophomonas Phage Pokken. Microbiology Resource Announcements, 2019, 8, .	0.6	7
41	Crowdsourcing biocuration: The Community Assessment of Community Annotation with Ontologies (CACAO). PLoS Computational Biology, 2021, 17, e1009463.	3.2	7
42	New Insights into the Structure and Assembly of Bacteriophage P1. Viruses, 2022, 14, 678.	3.3	6
43	Complete Genome Sequence of Salmonella enterica Serovar Typhimurium Siphophage Skate. Microbiology Resource Announcements, 2019, 8, .	0.6	5
44	Complete Genome Sequence of Klebsiella pneumoniae Phage Sweeny. Microbiology Resource Announcements, 2019, 8, .	0.6	5
45	Complete Genome Sequence of Klebsiella pneumoniae Siphophage Seifer. Microbiology Resource Announcements, 2019, 8, .	0.6	5
46	Complete Genome Sequence of Klebsiella pneumoniae Siphophage Sugarland. Microbiology Resource Announcements, 2018, 7, .	0.6	4
47	Complete Genome Sequence of Enterotoxigenic Escherichia coli Siphophage LL5. Microbiology Resource Announcements, 2019, 8, .	0.6	4
48	Complete Genome Sequence of Klebsiella pneumoniae Jumbo Phage Miami. Microbiology Resource Announcements, 2021, 10, .	0.6	4
49	Differential Bacteriophage Efficacy in Controlling Salmonella in Cattle Hide and Soil Models. Frontiers in Microbiology, 2021, 12, 657524.	3.5	4
50	Complete Genome Sequence of Escherichia coli Myophage Mangalitsa. Microbiology Resource Announcements, 2019, 8, .	0.6	4
51	Complete Genome Sequence of Escherichia coli Phage Paul. Microbiology Resource Announcements, 2019, 8, .	0.6	4
52	Complete Genome Sequence of Salmonella enterica Serovar Typhimurium Siphophage Seabear. Microbiology Resource Announcements, 2019, 8, .	0.6	4
53	Complete Genome Sequence of Salmonella enterica Serovar Enteritidis Siphophage Seafire. Microbiology Resource Announcements, 2019, 8, .	0.6	4
54	Complete Genome Sequence of the Novel Klebsiella pneumoniae Phage Marfa. Microbiology Resource Announcements, 2019, 8, .	0.6	3

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55	Complete Genome Sequence of Vibrio natriegens Phage Phriendly. Microbiology Resource Announcements, $2019, 8, \dots$	0.6	3
56	Complete Genome Sequence of Proteus mirabilis Phage Myduc. Microbiology Resource Announcements, 2019, 8, .	0.6	3
57	Complete Genome Sequence of Klebsiella pneumoniae Myophage Muenster. Microbiology Resource Announcements, 2021, 10, .	0.6	3
58	Complete Whole Genome Sequences of Escherichia coli Surrogate Strains and Comparison of Sequence Methods with Application to the Food Industry. Microorganisms, 2021, 9, 608.	3.6	3
59	Complete Genome Sequence of Klebsiella pneumoniae Podophage Pylas. Microbiology Resource Announcements, 2019, 8, .	0.6	3
60	Complete Genome Sequence of Klebsiella pneumoniae Myophage Mulock. Microbiology Resource Announcements, 2019, 8, .	0.6	3
61	Complete Genome Sequence of Klebsiella pneumoniae Myophage Magnus. Microbiology Resource Announcements, 2019, 8, .	0.6	3
62	Complete Genome Sequence of Staphylococcus aureus Siphophage Sebago. Microbiology Resource Announcements, 2019, 8, .	0.6	2
63	Complete Genome Sequence of Enterotoxigenic Escherichia coli Myophage LL12. Microbiology Resource Announcements, 2019, 8, .	0.6	2
64	Complete Genome Sequence of Salmonella enterica Serovar Typhimurium Myophage Mutine. Microbiology Resource Announcements, 2019, 8, .	0.6	2
65	Complete Genome Sequence of Salmonella enterica Serovar Typhimurium Siphophage Siskin. Microbiology Resource Announcements, 2019, 8, .	0.6	2
66	Complete Genome Sequence of Salmonella enterica Serovar Enteritidis Myophage Mooltan. Microbiology Resource Announcements, 2019, 8, .	0.6	2
67	Complete Genome Sequence of Shelby, a Siphophage Infecting Carbapenemase-Producing Klebsiella pneumoniae. Microbiology Resource Announcements, 2019, 8, .	0.6	2
68	Dual-function oleaginous biocatalysts for non-sterile cultivation and solvent-free biolipid bioextraction to reduce biolipid-based biofuel production costs. Science of the Total Environment, 2021, 758, 143969.	8.0	2
69	Isolation and characterization of novel phage (Podoviridae É.ParuNE1) and its efficacy against multi-drug-resistant Pseudomonas aeruginosa planktonic cells and biofilm. Beni-Suef University Journal of Basic and Applied Sciences, 2021, 10, .	2.0	2
70	Complete Genome Sequence of Staphylococcus aureus Myophage Maine. Microbiology Resource Announcements, 2019, 8, .	0.6	2
71	Complete Genome Sequence of Klebsiella pneumoniae Siphophage Skenny. Microbiology Resource Announcements, 2019, 8, .	0.6	2
72	Complete Genome Sequence of Xanthomonas Siphophage Samson. Microbiology Resource Announcements, 2019, 8, .	0.6	2

