Jiguo Qiu

List of Publications by Year in descending order

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331259 454577 1,381 89 21 30 citations h-index g-index papers 92 92 92 1189 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Characterization of a new heterotrophic nitrification bacterium Pseudomonas sp. strain JQ170 and functional identification of nap gene in nitrite production. Science of the Total Environment, 2022, 806, 150556.	3.9	7
2	Anaerobic biodegradation and detoxification of chloroacetamide herbicides by a novel Proteiniclasticum sediminis BAD-10T. Environmental Research, 2022, 209, 112859.	3.7	17
3	Genetic Foundations of Direct Ammonia Oxidation (Dirammox) to N ₂ and MocR-Like Transcriptional Regulator DnfR in Alcaligenes faecalis Strain JQ135. Applied and Environmental Microbiology, 2022, 88, aem0226121.	1.4	9
4	Degradation of dimethachlon by a newly isolated bacterium Paenarthrobacter sp. strain JH-1 relieves its toxicity against Chlorella ellipsoidea. Environmental Research, 2022, 208, 112706.	3.7	3
5	The TetR Family Repressor HpaR Negatively Regulates the Catabolism of 5-Hydroxypicolinic Acid in Alcaligenes faecalis JQ135 by Binding to Two Unique DNA Sequences in the Promoter of <i>Hpa</i> Operon. Applied and Environmental Microbiology, 2022, 88, aem0239021.	1.4	3
6	Two LysR Family Transcriptional Regulators, McbH and McbN, Activate the Operons Responsible for the Midstream and Downstream Pathways, Respectively, of Carbaryl Degradation in Pseudomonas sp. Strain XWY-1. Applied and Environmental Microbiology, 2022, 88, AEM0206021.	1.4	5
7	Pseudaminobacter soli sp. nov., Isolated from Paddy Soil Contaminated with Heavy Metals. Current Microbiology, 2022, 79, 19.	1.0	1
8	Biodegradation of Quinoline by a Newly Isolated Salt-Tolerating Bacterium Rhodococcus gordoniae Strain JH145. Microorganisms, 2022, 10, 797.	1.6	6
9	Biodegradation of Quinolinic acid by a Newly Isolated Bacterium $\langle i \rangle$ Alcaligenes faecalis $\langle i \rangle$ Strain JQ191. FEMS Microbiology Letters, 2022, , .	0.7	1
10	The Novel Amidase PcnH Initiates the Degradation of Phenazine-1-Carboxamide in Sphingomonas histidinilytica DS-9. Applied and Environmental Microbiology, 2022, 88, e0054322.	1.4	3
11	PicR as a MarR Family Transcriptional Repressor Multiply Controls the Transcription of Picolinic Acid Degradation Gene Cluster <i>pic</i> in Alcaligenes faecalis JQ135. Applied and Environmental Microbiology, 2022, 88, .	1.4	6
12	The Novel Monooxygenase Gene <i>dipD</i> in the <i>dip</i> Gene Cluster of <i>Alcaligenes faecalis</i> JQ135 Is Essential for the Initial Catabolism of Dipicolinic Acid. Applied and Environmental Microbiology, 2022, 88, .	1.4	2
13	Substrate preference of carbamate hydrolase CehA reveals its environmental behavior. Journal of Hazardous Materials, 2021, 403, 123677.	6.5	12
14	Enhanced recovery of hexavalent chromium by remodeling extracellular polymeric substances through engineering Agrobacterium tumefaciens F2. Journal of Cleaner Production, 2021, 279, 123829.	4.6	26
15	Comparative genomic analysis of iprodioneâ€degrading <i>Paenarthrobacter</i> strains reveals the iprodione catabolic molecular mechanism in <i>Paenarthrobacter</i> sp. strain <scp>YJN</scp> â€5. Environmental Microbiology, 2021, 23, 1079-1095.	1.8	8
