

Jumpei Uchiyama

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

669
citations

840119

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610482

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all docs

24
docs citations

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times ranked

1017
citing authors

#	ARTICLE	IF	CITATIONS
1	Heterogeneous IgE reactivities to <i>Staphylococcus pseudintermedius</i> strains in dogs with atopic dermatitis, and the identification of DM13-domain-containing protein as a bacterial IgE-reactive molecule. <i>FEMS Microbiology Letters</i> , 2022, 369, .	0.7	1
2	Purification of membrane vesicles from Gram-positive bacteria using flow cytometry, after iodixanol density-gradient ultracentrifugation. <i>Research in Microbiology</i> , 2021, 172, 103792.	1.0	1
3	Analyses of propagation processes of <i>Staphylococcus aureus</i> bacteriophages S13 and S25-3 in two different taxonomies by definitive screening design. <i>Virus Research</i> , 2021, 298, 198406.	1.1	3
4	Bacterial Viruses Subcommittee and Archaeal Viruses Subcommittee of the ICTV: update of taxonomy changes in 2021. <i>Archives of Virology</i> , 2021, 166, 3239-3244.	0.9	24
5	Use of Recombinant Endolysin to Improve Accuracy of Group B <i>Streptococcus</i> Tests. <i>Microbiology Spectrum</i> , 2021, 9, e0007721.	1.2	2
6	Specific antiviral effect of violaceoid E on bovine leukemia virus. <i>Virology</i> , 2021, 562, 1-8.	1.1	7
7	In Vitro and In Vivo Evaluation of Three Newly Isolated Bacteriophage Candidates, phiEF7H, phiEF14H1, phiEF19C, for Treatment of <i>Enterococcus faecalis</i> Endophthalmitis. <i>Microorganisms</i> , 2021, 9, 212.	1.6	6
8	Examination of the fecal microbiota in dairy cows infected with bovine leukemia virus. <i>Veterinary Microbiology</i> , 2020, 240, 108547.	0.8	27
9	Development of multipurpose recombinant reporter bovine leukemia virus. <i>Virology</i> , 2020, 548, 226-235.	1.1	3
10	Taxonomy of prokaryotic viruses: 2018-2019 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. <i>Archives of Virology</i> , 2020, 165, 1253-1260.	0.9	144
11	Novel neuroprotective hydroquinones with a vinyl alkyne from the fungus, <i>Pestalotiopsis microspora</i> . <i>Journal of Antibiotics</i> , 2019, 72, 793-799.	1.0	11
12	Therapeutic Potential of an Endolysin Derived from Kayvirus S25-3 for Staphylococcal Impetigo. <i>Viruses</i> , 2019, 11, 769.	1.5	25
13	Dark-Field Microscopic Detection of Bacteria using Bacteriophage-Immobilized SiO ₂ @AuNP Core-Shell Nanoparticles. <i>Analytical Chemistry</i> , 2019, 91, 12352-12357.	3.2	41
14	Taxonomy of prokaryotic viruses: 2017 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. <i>Archives of Virology</i> , 2018, 163, 1125-1129.	0.9	172
15	Piperacillin and ceftazidime produce the strongest synergistic phage-antibiotic effect in <i>Pseudomonas aeruginosa</i> . <i>Archives of Virology</i> , 2018, 163, 1941-1948.	0.9	58
16	Potential Application of Bacteriophages in Enrichment Culture for Improved Prenatal <i>Streptococcus agalactiae</i> Screening. <i>Viruses</i> , 2018, 10, 552.	1.5	7
17	Variations in the viral genome and biological properties of bovine leukemia virus wild-type strains. <i>Virus Research</i> , 2018, 253, 103-111.	1.1	21
18	Genome Sequences of 12 Mycobacteriophages Recovered from Archival Stocks in Japan. <i>Genome Announcements</i> , 2018, 6, .	0.8	4

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19	Virus purification by CsCl density gradient using general centrifugation. Archives of Virology, 2017, 162, 3523-3528.	0.9	45
20	Analyses of Short-Term Antagonistic Evolution of Pseudomonas aeruginosa Strain PAO1 and Phage KPP22 (Myoviridae Family, PB1-Like Virus Genus). Applied and Environmental Microbiology, 2016, 82, 4482-4491.	1.4	26
21	Characterization of <i>Pseudomonas aeruginosa</i> phage KPP21 belonging to family Podoviridae genus N4-like viruses isolated in Japan. Microbiology and Immunology, 2016, 60, 64-67.	0.7	12
22	In silico analysis of AHJD-like viruses, <i>Staphylococcus aureus</i> phages S24 and S13, and study of phage S24 adsorption. MicrobiologyOpen, 2014, 3, 257-270.	1.2	15
23	Characterization of a novel Pseudomonas aeruginosa bacteriophage, KPP25, of the family Podoviridae. Virus Research, 2014, 189, 43-46.	1.1	12
24	Analysis of deoxynucleosides in bacteriophages EF24C and K and the frequency of a specific restriction site in the genomes of members of the bacteriophage subfamily Spounavirinae. Archives of Virology, 2012, 157, 1587-1592.	0.9	2