

# Yanqin Ding

## List of Publications by Year in descending order

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Version: 2024-02-01

24  
papers

537  
citations

623734

14  
h-index

677142

22  
g-index

24  
all docs

24  
docs citations

24  
times ranked

597  
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Metabacillus dongyingensis</i> sp. nov. Is Represented by the Plant Growth-Promoting Bacterium BY2G20 Isolated from Saline-Alkaline Soil and Enhances the Growth of <i>Zea mays</i> L. under Salt Stress. <i>MSystems</i> , 2022, 7, e0142621.	3.8	9
2	Isolation and Genome Sequence of a Novel Phosphate-Solubilizing Rhizobacterium <i>Bacillus altitudinis</i> GQYP101 and Its Effects on Rhizosphere Microbial Community Structure and Functional Traits of Corn Seedling. <i>Current Microbiology</i> , 2022, 79, .	2.2	7
3	Interactional mechanisms of <i>Paenibacillus polymyxa</i> SC2 and pepper ( <i>Capsicum annuum</i> L.) suggested by transcriptomics. <i>BMC Microbiology</i> , 2021, 21, 70.	3.3	20
4	Plant metabolomics integrated with transcriptomics and rhizospheric bacterial community indicates the mitigation effects of <i>Klebsiella oxytoca</i> P620 on p-hydroxybenzoic acid stress in cucumber. <i>Journal of Hazardous Materials</i> , 2021, 415, 125756.	12.4	25
5	Transcriptome Profiles Reveal the Growth-Promoting Mechanisms of <i>Paenibacillus polymyxa</i> YC0136 on Tobacco ( <i>Nicotiana tabacum</i> L.). <i>Frontiers in Microbiology</i> , 2020, 11, 584174.	3.5	34
6	Identification of a native promoter P for gene expression in <i>Paenibacillus polymyxa</i> . <i>Journal of Biotechnology</i> , 2019, 295, 19-27.	3.8	7
7	Comparative genomic analysis of <i>Bacillus paralicheniformis</i> MDJK30 with its closely related species reveals an evolutionary relationship between <i>B. paralicheniformis</i> and <i>B. licheniformis</i> . <i>BMC Genomics</i> , 2019, 20, 283.	2.8	34
8	Complete Genome Sequence of <i>Bacillus velezensis</i> DSYZ, a Plant Growth-Promoting Rhizobacterium with Antifungal Properties. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.6	4
9	Effects of <i>Bacillus velezensis</i> FKM10 for Promoting the Growth of <i>Malus hupehensis</i> Rehd. and Inhibiting <i>Fusarium verticillioides</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 2889.	3.5	52
10	Screening and Whole-Genome Sequencing of Two <i>Streptomyces</i> Species from the Rhizosphere Soil of Peony Reveal Their Characteristics as Plant Growth-Promoting Rhizobacteria. <i>BioMed Research International</i> , 2018, 2018, 1-11.	1.9	12
11	Analysis of the Complete Genome Sequence of <i>Bacillus atrophaeus</i> GQJK17 Reveals Its Biocontrol Characteristics as a Plant Growth-Promoting Rhizobacterium. <i>BioMed Research International</i> , 2018, 2018, 1-9.	1.9	21
12	Complete Genome Sequence of <i>Paenibacillus polymyxa</i> YC0136, a Plant Growth-Promoting Rhizobacterium Isolated from Tobacco Rhizosphere. <i>Genome Announcements</i> , 2017, 5, .	0.8	18
13	Complete Genome Sequence of <i>Paenibacillus polymyxa</i> YC0573, a Plant Growth-Promoting Rhizobacterium with Antimicrobial Activity. <i>Genome Announcements</i> , 2017, 5, .	0.8	16
14	Promotion of iron nutrition and growth on peanut by <i>Paenibacillus illinoisensis</i> and <i>Bacillus</i> sp. strains in calcareous soil. <i>Brazilian Journal of Microbiology</i> , 2017, 48, 656-670.	2.0	63
15	Complete Genome Sequence of <i>Bacillus subtilis</i> GQJK2, a Plant Growth-Promoting Rhizobacterium with Antifungal Activity. <i>Genome Announcements</i> , 2017, 5, .	0.8	4
16	Complete Genome Sequence of <i>Bacillus velezensis</i> GQJK49, a Plant Growth-Promoting Rhizobacterium with Antifungal Activity. <i>Genome Announcements</i> , 2017, 5, .	0.8	4
17	Complete Genome Sequence of <i>Bacillus paralicheniformis</i> MDJK30, a Plant Growth-Promoting Rhizobacterium with Antifungal Activity. <i>Genome Announcements</i> , 2017, 5, .	0.8	20
18	Identification of Important Amino Acids in Gal2p for Improving the L-arabinose Transport and Metabolism in <i>Saccharomyces cerevisiae</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 1391.	3.5	22

#	ARTICLE	IF	CITATIONS
19	Complete Genome Sequence of Biocontroller <i>Bacillus velezensis</i> Strain JTYP2, Isolated from Leaves of <i>Echeveria laui</i> . <i>Genome Announcements</i> , 2017, 5, .	0.8	3
20	Coutilization of D-Glucose, D-Xylose, and L-Arabinose in <i>Saccharomyces cerevisiae</i> by Coexpressing the Metabolic Pathways and Evolutionary Engineering. <i>BioMed Research International</i> , 2017, 2017, 1-8.	1.9	15
21	Comparative Genomic Analysis of <i>Delftia tsuruhatensis</i> MTQ3 and the Identification of Functional NRPS Genes for Siderophore Production. <i>BioMed Research International</i> , 2016, 2016, 1-8.	1.9	17
22	A single amino acid mutation in Spo0A results in sporulation deficiency of <i>Paenibacillus polymyxa</i> SC2. <i>Research in Microbiology</i> , 2016, 167, 472-479.	2.1	22
23	Draft Genome Sequence of <i>Delftia tsuruhatensis</i> MTQ3, a Strain of Plant Growth-Promoting Rhizobacterium with Antimicrobial Activity. <i>Genome Announcements</i> , 2015, 3, .	0.8	14
24	Complete Genome Sequence of <i>Paenibacillus polymyxa</i> SC2, a Strain of Plant Growth-Promoting Rhizobacterium with Broad-Spectrum Antimicrobial Activity. <i>Journal of Bacteriology</i> , 2011, 193, 311-312.	2.2	94