16	Identification and characterization of Nornicotine degrading strain Arthrobacter sp. NOR5. Science of the Total Environment, 2021, 764, 142894.	3.9	5
17	Lysobacter gilvus sp. nov., isolated from activated sludge. Archives of Microbiology, 2021, 203, 7-11.	1.0	3
18	McbG, a LysR Family Transcriptional Regulator, Activates the <i>mcbBCDEF</i> Gene Cluster Involved in the Upstream Pathway of Carbaryl Degradation in <i>Pseudomonas</i> sp. Strain XWY-1. Applied and Environmental Microbiology, 2021, 87, .	1.4	13

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19	A novel hydrolase <scp>PyzH</scp> catalyses the cleavage of C=N double bond for pymetrozine degradation in <i>Pseudomonas</i> sp. <scp>BYT</scp> â€1. Environmental Microbiology, 2021, 23, 3265-3273.	1.8	6
20	Unveiling the CoA mediated salicylate catabolic mechanism in <i>Rhizobium </i> sp. X9. Molecular Microbiology, 2021, 116, 783-793.	1.2	4
21	Enhanced degradation of dicamba by an anaerobic sludge acclimated from river sediment. Science of the Total Environment, 2021, 777, 145931.	3.9	26
22	Cotinine Hydroxylase CotA Initiates Biodegradation of Wastewater Micropollutant Cotinine in <i>Nocardioides</i> sp. Strain JQ2195. Applied and Environmental Microbiology, 2021, 87, e0092321.	1.4	9
23	Sinanaerobacter chloroacetimidivorans gen. nov., sp. nov., an obligate anaerobic bacterium isolated from anaerobic sludge. Antonie Van Leeuwenhoek, 2021, 114, 1609-1617.	0.7	1
24	Rhodobacter kunshanensis sp. nov., a Novel Bacterium Isolated from Activated Sludge. Current Microbiology, 2021, 78, 3791-3797.	1.0	2
25	Catabolic characterization of dipicolinic acid in Alcaligenes faecalis strain JQ135. International Biodeterioration and Biodegradation, 2021, 165 , 105312 .	1.9	5
26	Identification of the key amino acid sites of the carbofuran hydrolase CehA from a newly isolated carbofuran-degrading strain Sphingbium sp. CFD-1. Ecotoxicology and Environmental Safety, 2020, 189, 109938.	2.9	26
27	An angular dioxygenase gene cluster responsible for the initial phenazine-1-carboxylic acid degradation step in Rhodococcus sp. WH99 can protect sensitive organisms from toxicity. Science of the Total Environment, 2020, 706, 135726.	3.9	7
28	Anaerobic biodegradation of acetochlor by acclimated sludge and its anaerobic catabolic pathway. Science of the Total Environment, 2020, 748, 141122.	3.9	31
29	Crenobacter caeni sp. nov. Isolated from Sludge. Current Microbiology, 2020, 77, 4180-4185.	1.0	1
30	Carbamate C-N Hydrolase Gene <i>ameH</i> Responsible for the Detoxification Step of Methomyl Degradation in Aminobacter aminovorans Strain MDW-2. Applied and Environmental Microbiology, 2020, 87, .	1.4	11
31	Caenimonas sedimenti sp. nov., Isolated from Sediment of the Wastewater Outlet of an Agricultural Chemical Plant. Current Microbiology, 2020, 77, 3767-3772.	1.0	2
32	Roles of the Gentisate 1,2-Dioxygenases DsmD and GtdA in the Catabolism of the Herbicide Dicamba in Rhizorhabdus dicambivorans Ndbn-20. Journal of Agricultural and Food Chemistry, 2020, 68, 9287-9298.	2.4	4
33	Applied microbiology and biotechnology uncovering the biosynthetic pathway of polysaccharide-based microbial flocculant in Agrobacterium tumefaciens F2. Applied Microbiology and Biotechnology, 2020, 104, 8479-8488.	1.7	7
