Cheng-Ying Jiang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7303662/publications.pdf Version: 2024-02-01



CHENC-YING LIANC

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Niabella beijingensis sp. nov. and Thermomonas beijingensis sp. nov., two bacteria from constructed wetland. International Journal of Systematic and Evolutionary Microbiology, 2022, 72, . | 1.7 | 12 |
| 2 | Alkaliphilus flagellatus sp. nov., Butyricicoccus intestinisimiae sp. nov., Clostridium mobile sp. nov., Clostridium simiarum sp. nov., Dysosmobacter acutus sp. nov., Paenibacillus brevis sp. nov., Peptoniphilus ovalis sp. nov. and Tissierella simiarum sp. nov., isolated from monkey faeces. International Journal of Systematic and Evolutionary Microbiology, 2022, 72, . | 1.7 | 34 |
| 3 | Alicyclobacillus curvatus sp. nov. and Alicyclobacillus mengziensis sp. nov., two acidophilic bacteria isolated from acid mine drainage. International Journal of Systematic and Evolutionary Microbiology, 2022, 72, . | 1.7 | 9 |
| 4 | PapA , a peptidoglycanâ€associated protein, interacts with OmpC and maintains cell envelope integrity. Environmental Microbiology, 2021, 23, 600-612. | 3.8 | 5 |
| 5 | Bacteria and Metabolic Potential in Karst Caves Revealed by Intensive Bacterial Cultivation and Genome Assembly. Applied and Environmental Microbiology, 2021, 87, . | 3.1 | 12 |
| 6 | Enlightening the taxonomy darkness of human gut microbiomes with a cultured biobank. Microbiome, 2021, 9, 119. | 11.1 | 479 |
| 7 | Submerged macrophytes recruit unique microbial communities and drive functional zonation in an aquatic system. Applied Microbiology and Biotechnology, 2021, 105, 7517-7528. | 3.6 | 9 |
| 8 | Responses of soil microbiome to steel corrosion. Npj Biofilms and Microbiomes, 2021, 7, 6. | 6.4 | 28 |
| 9 | Physiology, Taxonomy, and Sulfur Metabolism of the Sulfolobales, an Order of Thermoacidophilic Archaea. Frontiers in Microbiology, 2021, 12, 768283. | 3.5 | 4 |
| 10 | Key Factors Governing Microbial Community in Extremely Acidic Mine Drainage (pH <3). Frontiers in Microbiology, 2021, 12, 761579. | 3.5 | 12 |
| 11 | The Mouse Gut Microbial Biobank expands the coverage of cultured bacteria. Nature Communications, 2020, 11, 79. | 12.8 | 55 |
| 12 | Comparative Genomic Analysis Reveals the Metabolism and Evolution of the Thermophilic Archaeal Genus Metallosphaera. Frontiers in Microbiology, 2020, 11, 1192. | 3.5 | 8 |
| 13 | Diversity, Distribution and Co-occurrence Patterns of Bacterial Communities in a Karst Cave System. Frontiers in Microbiology, 2019, 10, 1726. | 3.5 | 80 |
| 14 | The ligandâ€binding domain of a chemoreceptor from <i>Comamonas testosteroni</i> has a previously unknown homotrimeric structure. Molecular Microbiology, 2019, 112, 906-917. | 2.5 | 13 |
| 15 | Cross Talk between Chemosensory Pathways That Modulate Chemotaxis and Biofilm Formation. MBio, 2019, 10, . | 4.1 | 49 |
| 16 | Vallitalea okinawensis sp. nov., isolated from Okinawa Trough sediment and emended description of the genus Vallitalea. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 404-410. | 1.7 | 17 |
| 17 | Crenobacter cavernae sp. nov., isolated from a karst cave, and emended description of the genus Crenobacter. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 476-480. | 1.7 | 11 |
| 18 | Cohnella faecalis sp. nov., isolated from animal faeces in a karst cave. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 572-577. | 1.7 | 13 |

