

Tessa E F Quax

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

1,552
citations

471061

17
h-index

377514

34
g-index

42
all docs

42
docs citations

42
times ranked

1798
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural insights into the mechanism of archaeal rotational switching. <i>Nature Communications</i> , 2022, 13, .	5.8	1
2	The Viral Susceptibility of the <i>Haloferax</i> Species. <i>Viruses</i> , 2022, 14, 1344.	1.5	4
3	The biology of thermoacidophilic archaea from the order <i>Sulfolobales</i> . <i>FEMS Microbiology Reviews</i> , 2021, 45, .	3.9	24
4	Viral Hijack of Filamentous Surface Structures in Archaea and Bacteria. <i>Viruses</i> , 2021, 13, 164.	1.5	15
5	Cellular and Genomic Properties of <i>Haloferax gibbonsii</i> LR2-5, the Host of Euryarchaeal Virus HFTV1. <i>Frontiers in Microbiology</i> , 2021, 12, 625599.	1.5	9
6	Viruses of Microbes 2020: The Latest Conquest on Viruses of Microbes. <i>Viruses</i> , 2021, 13, 802.	1.5	0
7	Growth Phase Dependent Cell Shape of <i>Haloarcula</i> . <i>Microorganisms</i> , 2021, 9, 231.	1.6	7
8	Insights into synthesis and function of KsgA/Dim1-dependent rRNA modifications in archaea. <i>Nucleic Acids Research</i> , 2021, 49, 1662-1687.	6.5	20
9	Motile ghosts of the halophilic archaeon, <i>Haloferax volcanii</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 26766-26772.	3.3	6
10	Interaction of two strongly divergent archaeellins stabilizes the structure of the <i>Halorubrum</i> archaeellum. <i>MicrobiologyOpen</i> , 2020, 9, e1047.	1.2	10
11	An Oscillating MinD Protein Determines the Cellular Positioning of the Motility Machinery in Archaea. <i>Current Biology</i> , 2020, 30, 4956-4972.e4.	1.8	19
12	The switch complex ArlCDE connects the chemotaxis system and the archaeellum. <i>Molecular Microbiology</i> , 2020, 114, 468-479.	1.2	19
13	Positioning of the Motility Machinery in Halophilic Archaea. <i>MBio</i> , 2019, 10, .	1.8	42
14	Cyclic nucleotides in archaea: Cyclic diAMP in the archaeon <i>Haloferax volcanii</i> and its putative role. <i>MicrobiologyOpen</i> , 2019, 8, e00829.	1.2	32
15	Architecture and modular assembly of <i>Sulfolobus</i> S-layers revealed by electron cryotomography. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 25278-25286.	3.3	33
16	Salt-dependent regulation of archaeellins in <i>Haloarcula marismortui</i> . <i>MicrobiologyOpen</i> , 2019, 8, e00718.	1.2	16
17	Structure and function of the archaeal response regulator CheY. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E1259-E1268.	3.3	43
18	Versatile cell surface structures of archaea. <i>Molecular Microbiology</i> , 2018, 107, 298-311.	1.2	50

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19	Structure and assembly mechanism of virus-associated pyramids. <i>Biophysical Reviews</i> , 2018, 10, 551-557.	1.5	8
20	Taxis in archaea. <i>Emerging Topics in Life Sciences</i> , 2018, 2, 535-546.	1.1	19
21	DNA-Interacting Characteristics of the Archaeal Rudiviral Protein SIRV2_Gp1. <i>Viruses</i> , 2017, 9, 190.	1.5	10
22	Viruses of Microbes. <i>Viruses</i> , 2017, 9, 263.	1.5	5
23	Improving heterologous membrane protein production in <i>Escherichia coli</i> by combining transcriptional tuning and codon usage algorithms. <i>PLoS ONE</i> , 2017, 12, e0184355.	1.1	37
24	Archaeal Surface Structures and Their Role in Communication with the Extracellular Environment. , 2017, , 67-84.		0
25	Archaeal viruses at the cell envelope: entry and egress. <i>Frontiers in Microbiology</i> , 2015, 6, 552.	1.5	23
26	Codon Bias as a Means to Fine-Tune Gene Expression. <i>Molecular Cell</i> , 2015, 59, 149-161.	4.5	554
27	Self-assembly of the general membrane-remodeling protein PVAP into sevenfold virus-associated pyramids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 3829-3834.	3.3	45
28	Unique genome replication mechanism of the archaeal virus <scp>AFV</scp>1. <i>Molecular Microbiology</i> , 2014, 92, 1313-1325.	1.2	16
29	Differential Translation Tunes Uneven Production of Operon-Encoded Proteins. <i>Cell Reports</i> , 2013, 4, 938-944.	2.9	64
30	First Insights into the Entry Process of Hyperthermophilic Archaeal Viruses. <i>Journal of Virology</i> , 2013, 87, 13379-13385.	1.5	66
31	Massive Activation of Archaeal Defense Genes during Viral Infection. <i>Journal of Virology</i> , 2013, 87, 8419-8428.	1.5	84
32	Insights into a Viral Lytic Pathway from an Archaeal Virus-Host System. <i>Journal of Virology</i> , 2013, 87, 2186-2192.	1.5	20
33	Exceptional virion release mechanism: one more surprise from archaeal viruses. <i>Current Opinion in Microbiology</i> , 2011, 14, 315-320.	2.3	26
34	Simple and elegant design of a virion egress structure in Archaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 3354-3359.	3.3	49
35	The <i>Sulfolobus</i> rod-shaped virus 2 encodes a prominent structural component of the unique virion release system in Archaea. <i>Virology</i> , 2010, 404, 1-4.	1.1	44
36	A unique virus release mechanism in the Archaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 11306-11311.	3.3	126