List of Publications by Year in descending order

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XIHIII SHEN

#	Article	IF	CITATIONS
1	A Pseudomonas T6SS effector recruits PQS-containing outer membrane vesicles for iron acquisition. Nature Communications, 2017, 8, 14888.	12.8	236
2	Legionella pneumophila inhibits macrophage apoptosis by targeting pro-death members of the Bcl2 protein family. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 5121-5126.	7.1	198
3	Inhibition of Host Vacuolar H+-ATPase Activity by a Legionella pneumophila Effector. PLoS Pathogens, 2010, 6, e1000822.	4.7	197
4	Secreted Bacterial Effectors That Inhibit Host Protein Synthesis Are Critical for Induction of the Innate Immune Response to Virulent Legionella pneumophila. PLoS Pathogens, 2011, 7, e1001289.	4.7	187
5	Manganese scavenging and oxidative stress response mediated by type VI secretion system in <i>Burkholderia thailandensis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E2233-E2242.	7.1	185
6	The Pseudomonas Quinolone Signal (PQS): Not Just for Quorum Sensing Anymore. Frontiers in Cellular and Infection Microbiology, 2018, 8, 230.	3.9	178
7	Type VI Secretion System Transports Zn2+ to Combat Multiple Stresses and Host Immunity. PLoS Pathogens, 2015, 11, e1005020.	4.7	169
8	Pantoea alhagi, a novel endophytic bacterium with ability to improve growth and drought tolerance in wheat. Scientific Reports, 2017, 7, 41564.	3.3	129
9	Targeting eEF1A by a <i>Legionella pneumophila</i> effector leads to inhibition of protein synthesis and induction of host stress response. Cellular Microbiology, 2009, 11, 911-926.	2.1	128
10	The Type VI Secretion System Engages a Redox-Regulated Dual-Functional Heme Transporter for Zinc Acquisition. Cell Reports, 2017, 20, 949-959.	6.4	107
11	A type <scp>VI</scp> secretion system regulated by <scp>OmpR</scp> in <i><scp>Y</scp>ersinia pseudotuberculosis</i> functions to maintain intracellular <scp>pH</scp> homeostasis. Environmental Microbiology, 2013, 15, 557-569.	3.8	99
12	Sensing of autoinducer-2 by functionally distinct receptors in prokaryotes. Nature Communications, 2020, 11, 5371.	12.8	86
13	Effector–Immunity Pairs Provide the T6SS Nanomachine its Offensive and Defensive Capabilities. Molecules, 2018, 23, 1009.	3.8	69
14	Roles of RpoS in Yersinia pseudotuberculosis stress survival, motility, biofilm formation and type VI secretion system expression. Journal of Microbiology, 2015, 53, 633-642.	2.8	59
15	Taibaiella smilacinae gen. nov., sp. nov., an endophytic member of the family Chitinophagaceae isolated from the stem of Smilacina japonica, and emended description of Flavihumibacter petaseus. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3769-3776.	1.7	55
16	Modulation of a thermoregulated type VI secretion system by AHL-dependent Quorum Sensing in Yersinia pseudotuberculosis. Archives of Microbiology, 2011, 193, 351-63.	2.2	50
17	<scp>FliS</scp> modulates <scp>FlgM</scp> activity by acting as a nonâ€canonical chaperone to control late flagellar gene expression, motility and biofilm formation in <scp><i>Y</i></scp> <i>ersinia pseudotuberculosis</i> . Environmental Microbiology, 2014, 16, 1090-1104.	3.8	50
18	Extracellular matrix-associated proteins form an integral and dynamic system during Pseudomonas aeruginosa biofilm development. Frontiers in Cellular and Infection Microbiology, 2015, 5, 40.	3.9	48

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19	Deciphering the Root Endosphere Microbiome of the Desert Plant <i>Alhagi sparsifolia</i> for Drought Resistance-Promoting Bacteria. Applied and Environmental Microbiology, 2020, 86, .	3.1	44