34	Pedobacter puniceum sp. nov. Isolated from Sludge. Current Microbiology, 2020, 77, 4186-4191.	1.0	3
35	Rudanella paleaurantiibacter sp. nov., Isolated from Activated Sludge. Current Microbiology, 2020, 77, 2016-2022.	1.0	1
36	Bacterial catabolism of nicotine: Catabolic strains, pathways and modules. Environmental Research, 2020, 183, 109258.	3.7	24

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37	Sphingobacterium olei sp. nov., isolated from oil-contaminated soil. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 1931-1939.	0.8	13
38	Niastella caeni sp. nov., isolated from activated sludge. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2261-2268.	0.8	6
39	Humidity Control Strategies for Solid-State Fermentation: Capillary Water Supply by Water-Retention Materials and Negative-Pressure Auto-controlled Irrigation. Frontiers in Bioengineering and Biotechnology, 2019, 7, 263.	2.0	27
40	Novel 3,6-Dihydroxypicolinic Acid Decarboxylase-Mediated Picolinic Acid Catabolism in <i>Alcaligenes faecalis</i> JQ135. Journal of Bacteriology, 2019, 201, .	1.0	8
41	Genome Analysis of Carbaryl-Degrading Strain Pseudomonas putida XWY-1. Current Microbiology, 2019, 76, 927-929.	1.0	24
42	The Properties of 5-Methyltetrahydrofolate Dehydrogenase (MetF1) and Its Role in the Tetrahydrofolate-Dependent Dicamba Demethylation System in Rhizorhabdus dicambivorans Ndbn-20. Journal of Bacteriology, 2019, 201, .	1.0	5
43	Identification and Characterization of a Novel <i>pic</i> Gene Cluster Responsible for Picolinic Acid Degradation in Alcaligenes faecalis JQ135. Journal of Bacteriology, 2019, 201, .	1.0	18
44	Comparative genome analysis reveals the evolution of chloroacetanilide herbicide mineralization in Sphingomonas wittichii DC-6. Archives of Microbiology, 2019, 201, 907-918.	1.0	10
45	Biodegradation of Picolinic Acid by Rhodococcus sp. PA18. Applied Sciences (Switzerland), 2019, 9, 1006.	1.3	4
46	Isolation and Characterization of the Pymetrozine-Degrading Strain <i>Pseudomonas</i> sp. BYT-1. Journal of Agricultural and Food Chemistry, 2019, 67, 4170-4176.	2.4	14
47	Complete Genome Sequence of Alcaligenes Faecalis Strain JQ135, a Bacterium Capable of Efficiently Degrading Nicotinic Acid. Current Microbiology, 2018, 75, 1551-1554.	1.0	21
48	Biodegradation of Pendimethalin by Paracoccus sp. P13. Current Microbiology, 2018, 75, 1077-1083.	1.0	14
49	Isolation and characterization of the cotinine-degrading bacterium Nocardioides sp. Strain JQ2195. Journal of Hazardous Materials, 2018, 353, 158-165.	6.5	22
50	Identification and characterization of a new three-component nicotinic acid hydroxylase NahAB1B2fromPusillimonassp. strain T2. Letters in Applied Microbiology, 2018, 66, 321-328.	1.0	5
51	Cloning and expression of the carbaryl hydrolase gene mcbA and the identification of a key amino acid necessary for carbaryl hydrolysis. Journal of Hazardous Materials, 2018, 344, 1126-1135.	6.5	36
52	3,6-Dichlorosalicylate Catabolism Is Initiated by the DsmABC Cytochrome P450 Monooxygenase System in Rhizorhabdus dicambivorans Ndbn-20. Applied and Environmental Microbiology, 2018, 84, .	1.4	13
53	A Novel Degradation Mechanism for Pyridine Derivatives in Alcaligenes faecalis JQ135. Applied and Environmental Microbiology, 2018, 84, .	1.4	30