CHENG-YING JIANG

| # | Article | IF | CITATIONS |
|----|---|------|-----------|
| 19 | Thauera hydrothermalis sp. nov., a thermophilic bacterium isolated from hot spring. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3163-3168. | 1.7 | 17 |
| 20 | Direct sensing and signal transduction during bacterial chemotaxis toward aromatic compounds in <i>Comamonas testosteroni</i> . Molecular Microbiology, 2016, 101, 224-237. | 2.5 | 34 |
| 21 | Reconstruction of metabolic networks in a fluoranthene-degrading enrichments from polycyclic aromatic hydrocarbon polluted soil. Journal of Hazardous Materials, 2016, 318, 90-98. | 12.4 | 44 |
| 22 | High-Throughput Single-Cell Cultivation on Microfluidic Streak Plates. Applied and Environmental Microbiology, 2016, 82, 2210-2218. | 3.1 | 136 |
| 23 | Ribosome binding site libraries and pathway modules for shikimic acid synthesis with Corynebacterium glutamicum. Microbial Cell Factories, 2015, 14, 71. | 4.0 | 78 |
| 24 | Unraveling the kinetic diversity of microbial 3-dehydroquinate dehydratases of shikimate pathway. AMB Express, 2015, 5, 7. | 3.0 | 5 |
| 25 | A novel chemoreceptor MCP2983 from Comamonas testosteroni specifically binds to cis-aconitate and triggers chemotaxis towards diverse organic compounds. Applied Microbiology and Biotechnology, 2015, 99, 2773-2781. | 3.6 | 25 |
| 26 | Metallosphaera tengchongensis sp. nov., an acidothermophilic archaeon isolated from a hot spring. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 537-542. | 1.7 | 19 |
| 27 | Alicyclobacillus fodiniaquatilis sp. nov., isolated from acid mine water. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 4915-4920. | 1.7 | 15 |
| 28 | Benzoate Metabolism Intermediate Benzoyl Coenzyme A Affects Gentisate Pathway Regulation in Comamonas testosteroni. Applied and Environmental Microbiology, 2014, 80, 4051-4062. | 3.1 | 27 |
| 29 | Thiosulfate Transfer Mediated by DsrE/TusA Homologs from Acidothermophilic Sulfur-oxidizing Archaeon Metallosphaera cuprina. Journal of Biological Chemistry, 2014, 289, 26949-26959. | 3.4 | 53 |
| 30 | Genetic characterization of 4-cresol catabolism in Corynebacterium glutamicum. Journal of Biotechnology, 2014, 192, 355-365. | 3.8 | 15 |
| 31 | Construction and application of an expression vector from the new plasmid pLAtc1 of Acidithiobacillus caldus. Applied Microbiology and Biotechnology, 2014, 98, 4083-4094. | 3.6 | 14 |
| 32 | Resolution of carbon metabolism and sulfur-oxidation pathways of Metallosphaera cuprina Ar-4 via comparative proteomics. Journal of Proteomics, 2014, 109, 276-289. | 2.4 | 30 |
| 33 | Parapedobacter pyrenivorans sp. nov., isolated from a pyrene-degrading microbial enrichment, and emended description of the genus Parapedobacter. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3994-3999. | 1.7 | 27 |
| 34 | Metallosphaera cuprina sp. nov., an acidothermophilic, metal-mobilizing archaeon. International Journal of Systematic and Evolutionary Microbiology, 2011, 61, 2395-2400. | 1.7 | 50 |
| 35 | Alicyclobacillus ferrooxydans sp. nov., a ferrous-oxidizing bacterium from solfataric soil. International Journal of Systematic and Evolutionary Microbiology, 2008, 58, 2898-2903. | 1.7 | 75 |
| 36 | Genome-wide investigation of aromatic acid transporters in Corynebacterium glutamicum. Microbiology (United Kingdom), 2007, 153, 857-865. | 1.8 | 63 |

CHENG-YING JIANG

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 37 | Micrococcus flavus sp. nov., isolated from activated sludge in a bioreactor. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 66-69. | 1.7 | 38 |
| 38 | Roseomonas lacus sp. nov., isolated from freshwater lake sediment. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 25-28. | 1.7 | 69 |
| 39 | Paracoccus sulfuroxidans sp. nov., a sulfur oxidizer from activated sludge. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 2693-2695. | 1.7 | 44 |
| 40 | Flavobacterium saliperosum sp. nov., isolated from freshwater lake sediment. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 439-442. | 1.7 | 54 |