Louis S Tisa

List of Publications by Year in descending order

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101384 149479 3,960 138 36 56 citations h-index g-index papers 140 140 140 2891 docs citations times ranked citing authors all docs

| # | Article | IF | CITATIONS |
|----|---|-------------|-----------|
| 1 | Genome characteristics of facultatively symbiotic Frankia sp. strains reflect host range and host plant biogeography. Genome Research, 2006, 17, 7-15. | 2.4 | 352 |
| 2 | Melanin Production and Use as a Soluble Electron Shuttle for Fe(III) Oxide Reduction and as a Terminal Electron Acceptor by Shewanella algae BrY. Applied and Environmental Microbiology, 2002, 68, 2436-2444. | 1.4 | 192 |
| 3 | Identification of the metalloregulatory element of the plasmid-encoded arsenical resistance operon. Nucleic Acids Research, 1990, 18, 619-624. | 6.5 | 137 |
| 4 | Friend or foe? A review of the mechanisms that drive <i>Serratia</i> towards diverse lifestyles. Canadian Journal of Microbiology, 2013, 59, 627-640. | 0.8 | 103 |
| 5 | Heavy Metal Resistance Patterns of Frankia Strains. Applied and Environmental Microbiology, 2002, 68, 923-927. | 1.4 | 97 |
| 6 | Significant Natural Product Biosynthetic Potential of Actinorhizal Symbionts of the Genus Frankia, as Revealed by Comparative Genomic and Proteomic Analyses. Applied and Environmental Microbiology, 2011, 77, 3617-3625. | 1.4 | 94 |
| 7 | Phylogenetic perspectives of nitrogen-fixing actinobacteria. Archives of Microbiology, 2012, 194, 3-11. | 1.0 | 92 |
| 8 | Cultivating the uncultured: growing the recalcitrant cluster-2 Frankia strains. Scientific Reports, 2015, 5, 13112. | 1.6 | 90 |
| 9 | The <i>Casuarina <scp>NIN</scp></i> gene is transcriptionally activated throughout <i>Frankia</i> root infection as well as in response to bacterial diffusible signals. New Phytologist, 2015, 208, 887-903. | 3.5 | 87 |
| 10 | Chronic N-amended soils exhibit an altered bacterial community structure in Harvard Forest, MA, USA. FEMS Microbiology Ecology, 2013, 83, 478-493. | 1.3 | 85 |
| 11 | Auxin Carriers Localization Drives Auxin Accumulation in Plant Cells Infected by <i>Frankia</i> in <i>Casuarina glauca</i> Actinorhizal Nodules. Plant Physiology, 2010, 154, 1372-1380. | 2.3 | 75 |
| 12 | Stone-dwelling actinobacteria <i>Blastococcus saxobsidens</i> , <i>Modestobacter marinus</i> and <i>Geodermatophilus obscurus</i> proteogenomes. ISME Journal, 2016, 10, 21-29. | 4.4 | 71 |
| 13 | Studies of growth and morphology of <i>Frankia</i> strains EAN1 _{pec} , Eul1 _c , Cpl1, and ACN1 ^{AG} . Canadian Journal of Botany, 1983, 61, 2768-2773. | 1.2 | 70 |
| 14 | Proposal of a type strain for Frankia alni (Woronin 1866) Von Tubeuf 1895, emended description of Frankia alni, and recognition of Frankia casuarinae sp. nov. and Frankia elaeagni sp. nov International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 5201-5210. | 0.8 | 68 |
| 15 | Exploring the genomes of Frankia. Physiologia Plantarum, 2007, 130, 331-343. | 2.6 | 62 |
| 16 | Phylogeny of members of the Frankia genus based on gyrB, nifH and glnII sequences. Antonie Van Leeuwenhoek, 2011, 100, 579-587. | 0.7 | 62 |
| 17 | Chitinaseâ€resistant hydrophilic symbiotic factors secreted by <i>Frankia</i> activate both Ca ²⁺ spiking and <i><scp>NIN</scp></i> gene expression in the actinorhizal plant <i>Casuarina glauca</i> New Phytologist, 2016, 209, 86-93. | 3. 5 | 62 |
| 18 | Electron transfer from Shewanella algae BrY to hydrous ferric oxide is mediated by cell-associated melanin. FEMS Microbiology Letters, 2003, 220, 99-104. | 0.7 | 60 |

