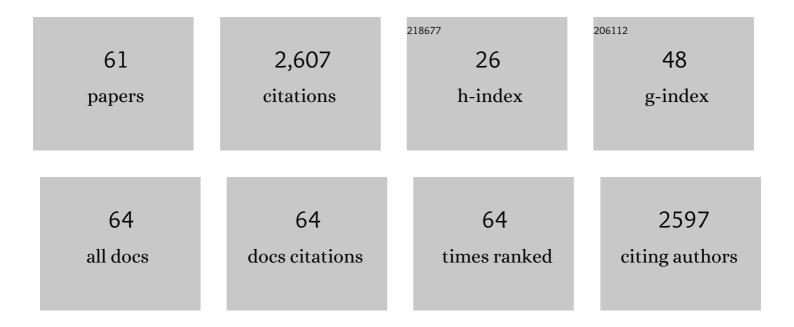
## Yao Wang

## List of Publications by Year in descending order

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#	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	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
3	The Pseudomonas Quinolone Signal (PQS): Not Just for Quorum Sensing Anymore. Frontiers in Cellular and Infection Microbiology, 2018, 8, 230.	3.9	178
4	Type VI Secretion System Transports Zn2+ to Combat Multiple Stresses and Host Immunity. PLoS Pathogens, 2015, 11, e1005020.	4.7	169
5	Pantoea alhagi, a novel endophytic bacterium with ability to improve growth and drought tolerance in wheat. Scientific Reports, 2017, 7, 41564.	3.3	129
6	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
7	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
8	Sensing of autoinducer-2 by functionally distinct receptors in prokaryotes. Nature Communications, 2020, 11, 5371.	12.8	86
9	Physiological roles of mycothiol in detoxification and tolerance to multiple poisonous chemicals in Corynebacterium glutamicum. Archives of Microbiology, 2013, 195, 419-429.	2.2	80
10	OmpR positively regulates urease expression to enhance acid survival of Yersinia pseudotuberculosis. Microbiology (United Kingdom), 2009, 155, 2522-2531.	1.8	66
11	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
12	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
13	Functional interplay between the <i>Yersinia pseudotuberculosis</i> YpsRI and YtbRI quorum sensing systems modulates swimming motility by controlling expression of <i>flhDC</i> and <i>fliA</i> . Molecular Microbiology, 2008, 69, 137-151.	2.5	53
14	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
15	<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
16	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
17	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
18	NrdH Redoxin Enhances Resistance to Multiple Oxidative Stresses by Acting as a Peroxidase Cofactor in Corynebacterium glutamicum. Applied and Environmental Microbiology, 2014, 80, 1750-1762.	3.1	43

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19	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
20	Genome-wide identification of and functional insights into the late embryogenesis abundant (LEA) gene family in bread wheat (Triticum aestivum). Scientific Reports, 2019, 9, 13375.	3.3	42
21	Contact-independent killing mediated by a T6SS effector with intrinsic cell-entry properties. Nature Communications, 2021, 12, 423.	12.8	42
22	Biosorption of Cadmium and Manganese Using Free Cells of Klebsiella sp. Isolated from Waste Water. PLoS ONE, 2015, 10, e0140962.	2.5	39
23	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
24	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
25	Type VI Secretion Systems Present New Insights on Pathogenic Yersinia. Frontiers in Cellular and Infection Microbiology, 2018, 8, 260.	3.9	33
26	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
27	Effects of quorum sensing autoinducer degradation gene on virulence and biofilm formation of Pseudomonas aeruginosa. Science in China Series C: Life Sciences, 2007, 50, 385-391.	1.3	29
28	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
29	Ohr Protects Corynebacterium glutamicum against Organic Hydroperoxide Induced Oxidative Stress. PLoS ONE, 2015, 10, e0131634.	2.5	28
30	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
31	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
32	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
33	Positive regulation of flhDC expression by OmpR in Yersinia pseudotuberculosis. Microbiology (United Kingdom), 2009, 155, 3622-3631.	1.8	21
34	Rhizobium smilacinae sp. nov., an endophytic bacterium isolated from the leaf of Smilacina japonica. Antonie Van Leeuwenhoek, 2014, 106, 715-723.	1.7	21
35	Functional characterization of FlgM in the regulation of flagellar synthesis and motility in Yersinia pseudotuberculosis. Microbiology (United Kingdom), 2009, 155, 1890-1900.	1.8	20
36	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

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37	ZntR positively regulates T6SS4 expression in Yersinia pseudotuberculosis. Journal of Microbiology, 2017, 55, 448-456.	2.8	20
38	Transgenic Paulownia Expressing shiva-1 Gene Has Increased Resistance to Paulownia Witches' Broom Disease. Journal of Integrative Plant Biology, 2005, 47, 1500-1506.	8.5	19
39	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
40	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
41	Enhancing Corynebacterium glutamicum robustness by over-expressing a gene, mshA, for mycothiol glycosyltransferase. Biotechnology Letters, 2014, 36, 1453-1459.	2.2	18
42	Global Transcriptomic Analysis of the Response of Corynebacterium glutamicum to Vanillin. PLoS ONE, 2016, 11, e0164955.	2.5	18
43	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
44	Complete genome sequence of the drought resistance-promoting endophyte Klebsiella sp. LTGPAF-6F. Journal of Biotechnology, 2017, 246, 36-39.	3.8	17
45	Engineering an Enhanced, Thermostable, Monomeric Bacterial Luciferase Gene As a Reporter in Plant Protoplasts. PLoS ONE, 2014, 9, e107885.	2.5	16
46	Global transcriptomic analysis of the response of Corynebacterium glutamicum to ferulic acid. Archives of Microbiology, 2017, 199, 325-334.	2.2	16
47	RovM and CsrA Negatively Regulate Urease Expression in Yersinia pseudotuberculosis. Frontiers in Microbiology, 2018, 9, 348.	3.5	16
48	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
49	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
50	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
51	Aerobactin-Mediated Iron Acquisition Enhances Biofilm Formation, Oxidative Stress Resistance, and Virulence of Yersinia pseudotuberculosis. Frontiers in Microbiology, 2021, 12, 699913.	3.5	13
52	Rhizobium gei sp. nov., a bacterial endophyte of Geum aleppicum. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 4282-4288.	1.7	13
53	The flhDC gene affects motility and biofilm formation in Yersinia pseudotuberculosis. Science in China Series C: Life Sciences, 2007, 50, 814-821.	1.3	11
54	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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55	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
56	Bioluminescence Resonance Energy Transfer System for Measuring Dynamic Protein-Protein Interactions in Bacteria. MBio, 2014, 5, e01050-14.	4.1	8
57	The stringent response factor, RelA, positively regulates T6SS4 expression through the RovM/RovA pathway in Yersinia pseudotuberculosis. Microbiological Research, 2019, 220, 32-41.	5.3	8
58	Zinc acquisition via ZnuABC in Yersinia pseudotuberculosis facilitates resistance to oxidative stress. Annals of Microbiology, 2016, 66, 1189-1197.	2.6	7
59	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
60	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
61	Isolation and identification of canine parvovirus serotype 2a and its VP2 protein expression in transgenic tobacco. Virologica Sinica, 2008, 23, 203-210.	3.0	2