List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Sulfolobus tengchongensis Spindle-Shaped Virus STSV1: Virus-Host Interactions and Genomic Features. Journal of Virology, 2005, 79, 8677-8686.	3.4	119
2	An Abundant DNA Binding Protein from the Hyperthermophilic Archaeon Sulfolobus shibatae Affects DNA Supercoiling in a Temperature-Dependent Fashion. Journal of Bacteriology, 2000, 182, 3929-3933.	2.2	92
3	Ssh10b, a conserved thermophilic archaeal protein, binds RNA in vivo. Molecular Microbiology, 2003, 50, 1605-1615.	2.5	83
4	Biochemical and structural characterization of Cren7, a novel chromatin protein conserved among Crenarchaea. Nucleic Acids Research, 2008, 36, 1129-1137.	14.5	82
5	Diversity, Biogeography, and Biodegradation Potential of Actinobacteria in the Deep-Sea Sediments along the Southwest Indian Ridge. Frontiers in Microbiology, 2016, 7, 1340.	3.5	76
6	Archaeal Extrachromosomal Genetic Elements. Microbiology and Molecular Biology Reviews, 2015, 79, 117-152.	6.6	64
7	Expansion of <i>Thaumarchaeota</i> habitat range is correlated with horizontal transfer of ATPase operons. ISME Journal, 2019, 13, 3067-3079.	9.8	59
8	Revealing the essentiality of multiple archaeal pcna genes using a mutant propagation assay based on an improved knockout method. Microbiology (United Kingdom), 2010, 156, 3386-3397.	1.8	58
9	Structural insights into the interaction of the crenarchaeal chromatin protein Cren7 with DNA. Molecular Microbiology, 2010, 76, 749-759.	2.5	44
10	High-throughput single-cell cultivation reveals the underexplored rare biosphere in deep-sea sediments along the Southwest Indian Ridge. Lab on A Chip, 2020, 20, 363-372.	6.0	31
11	Biochemical and Structural Insights into RNA Binding by Ssh10b, a Member of the Highly Conserved Sac10b Protein Family in Archaea. Journal of Biological Chemistry, 2014, 289, 1478-1490.	3.4	30
12	Archaeal chromatin proteins. Science China Life Sciences, 2012, 55, 377-385.	4.9	29
13	One cell at a time: droplet-based microbial cultivation, screening and sequencing. Marine Life Science and Technology, 2021, 3, 169-188.	4.6	29
14	Extraordinary diversity of viruses in deepâ€sea sediments as revealed by metagenomics without prior virion separation. Environmental Microbiology, 2021, 23, 728-743.	3.8	27
15	Identification and Characterization of a Highly Conserved Crenarchaeal Protein Lysine Methyltransferase with Broad Substrate Specificity. Journal of Bacteriology, 2012, 194, 6917-6926.	2.2	26
16	Genome Sequencing of Sulfolobus sp. A20 from Costa Rica and Comparative Analyses of the Putative Pathways of Carbon, Nitrogen, and Sulfur Metabolism in Various Sulfolobus Strains. Frontiers in Microbiology, 2016, 7, 1902.	3.5	26
17	The Sac10b Homolog in <i>Methanococcus maripaludis</i> Binds DNA at Specific Sites. Journal of Bacteriology, 2009, 191, 2315-2329.	2.2	24
18	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

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19	Two Conformations of Archaeal Ssh10b. Journal of Biological Chemistry, 2003, 278, 51015-51022.	3.4	23
20	NQO-Induced DNA-Less Cell Formation Is Associated with Chromatin Protein Degradation and Dependent on A0A1-ATPase in Sulfolobus. Frontiers in Microbiology, 2017, 8, 1480.	3.5	23
21	Cultivation of uncultured marine microorganisms. Marine Life Science and Technology, 2021, 3, 117-120.	4.6	23
22	Crystal structure of a DNA binding protein from the hyperthermophilic euryarchaeonMethanococcus jannaschii. Protein Science, 2003, 12, 2815-2822.	7.6	22
23	A primase subunit essential for efficient primer synthesis by an archaeal eukaryotic-type primase. Nature Communications, 2015, 6, 7300.	12.8	18
24	Functional Insights Into Protein Acetylation in the Hyperthermophilic Archaeon Sulfolobus islandicus*. Molecular and Cellular Proteomics, 2019, 18, 1572-1587.	3.8	18
25	The Sulfolobus solfataricus GINS Complex Stimulates DNA Binding and Processive DNA Unwinding by Minichromosome Maintenance Helicase. Journal of Bacteriology, 2015, 197, 3409-3420.	2.2	17
26	Virgibacillus indicus sp. nov. and Virgibacillus profundi sp. nov, two moderately halophilic bacteria isolated from marine sediment by using microfluidic streak plates. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2015-2023.	1.7	17
27	aKMT Catalyzes Extensive Protein Lysine Methylation in the Hyperthermophilic Archaeon Sulfolobus islandicus but is Dispensable for the Growth of the Organism. Molecular and Cellular Proteomics, 2016, 15, 2908-2923.	3.8	16
28	Fulvimarina manganoxydans sp. nov., isolated from a deep-sea hydrothermal plume in the south-west Indian Ocean. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 2920-2925.	1.7	15
29	Novel Sulfolobus Virus with an Exceptional Capsid Architecture. Journal of Virology, 2018, 92, .	3.4	15
30	Insights into the postâ€translational modifications of archaeal Sis10b (Alba): lysineâ€16 is methylated, not acetylated, and this does not regulate transcription or growth. Molecular Microbiology, 2018, 109, 192-208.	2.5	14
31	Gimesia benthica sp. nov., a planctomycete isolated from a deep-sea water sample of the Northwest Indian Ocean. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4384-4389.	1.7	14
32	Structural and functional characterization of the C-terminal catalytic domain of SSV1 integrase. Acta Crystallographica Section D: Biological Crystallography, 2012, 68, 659-670.	2.5	13
33	A Stabilizing α/β-Hydrophobic Core Greatly Contributes to Hyperthermostability of Archaeal [P62A]Ssh10b. Biochemistry, 2008, 47, 11212-11221.	2.5	12
34	Site-Specific Recombination by SSV2 Integrase: Substrate Requirement and Domain Functions. Journal of Virology, 2015, 89, 10934-10944.	3.4	11
35	Architectural roles of Cren7 in folding crenarchaeal chromatin filament. Molecular Microbiology, 2019, 111, 556-569.	2.5	11
36	Archaeal Chromatin Proteins Cren7 and Sul7d Compact DNA by Bending and Bridging. MBio, 2020, 11, .	4.1	11

