Jakub Barylski

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Plant antimicrobial peptides. Folia Microbiologica, 2014, 59, 181-196.	2.3	305
2	Taxonomy of prokaryotic viruses: 2017 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. Archives of Virology, 2018, 163, 1125-1129.	2.1	172
3	Taxonomy of prokaryotic viruses: 2018-2019 update from the ICTV Bacterial and Archaeal Viruses Subcommittee. Archives of Virology, 2020, 165, 1253-1260.	2.1	144
4	Analysis of Spounaviruses as a Case Study for the Overdue Reclassification of Tailed Phages. Systematic Biology, 2020, 69, 110-123.	5.6	89
5	Taxonomy of prokaryotic viruses: update from the ICTV bacterial and archaeal viruses subcommittee. Archives of Virology, 2016, 161, 1095-1099.	2.1	83
6	Bacteriophages are more virulent to bacteria with human cells than they are in bacterial culture; insights from HT-29 cells. Scientific Reports, 2018, 8, 5091.	3.3	79
7	Viral and Other Cell-Penetrating Peptides as Vectors of Therapeutic Agents in Medicine. Journal of Pharmacology and Experimental Therapeutics, 2015, 354, 32-42.	2.5	62
8	Taxonomy of prokaryotic viruses: 2016 update from the ICTV bacterial and archaeal viruses subcommittee. Archives of Virology, 2017, 162, 1153-1157.	2.1	57
9	RaFAH: Host prediction for viruses of Bacteria and Archaea based on protein content. Patterns, 2021, 2, 100274.	5.9	53
10	Effect of temperature on the pathogenesis, accumulation of viral and satellite RNAs and on plant proteome in peanut stunt virus and satellite RNA-infected plants. Frontiers in Plant Science, 2015, 6, 903.	3.6	40
11	ICTV Virus Taxonomy Profile: Herelleviridae. Journal of General Virology, 2020, 101, 362-363.	2.9	37
12	Two Novel Myoviruses from the North of Iraq Reveal Insights into Clostridium difficile Phage Diversity and Biology. Viruses, 2016, 8, 310.	3.3	31
13	The Discovery of phiAGATE, A Novel Phage Infecting Bacillus pumilus, Leads to New Insights into the Phylogeny of the Subfamily Spounavirinae. PLoS ONE, 2014, 9, e86632.	2.5	29
14	Cylindrospermopsin Biodegradation Abilities of Aeromonas sp. Isolated from RusaÅ,ka Lake. Toxins, 2016, 8, 55.	3.4	24
15	Bacterial Viruses Subcommittee and Archaeal Viruses Subcommittee of the ICTV: update of taxonomy changes in 2021. Archives of Virology, 2021, 166, 3239-3244.	2.1	24
16	Combination of transcriptomic and proteomic approaches helps to unravel the protein composition of Chelidonium majus L. milky sap. Planta, 2016, 244, 1055-1064.	3.2	22
17	Uncoupling proteins of invertebrates: A review. IUBMB Life, 2016, 68, 691-699.	3.4	22
18	Correlation between specific groups of heterotrophic bacteria and microcystin biodegradation in freshwater bodies of central Europe. FEMS Microbiology Ecology, 2019, 95, .	2.7	18

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19	Taxonomy-aware, sequence similarity ranking reliably predicts phage–host relationships. BMC Biology, 2021, 19, 223.	3.8	16
20	Two novel temperate bacteriophages infecting Streptococcus pyogenes: Their genomes, morphology and stability. PLoS ONE, 2018, 13, e0205995.	2.5	15
21	Characterization of the complete chloroplast genome of Pinus uliginosa (Neumann) from the Pinus mugo complex. Conservation Genetics Resources, 2017, 9, 209-212.	0.8	10
22	Incorrectly annotated keratin derived peptide sequences lead to misleading MS/MS data interpretation. Journal of Proteomics, 2013, 91, 270-273.	2.4	7
23	Complete genome sequences of two novel autographiviruses infecting a bacterium from the Pseudomonas fluorescens group. Archives of Virology, 2017, 162, 2907-2911.	2.1	7
24	Complete Genome Sequence of <i>Lelliottia</i> Podophage phD2B. Genome Announcements, 2014, 2, .	0.8	3
25	Characterization and expression of a novel thaumatin-like protein (CcTLP1) from papaveraceous plant Corydalis cava. International Journal of Biological Macromolecules, 2021, 189, 678-689.	7.5	2
26	Molecular, morphological, and ecological differences between the terrestrial and aquatic forms ofOxyrrhynchium speciosum(Brid.) Warnst. (Brachytheciaceae). Journal of Bryology, 2014, 36, 180-190.	1.2	1