20	T6SS secretes an LPS-binding effector to recruit OMVs for exploitative competition and horizontal gene transfer. ISME Journal, 2022, 16, 500-510.	9.8	44
21	Functional characterization of a mycothiol peroxidase in <i>Corynebacterium glutamicum</i> that uses both mycoredoxin and thioredoxin reducing systems in the response to oxidative stress. Biochemical Journal, 2015, 469, 45-57.	3.7	43
22	Corynebacterium glutamicum Methionine Sulfoxide Reductase A Uses both Mycoredoxin and Thioredoxin for Regeneration and Oxidative Stress Resistance. Applied and Environmental Microbiology, 2015, 81, 2781-2796.	3.1	42
23	Confirmed and Potential Roles of Bacterial T6SSs in the Intestinal Ecosystem. Frontiers in Microbiology, 2019, 10, 1484.	3.5	42
24	Contact-independent killing mediated by a T6SS effector with intrinsic cell-entry properties. Nature Communications, 2021, 12, 423.	12.8	42
25	Functional characterization of a vanillin dehydrogenase in Corynebacterium glutamicum. Scientific Reports, 2015, 5, 8044.	3.3	39
26	In Vivo Analysis of Protein–Protein Interactions with Bioluminescence Resonance Energy Transfer (BRET): Progress and Prospects. International Journal of Molecular Sciences, 2016, 17, 1704.	4.1	37
27	The icmF3 locus is involved in multiple adaptation- and virulence-related characteristics in Pseudomonas aeruginosa PAO1. Frontiers in Cellular and Infection Microbiology, 2015, 5, 70.	3.9	35
28	Type VI Secretion Systems Present New Insights on Pathogenic Yersinia. Frontiers in Cellular and Infection Microbiology, 2018, 8, 260.	3.9	33
29	Overexpression of Mycothiol Disulfide Reductase Enhances Corynebacterium glutamicum Robustness by Modulating Cellular Redox Homeostasis and Antioxidant Proteins under Oxidative Stress. Scientific Reports, 2016, 6, 29491.	3.3	32
30	ISCR2 is associated with the dissemination of multiple resistance genes among Vibrio spp. and Pseudoalteromonas spp. isolated from farmed fish. Archives of Microbiology, 2017, 199, 891-896.	2.2	31
31	Mycothiol protects <i>Corynebacterium glutamicum</i> against acid stress via maintaining intracellular pH homeostasis, scavenging ROS, and <i>S</i> -mycothiolating MetE. Journal of General and Applied Microbiology, 2016, 62, 144-153.	0.7	30
32	Siderophore-Mediated Iron Acquisition Enhances Resistance to Oxidative and Aromatic Compound Stress in <i>Cupriavidus necator</i> JMP134. Applied and Environmental Microbiology, 2019, 85, .	3.1	30
33	Graded Response of the Multifunctional 2-Cysteine Peroxiredoxin, CgPrx, to Increasing Levels of Hydrogen Peroxide in <i>Corynebacterium glutamicum</i> . Antioxidants and Redox Signaling, 2017, 26, 1-14.	5.4	28
34	A starvation-induced regulator, RovM, acts as a switch for planktonic/biofilm state transition in Yersinia pseudotuberculosis. Scientific Reports, 2017, 7, 639.	3.3	28
35	Ohr Protects Corynebacterium glutamicum against Organic Hydroperoxide Induced Oxidative Stress. PLoS ONE, 2015, 10, e0131634.	2.5	28
36	T6SS translocates a micropeptide to suppress STING-mediated innate immunity by sequestering manganese. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	27

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37	The dual transcriptional regulator <scp>RovM</scp> regulates the expression of <scp>AR3</scp> ―and <scp>T6SS4</scp> â€dependent acid survival systems in response to nutritional status in <scp><i>Y</i></scp> <i>ersinia pseudotuberculosis</i> . Environmental Microbiology, 2015, 17, 4631-4645.	3.8	24