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73	Complete Genome Sequence of Klebsiella pneumoniae Siphophage Sanco. Microbiology Resource Announcements, 2019, 8 , .	0.6	2
74	Complete Genome Sequence of Citrobacter freundii Myophage Maleficent. Microbiology Resource Announcements, $2019, 8, \ldots$	0.6	2
75	Complete Genome Sequence of Citrobacter freundii Myophage Mijalis. Genome Announcements, 2017, 5,	0.8	1
76	Complete Genome Sequence of Enterotoxigenic Escherichia coli Podophage LL11. Microbiology Resource Announcements, 2019, 8, .	0.6	1
77	Complete Genome Sequence of Salmonella enterica Serovar Heidelberg Myophage Meda. Microbiology Resource Announcements, 2019, 8, .	0.6	1
78	Complete Genome Sequence of Burkholderia cenocepacia Phage Mica. Microbiology Resource Announcements, 2021, 10, .	0.6	1
79	Complete Genome Sequence of Burkholderia gladioli Myophage Mana. Microbiology Resource Announcements, 2021, 10, .	0.6	1
80	The Selection and Optimization of Phage Hosts. , 2021, , 689-698.		1
81	Complete Genome Sequence of Sin4, a Siphophage Infecting Carbapenemase-Producing Klebsiella pneumoniae. Microbiology Resource Announcements, 2019, 8, .	0.6	1
82	Complete Genome Sequence of Salmonella enterica Siphophage Shelanagig. Microbiology Resource Announcements, $2019, 8, \ldots$	0.6	1
83	Complete Genome Sequence of Salmonella enterica Myophage Matapan. Microbiology Resource Announcements, 2019, 8, .	0.6	1
84	Complete Genome Sequence of Proteus mirabilis Siphophage Saba. Microbiology Resource Announcements, 2019, 8, .	0.6	1
85	Complete Genome Sequence of Burkholderia cenocepacia Phage Paku. Microbiology Resource Announcements, 2022, , e0122021.	0.6	1
86	Complete Genome Sequence of Salmonella enterica Serovar Heidelberg Siphophage Sepoy. Microbiology Resource Announcements, 2019, 8, .	0.6	0
87	Complete Genome Sequence of Salmonella enterica Serovar Newport Myophage Melville. Microbiology Resource Announcements, 2019, 8, .	0.6	0
88	Complete Genome Sequence of Proteus mirabilis Phage Mydo. Microbiology Resource Announcements, 2019, 8, .	0.6	0
89	Complete Genome Sequence of Citrobacter freundii Siphophage Sazh. Microbiology Resource Announcements, 2019, 8, .	0.6	0
90	Complete Genome Sequence of Salmonella enterica Siphophage Shemara. Microbiology Resource Announcements, 2020, 9, .	0.6	0

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91	Complete Genome Sequence of Klebsiella aerogenes Siphophage Solomon. Microbiology Resource Announcements, 2021, 10, .	0.6	0
92	Complete Genome Sequence of Klebsiella aerogenes Myophage Metamorpho. Microbiology Resource Announcements, 2021, 10 , .	0.6	0
93	Complete Genome Sequence of Burkholderia cenocepacia Phage Magia. Microbiology Resource Announcements, 2021, 10, .	0.6	0
94	Complete Genome Sequence of Klebsiella pneumoniae Podophage Pone. Microbiology Resource Announcements, 2021, 10, e0140520.	0.6	0
95	Solvent Extraction of Klebsiella pneumoniae Bacteriophage Lysates with 1-Dodecanol Results in Endotoxin Reduction with Low Risk of Solvent Contamination. Phage, 2021, 2, 112-119.	1.7	0
96	Coding-Complete Genome Sequence of Staphylococcus aureus Podophage Portland. Microbiology Resource Announcements, 2019, 8, .	0.6	0
97	The Selection and Optimization of Phage Hosts. , 2020, , 1-10.		O