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| 19 | Genomic approaches toward understanding the actinorhizal symbiosis: an update on the status ofÂthe Frankia genomes. Symbiosis, 2016, 70, 5-16. | 1.2 | 57 |
| 20 | Draft Genome Sequence of <i>Frankia</i> sp. Strain CN3, an Atypical, Noninfective (Nod [–]) Ineffective (Fix [–]) Isolate from <i>Coriaria nepalensis</i> . Genome Announcements, 2013, 1, e0008513. | 0.8 | 51 |
| 21 | Comparative secretome analysis suggests low plant cell wall degrading capacity in Frankia symbionts. BMC Genomics, 2008, 9, 47. | 1.2 | 49 |
| 22 | Frankia inefficax sp. nov., an actinobacterial endophyte inducing ineffective, non nitrogen-fixing, root nodules on its actinorhizal host plants. Antonie Van Leeuwenhoek, 2017, 110, 313-320. | 0.7 | 48 |
| 23 | Genomic, transcriptomic, and proteomic approaches towards understanding the molecular mechanisms of salt tolerance in Frankia strains isolated from Casuarina trees. BMC Genomics, 2017, 18, 633. | 1.2 | 46 |
| 24 | Actinorhizal Signaling Molecules: Frankia Root Hair Deforming Factor Shares Properties With NIN Inducing Factor. Frontiers in Plant Science, 2018, 9, 1494. | 1.7 | 46 |
| 25 | Genomic Insights Into Plant-Growth-Promoting Potentialities of the Genus Frankia. Frontiers in Microbiology, 2019, 10, 1457. | 1.5 | 46 |
| 26 | What stories can the Frankia genomes start to tell us?. Journal of Biosciences, 2013, 38, 719-726. | 0.5 | 44 |
| 27 | Casuarina Root Exudates Alter the Physiology, Surface Properties, and Plant Infectivity of Frankia sp. Strain Ccl3. Applied and Environmental Microbiology, 2012, 78, 575-580. | 1.4 | 43 |
| 28 | Symbiotic Performance of Diverse Frankia Strains on Salt-Stressed Casuarina glauca and Casuarina equisetifolia Plants. Frontiers in Plant Science, 2016, 7, 1331. | 1.7 | 43 |
| 29 | Tolerance to environmental stress by the nitrogen-fixing actinobacterium Frankia and its role in actinorhizal plants adaptation. Symbiosis, 2016, 70, 17-29. | 1.2 | 42 |
| 30 | Transport systems encoded by bacterial plasmids. Journal of Bioenergetics and Biomembranes, 1990, 22, 493-507. | 1.0 | 41 |
| 31 | Effects of Organic Antagonists of Ca2+, Na+, and K+ on Chemotaxis and Motility ofEscherichia coli. Journal of Bacteriology, 2000, 182, 4856-4861. | 1.0 | 40 |
| 32 | Draft Genome Sequence of <i>Frankia</i> sp. Strain BCU110501, a Nitrogen-Fixing Actinobacterium Isolated from Nodules of <i>Discaria trinevis</i> . Genome Announcements, 2013, 1, . | 0.8 | 40 |
| 33 | Contrasted evolutionary constraints on secreted and non-secreted proteomes of selected Actinobacteria. BMC Genomics, 2013, 14, 474. | 1.2 | 39 |
| 34 | Draft Genome Sequence of <i>Frankia</i> sp. Strain QA3, a Nitrogen-Fixing Actinobacterium Isolated from the Root Nodule of <i>Alnus nitida</i> Genome Announcements, 2013, 1, e0010313. | 0.8 | 39 |
| 35 | Draft Genome Sequence of <i>Frankia</i> sp. Strain BMG5.12, a Nitrogen-Fixing Actinobacterium Isolated from Tunisian Soils. Genome Announcements, 2013, 1 , . | 0.8 | 39 |
| 36 | Elucidating the ecological networks in stoneâ€dwelling microbiomes. Environmental Microbiology, 2020, 22, 1467-1480. | 1.8 | 38 |