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37	Phaeocystidibacter marisrubri sp. nov., a member of the family Cryomorphaceae isolated from Red Sea sediment. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 2199-2203.	1.7	11
38	Hyphobacterium indicum sp. nov., isolated from deep seawater, and emended description of the genus Hyphobacterium. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3760-3765.	1.7	11
39	Interfacial Nanoinjectionâ€Based Nanoliter Singleâ€Cell Analysis. Small, 2020, 16, e1903739.	10.0	9
40	Novel <i>Sulfolobus</i> Fuselloviruses with Extensive Genomic Variations. Journal of Virology, 2020, 94, .	3.4	9
41	Detection of the deep biosphere in metamorphic rocks from the Chinese continental scientific drilling. Geobiology, 2021, 19, 278-291.	2.4	9
42	The extraordinary thermal stability of EstA from <i>S. islandicus</i> is independent of post translational modifications. Protein Science, 2017, 26, 1819-1827.	7.6	8
43	Insights into the interaction between Cren7 and DNA: the role of loop β3–β4. Extremophiles, 2015, 19, 395-406.	2.3	7
44	Halovulum marinum sp. nov., isolated from deep-sea water of the Indian Ocean, and emended description of the genus Halovulum. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4523-4530.	1.7	7
45	Sulfolobus Replication Factor C Stimulates the Activity of DNA Polymerase B1. Journal of Bacteriology, 2014, 196, 2367-2375.	2.2	6
46	Isolation and Characterization of the First Temperate Virus Infecting Psychrobacillus from Marine Sediments. Viruses, 2022, 14, 108.	3.3	6
47	Roles of Leu28 side chain intercalation in the interaction between Cren7 and DNA. Biochemical Journal, 2017, 474, 1727-1739.	3.7	5
48	The Sac10b homolog from Sulfolobus islandicus is an RNA chaperone. Nucleic Acids Research, 2020, 48, 9273-9284.	14.5	5
49	Lysine Methylation Modulates the Interaction of Archaeal Chromatin Protein Cren7 With DNA. Frontiers in Microbiology, 2022, 13, 837737.	3.5	5
50	The Archaeal Transcription Termination Factor aCPSF1 is a Robust Phylogenetic Marker for Archaeal Taxonomy. Microbiology Spectrum, 2021, 9, e0153921.	3.0	5
51	A euryarchaeal histone modulates strand displacement synthesis by replicative DNA polymerases. Science China Life Sciences, 2016, 59, 709-716.	4.9	4
52	Microbes: the hidden giant behind the biogeochemical cycling of elements in the hydrosphere. Science China Life Sciences, 2019, 62, 1271-1274.	4.9	4
53	A Novel Family of Winged-Helix Single-Stranded DNA-Binding Proteins from Archaea. International Journal of Molecular Sciences, 2022, 23, 3455.	4.1	4
54	pTC Plasmids from Sulfolobus Species in the Geothermal Area of Tengchong, China: Genomic Conservation and Naturally-Occurring Variations as a Result of Transposition by Mobile Genetic Elements. Life, 2015, 5, 506-520.	2.4	2

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55	Kinetic insights into the temperature dependence of DNA strand cleavage and religation by topoisomerase III from the hyperthermophile Sulfolobus solfataricus. Scientific Reports, 2017, 7, 5494.	3.3	2
56	Self-assembled bionanoparticles based on the Sulfolobus tengchongensis spindle-shaped virus 1 (STSV1) coat protein as a prospective bioscaffold for nanotechnological applications. Extremophiles, 2014, 18, 745-754.	2.3	1
57	ICTV Virus Taxonomy Profile: Ovaliviridae. Journal of General Virology, 2021, 102, .	2.9	1
58	Construction of a Fusellovirus with a Minimal Set of Genes. ACS Synthetic Biology, 2021, 10, 2617-2627.	3.8	1
59	Effect of DNA binding protein Ssh12 from hyperthermophilic archaeonSulfolobus shibatae on DNA supercoiling. Science in China Series C: Life Sciences, 1999, 42, 401-408.	1.3	Ο
60	<i>>mLife</i> : Your journal for cuttingâ€edge research in all microbiological disciplines. , 2022, 1, 1-2.		0
61	Reply to Zrelovs et al. PVJ1 Is Not the First Tailed Temperate Phage Infecting Bacteria from Genus Psychrobacillus. Comment on "Liu et al. Isolation and Characterization of the First Temperate Virus Infecting Psychrobacillus from Marine Sediments. Viruses 2022, 14, 108― Viruses, 2022, 14, 866.	3.3	0