54	Roles of Two Glutathione-Dependent 3,6-Dichlorogentisate Dehalogenases in Rhizorhabdus dicambivorans Ndbn-20 in the Catabolism of the Herbicide Dicamba. Applied and Environmental Microbiology, 2018, 84, .	1.4	12

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55	Optimization of fed-batch fermentation and direct spray drying in the preparation of microbial inoculant of acetochlor-degrading strain Sphingomonas sp. DC-6. 3 Biotech, 2018, 8, 294.	1.1	6
56	Flavobacterium zaozhuangense sp. nov., a new member of the family Flavobacteriaceae, isolated from metolachlor-contaminated soil. Antonie Van Leeuwenhoek, 2018, 111, 1977-1984.	0.7	6
57	Stereoselective accumulations of hexachlorocyclohexanes (HCHs) are correlated with Sphingomonas spp. in agricultural soils across China. Environmental Pollution, 2018, 240, 27-33.	3.7	20
58	An Amidase Gene, <i>ipaH</i> , Is Responsible for the Initial Step in the Iprodione Degradation Pathway of Paenarthrobacter sp. Strain YJN-5. Applied and Environmental Microbiology, 2018, 84, .	1.4	34
59	Hydrolase CehA and Monooxygenase CfdC Are Responsible for Carbofuran Degradation in Sphingomonas sp. Strain CDS-1. Applied and Environmental Microbiology, 2018, 84, .	1.4	32
60	Complete Genome Sequence of Sphingobium baderi DE-13, an Alkyl-Substituted Aniline-Mineralizing Bacterium. Current Microbiology, 2018, 75, 27-31.	1.0	4
61	Paenibacillus shunpengii sp. nov., isolated from farmland soil. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 211-216.	0.8	6
62	Identification and characterization of a novel carboxylesterase (FpbH) that hydrolyzes aryloxyphenoxypropionate herbicides. Biotechnology Letters, 2017, 39, 553-560.	1.1	18
63	Degradation of Diphenyl Ether in Sphingobium phenoxybenzoativorans SC_3 Is Initiated by a Novel Ring Cleavage Dioxygenase. Applied and Environmental Microbiology, 2017, 83, .	1.4	14
64	Biodegradation of Picolinic Acid by a Newly Isolated Bacterium Alcaligenes faecalis Strain JQ135. Current Microbiology, 2017, 74, 508-514.	1.0	28
65	Functional Characterization of a Novel Amidase Involved in Biotransformation of Triclocarban and its Dehalogenated Congeners in <i>Ochrobactrum</i> sp. TCC-2. Environmental Science & Environmental S	4.6	79
66	A Novel Aerobic Degradation Pathway for Thiobencarb Is Initiated by the TmoAB Two-Component Flavin Mononucleotide-Dependent Monooxygenase System in Acidovorax sp. Strain T1. Applied and Environmental Microbiology, 2017, 83, .	1.4	14
67	Molecular Mechanism and Genetic Determinants of Buprofezin Degradation. Applied and Environmental Microbiology, 2017, 83, .	1.4	14
68	Characterization and Genome Analysis of a Nicotine and Nicotinic Acid-Degrading Strain Pseudomonas putida JQ581 Isolated from Marine. Marine Drugs, 2017, 15, 156.	2.2	15
69	Characterization of a Novel Nicotine Degradation Gene Cluster ndp in Sphingomonas melonis TY and Its Evolutionary Analysis. Frontiers in Microbiology, 2017, 8, 337.	1.5	16
70	The Complete Genome Sequence of the Nicotine-Degrading Bacterium Shinella sp. HZN7. Frontiers in Microbiology, 2016, 7, 1348.	1.5	24
71	Isolation, identification, and acetochlor-degrading potential of a novel <i>Rhodococcus</i> sp. MZ-3. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2016, 51, 688-694.	0.7	4
