

# Jolene Ramsey

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

35  
papers

175  
citations

5  
h-index

12  
g-index

52  
ext. papers

276  
ext. citations

2.2  
avg, IF

3.3  
L-index

#	Paper	IF	Citations
35	Complete Genome Sequences of Lambdoid Phages 21, 434, and 434B and Several Lambda Hybrids.. <i>Microbiology Resource Announcements</i> , <b>2022</b> , e0012022	1.3	0
34	Endolysin Regulation in Phage Mu Lysis.. <i>MBio</i> , <b>2022</b> , e0081322	7.8	
33	Crowdsourcing biocuration: The Community Assessment of Community Annotation with Ontologies (CACAO). <i>PLoS Computational Biology</i> , <b>2021</b> , 17, e1009463	5	3
32	Characterization and complete genome sequence of Privateer, a highly prolate podophage. <i>PeerJ</i> , <b>2021</b> , 9, e10645	3.1	1
31	Phage-encoded cationic antimicrobial peptide required for lysis. <i>Journal of Bacteriology</i> , <b>2021</b> , JB002142	3.5	3
30	Complete Genome Sequence of Serratia marcescens Podophage Pila. <i>Microbiology Resource Announcements</i> , <b>2020</b> , 9,	1.3	1
29	Complete Genome Sequence of Stenotrophomonas Phage Mendera. <i>Microbiology Resource Announcements</i> , <b>2020</b> , 9,	1.3	3
28	Complete Genome Sequence of Stenotrophomonas maltophilia Myophage Moby. <i>Microbiology Resource Announcements</i> , <b>2020</b> , 9,	1.3	2
27	Complete Genome Sequence of Phage Muldoon. <i>Microbiology Resource Announcements</i> , <b>2020</b> , 9,	1.3	1
26	Galaxy and Apollo as a biologist-friendly interface for high-quality cooperative phage genome annotation. <i>PLoS Computational Biology</i> , <b>2020</b> , 16, e1008214	5	44
25	Complete Genome Sequence of Serratia marcescens Siphophage Slocum. <i>Microbiology Resource Announcements</i> , <b>2020</b> , 9,	1.3	1
24	Complete Genome Sequence of Serratia marcescens Phage MTx. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
23	Domains of the TF protein important in regulating its own palmitoylation. <i>Virology</i> , <b>2019</b> , 531, 31-39	3.6	3
22	Complete Genome Sequence of Agrobacterium tumefaciens Myophage Milano. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
21	Complete Genome Sequence of Serratia marcescens Myophage Moabite. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	3
20	Complete Genome Sequence of the Novel Klebsiella pneumoniae Phage Marfa. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
19	Complete Genome Sequence of Serratia marcescens Podophage Parlo. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	3

18	Complete Genome Sequence of <i>Stenotrophomonas maltophilia</i> Podophage Ponderosa. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	5
17	Complete Genome Sequence of <i>Escherichia coli</i> Myophage Mangalitsa. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
16	Complete Genome Sequence of <i>Serratia marcescens</i> Myophage MyoSmar. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
15	Complete Genome Sequence of <i>Klebsiella pneumoniae</i> Phage Sweeny. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	5
14	Complete Genome Sequence of <i>Escherichia coli</i> Phage Pisces. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
13	Complete Genome Sequence of <i>Escherichia coli</i> Phage Paul. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
12	Complete Genome Sequence of <i>Stenotrophomonas</i> Phage Pokken. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	6
11	Complete Genome Sequence of <i>Klebsiella pneumoniae</i> Myophage Magnus. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
10	Complete Genome Sequence of Sin4, a Siphophage Infecting Carbapenemase-Producing <i>Klebsiella pneumoniae</i> . <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
9	Complete Genome Sequence of <i>Klebsiella pneumoniae</i> Siphophage Skenny. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
8	Complete Genome Sequence of Siphophage Samson. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
7	Complete Genome Sequence of Phage Pagan. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
6	Complete Genome Sequence of <i>Vibrio natriegens</i> Phage Phriendly. <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	1
5	Complete Genome Sequence of Shelby, a Siphophage Infecting Carbapenemase-Producing <i>Klebsiella pneumoniae</i> . <i>Microbiology Resource Announcements</i> , <b>2019</b> , 8,	1.3	2
4	Alphavirus Purification Using Low-speed Spin Centrifugation. <i>Bio-protocol</i> , <b>2018</b> , 8, e2772	0.9	1
3	Palmitoylation of Sindbis Virus TF Protein Regulates Its Plasma Membrane Localization and Subsequent Incorporation into Virions. <i>Journal of Virology</i> , <b>2017</b> , 91,	6.6	21
2	Disentangling the Frames, the State of Research on the Alphavirus 6K and TF Proteins. <i>Viruses</i> , <b>2017</b> , 9,	6.2	27
1	Comprehensive in vivo RNA-binding site analyses reveal a role of Prp8 in spliceosomal assembly. <i>Nucleic Acids Research</i> , <b>2013</b> , 41, 3805-18	20.1	21