38	Sphingomonas gei sp. nov., isolated from roots of Geum aleppicum. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 1160-1166.	1.7	24
39	Solirubrobacter phytolaccae sp. nov., an endophytic bacterium isolated from roots of Phytolacca acinosa Roxb International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 858-862.	1.7	23
40	Roles of Type VI Secretion System in Transport of Metal Ions. Frontiers in Microbiology, 2021, 12, 756136.	3.5	23
41	Molecular Mechanisms of AhpC in Resistance to Oxidative Stress in Burkholderia thailandensis. Frontiers in Microbiology, 2019, 10, 1483.	3.5	22
42	Rhizobium smilacinae sp. nov., an endophytic bacterium isolated from the leaf of Smilacina japonica. Antonie Van Leeuwenhoek, 2014, 106, 715-723.	1.7	21
43	Pseudoxanthomonas gei sp. nov., a novel endophytic bacterium isolated from the stem of Geum aleppicum. Antonie Van Leeuwenhoek, 2014, 105, 653-661.	1.7	20
44	ZntR positively regulates T6SS4 expression in Yersinia pseudotuberculosis. Journal of Microbiology, 2017, 55, 448-456.	2.8	20
45	Pontibacter toksunensis sp. nov., isolated from soil, and emended descriptions of Pontibacter roseus and Pontibacter akesuensis. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 4462-4468.	1.7	19
46	Functional Characterization of Corynebacterium glutamicum Mycothiol S-Conjugate Amidase. PLoS ONE, 2014, 9, e115075.	2.5	19
47	Rhizobacter bergeniae sp. nov., isolated from the root of Bergenia scopulosa. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 479-484.	1.7	19
48	Nafulsella turpanensis gen. nov., sp. nov., a member of the phylum Bacteroidetes isolated from soil. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 1639-1645.	1.7	18
49	Transcriptional control of the phenol hydroxylase gene phe of Corynebacterium glutamicum by the AraC-type regulator PheR. Microbiological Research, 2018, 209, 14-20.	5.3	18
50	Global Transcriptomic Analysis of the Response of Corynebacterium glutamicum to Vanillin. PLoS ONE, 2016, 11, e0164955.	2.5	18
51	Asticcacaulis endophyticus sp. nov., a prosthecate bacterium isolated from the root of Geum aleppicum. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 3964-3969.	1.7	17
52	Complete genome sequence of the drought resistance-promoting endophyte Klebsiella sp. LTGPAF-6F. Journal of Biotechnology, 2017, 246, 36-39.	3.8	17
53	Engineering an Enhanced, Thermostable, Monomeric Bacterial Luciferase Gene As a Reporter in Plant Protoplasts. PLoS ONE, 2014, 9, e107885.	2.5	16
54	Sphingobium endophyticus sp. nov., isolated from the root of Hylomecon japonica. Antonie Van Leeuwenhoek, 2015, 107, 1001-1008.	1.7	16

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55	Global transcriptomic analysis of the response of Corynebacterium glutamicum to ferulic acid. Archives of Microbiology, 2017, 199, 325-334.	2.2	16
56	RovM and CsrA Negatively Regulate Urease Expression in Yersinia pseudotuberculosis. Frontiers in Microbiology, 2018, 9, 348.	3.5	16
57	The Catabolite Repressor/Activator Cra Is a Bridge Connecting Carbon Metabolism and Host Colonization in the Plant Drought Resistance-Promoting Bacterium Pantoea alhagi LTYR-11Z. Applied and Environmental Microbiology, 2018, 84, .	3.1	15
58	An Osmoregulatory Mechanism Operating through OmpR and LrhA Controls the Motile-Sessile Switch in the Plant Growth-Promoting Bacterium <i>Pantoea alhagi</i> . Applied and Environmental Microbiology, 2019, 85, .	3.1	15
59	The transcriptional regulator Zur regulates the expression of ZnuABC and T6SS4 in response to stresses in Yersinia pseudotuberculosis. Microbiological Research, 2021, 249, 126787.	5.3	15