72	Enantiomer signature and carbon isotope evidence for the migration and transformation of DDTs in arable soils across China. Scientific Reports, 2016, 6, 38475.	1.6	16

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73	Characterization of the novel dimethyl sulfide-degrading bacterium Alcaligenes sp . SY1 and its biochemical degradation pathway. Journal of Hazardous Materials, 2016, 304, 543-552.	6.5	23
74	Conversion of nornicotine to 6-hydroxy-nornicotine and 6-hydroxy-myosmine by Shinella sp. strain HZN7. Applied Microbiology and Biotechnology, 2016, 100, 10019-10029.	1.7	14
75	A Tetrahydrofolate-Dependent Methyltransferase Catalyzing the Demethylation of Dicamba in Sphingomonas sp. Strain Ndbn-20. Applied and Environmental Microbiology, 2016, 82, 5621-5630.	1.4	16
76	Co-exposure of silver nanoparticles and chiral herbicide imazethapyr to Arabidopsis thaliana: Enantioselective effects. Chemosphere, 2016, 145, 207-214.	4.2	47
77	Biodegradation of nicotine by a novel strain Pusillimonas. Research in Microbiology, 2015, 166, 67-71.	1.0	19
78	Regulators essential for nicotine degradation in Shinella sp. HZN7. Process Biochemistry, 2015, 50, 1947-1950.	1.8	10
79	Are Nutrient Stresses Associated with Enantioselectivity of the Chiral Herbicide Imazethapyr in <i>Arabidopsis thaliana</i> ?. Journal of Agricultural and Food Chemistry, 2015, 63, 10209-10217.	2.4	19
80	Isolation, transposon mutagenesis, and characterization of the novel nicotine-degrading strain Shinella sp. HZN7. Applied Microbiology and Biotechnology, 2014, 98, 2625-2636.	1.7	45
81	A Novel (<i>S</i>)-6-Hydroxynicotine Oxidase Gene from Shinella sp. Strain HZN7. Applied and Environmental Microbiology, 2014, 80, 5552-5560.	1.4	29
82	Biodegradation of beta-cypermethrin by a novel <i>Azoarcus indigens</i> strain HZ5. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2013, 48, 851-859.	0.7	19
83	Biodegradation of acetochlor by a newly isolated <i> Achromobacter </i> sp. strain D-12. Journal of Environmental Science and Health - Part B Pesticides, Food Contaminants, and Agricultural Wastes, 2013, 48, 960-966.	0.7	15
84	Cloning of a Novel Nicotine Oxidase Gene from Pseudomonas sp. Strain HZN6 Whose Product Nonenantioselectively Degrades Nicotine to Pseudooxynicotine. Applied and Environmental Microbiology, 2013, 79, 2164-2171.	1.4	31
85	Functional Identification of Two Novel Genes from Pseudomonas sp. Strain HZN6 Involved in the Catabolism of Nicotine. Applied and Environmental Microbiology, 2012, 78, 2154-2160.	1.4	49
86	A sirA-like gene, sirA2, is essential for 3-succinoyl-pyridine metabolism in the newly isolated nicotine-degrading Pseudomonas sp. HZN6 strain. Applied Microbiology and Biotechnology, 2011, 92, 1023-1032.	1.7	28
87	Hydrolytic Dechlorination of Chlorothalonil by Ochrobactrum sp. CTN-11 Isolated from a Chlorothalonil-Contaminated Soil. Current Microbiology, 2010, 61, 226-233.	1.0	39
88	Isolation, Identification and Characteristics of a Fluoroglycofen-ethyl-degrading Bacterium YF1*. Ying Yong Yu Huan Jing Sheng Wu Xue Bao = Chinese Journal of Applied and Environmental Biology, 2010, 2009, 686-691.	0.1	1
89	Isolation and characterization of three Sphingobium sp. strains capable of degrading isoproturon and cloning of the catechol 1,2-dioxygenase gene from these strains. World Journal of Microbiology and Biotechnology, 2009, 25, 259-268.	1.7	43