60	Paenibacillus sinopodophylli sp. nov., a siderophore-producing endophytic bacterium isolated from roots of Sinopodophyllum hexandrum (Royle) Ying. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4993-4999.	1.7	15
61	Paenibacillus qinlingensis sp. nov., an indole-3-acetic acid-producing bacterium isolated from roots of Sinopodophyllum hexandrum (Royle) Ying. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 589-595.	1.7	14
62	Propioniciclava sinopodophylli sp. nov., isolated from leaves of Sinopodophyllum hexandrum (Royle) Ying. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 4111-4115.	1.7	14
63	Solirubrobacter taibaiensis sp. nov., isolated from a stem of Phytolacca acinosa Roxb Antonie Van Leeuwenhoek, 2014, 106, 279-285.	1.7	13
64	Aerobactin-Mediated Iron Acquisition Enhances Biofilm Formation, Oxidative Stress Resistance, and Virulence of Yersinia pseudotuberculosis. Frontiers in Microbiology, 2021, 12, 699913.	3.5	13
65	Sphingomonas hylomeconis sp. nov., isolated from the stem of Hylomecon japonica. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 4025-4031.	1.7	13
66	Rhizobium gei sp. nov., a bacterial endophyte of Geum aleppicum. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4282-4288.	1.7	13
67	Ficolin-2 binds to HIV-1 gp120 and blocks viral infection. Virologica Sinica, 2016, 31, 406-414.	3.0	12
68	Functional characterization of a csoR-cueA divergon in Bradyrhizobium liaoningense CCNWSX0360, involved in copper, zinc and cadmium cotolerance. Scientific Reports, 2016, 6, 35155.	3.3	12
69	Mycothiol peroxidase MPx protects Corynebacterium glutamicum against acid stress by scavenging ROS. Biotechnology Letters, 2016, 38, 1221-1228.	2.2	10
70	Manipulation of the silkworm immune system by a metalloprotease from the pathogenic bacterium Pseudomonas aeruginosa. Developmental and Comparative Immunology, 2019, 90, 176-185.	2.3	10
71	Identification of Robinia pseudoacacia target proteins responsive to Mesorhizobium amphore CCNWGS0123 effector protein NopT. Journal of Experimental Botany, 2020, 71, 7347-7363.	4.8	10
72	Salmonella Induces the cGAS-STING-Dependent Type I Interferon Response in Murine Macrophages by Triggering mtDNA Release. MBio, 2022, 13, .	4.1	10

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73	HpaR, the Repressor of Aromatic Compound Metabolism, Positively Regulates the Expression of T6SS4 to Resist Oxidative Stress in Yersinia pseudotuberculosis. Frontiers in Microbiology, 2020, 11, 705.	3.5	9
74	Bioluminescence Resonance Energy Transfer System for Measuring Dynamic Protein-Protein Interactions in Bacteria. MBio, 2014, 5, e01050-14.	4.1	8
75	Beyond dueling: roles of the type VI secretion system in microbiome modulation, pathogenesis and stress resistance. Stress Biology, 2021, 1, 1.	3.1	8
76	Zinc acquisition via ZnuABC in Yersinia pseudotuberculosis facilitates resistance to oxidative stress. Annals of Microbiology, 2016, 66, 1189-1197.	2.6	7
77	Molecular characterization of a eukaryotic-like phenol hydroxylase from <i>Corynebacterium glutamicum</i> . Journal of General and Applied Microbiology, 2015, 61, 99-107.	0.7	6
78	Functional comparison of methionine sulphoxide reductase A and B in <i>Corynebacterium glutamicum</i> . Journal of General and Applied Microbiology, 2017, 63, 280-286.	0.7	5
79	A c-di-GMP Signaling Cascade Controls Motility, Biofilm Formation, and Virulence in Burkholderia thailandensis. Applied and Environmental Microbiology, 2022, 88, e0252921.	3.1	4