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| 37 | The implication of life style on codon usage patterns and predicted highly expressed genes for three Frankia genomes. Antonie Van Leeuwenhoek, 2008, 93, 335-346. | 0.7 | 37 |
| 38 | Photorhabdus heterorhabditis subsp. aluminescens subsp. nov., Photorhabdus heterorhabditis subsp. heterorhabditis subsp. nov., Photorhabdus australis subsp. thailandensis subsp. nov., Photorhabdus australis subsp. australis subsp. nov., and Photorhabdus aegyptia sp. nov. isolated from Heterorhabditis entomopathogenic nematodes. International Journal of Systematic and Evolutionary Microbiology, 2021, 71,. | 0.8 | 37 |
| 39 | Draft Genome Sequence of <i>Frankia</i> sp. Strain Ccl6, a Salt-Tolerant Nitrogen-Fixing Actinobacterium Isolated from the Root Nodule of <i>Casuarina cunninghamiana</i> Genome Announcements, 2014, 2, . | 0.8 | 36 |
| 40 | Draft genome sequence of the symbiotic Frankia sp. strain BMG5.30 isolated from root nodules of Coriaria myrtifolia in Tunisia. Antonie Van Leeuwenhoek, 2019, 112, 67-74. | 0.7 | 35 |
| 41 | Nitric Oxide and Oxygen Regulate Truncated Hemoglobin Gene Expression in (i>Frankia (/i>Strain Ccl3. Journal of Bacteriology, 2008, 190, 7864-7867. | 1.0 | 34 |
| 42 | Insertion sequence content reflects genome plasticity in strains of the root nodule actinobacterium Frankia. BMC Genomics, 2009, 10, 468. | 1.2 | 34 |
| 43 | Soil bacterial communities of a calcium-supplemented and a reference watershed at the Hubbard Brook Experimental Forest (HBEF), New Hampshire, USA. FEMS Microbiology Ecology, 2012, 79, 728-740. | 1.3 | 34 |
| 44 | Draft Genome Sequence of <i>Frankia</i> sp. Strain Thr, a Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Casuarina cunninghamiana</i> Grown in Egypt. Genome Announcements, 2014, 2, . | 0.8 | 33 |
| 45 | Draft Genome Sequence of <i>Frankia</i> sp. Strain BMG5.23, a Salt-Tolerant Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Casuarina glauca</i> Grown in Tunisia. Genome Announcements, 2014, 2, . | 0.8 | 33 |
| 46 | Frankia discariae sp. nov.: an infective and effective microsymbiont isolated from the root nodule of Discaria trinervis. Archives of Microbiology, 2017, 199, 641-647. | 1.0 | 33 |
| 47 | Pyomelanin is produced by Shewanella algae BrY and affected by exogenous iron. Canadian Journal of Microbiology, 2008, 54, 334-339. | 0.8 | 30 |
| 48 | Copper tolerance in Frankia sp. strain Eullc involves surface binding and copper transport. Applied Microbiology and Biotechnology, 2014, 98, 8005-8015. | 1.7 | 29 |
| 49 | Frankia torreyi sp. nov., the first actinobacterium of the genus Frankia Brunchorst 1886, 174AL isolated in axenic culture. Antonie Van Leeuwenhoek, 2019, 112, 57-65. | 0.7 | 29 |
| 50 | An update on the taxonomy of the genus Frankia Brunchorst, 1886, 174AL. Antonie Van Leeuwenhoek, 2019, 112, 5-21. | 0.7 | 29 |
| 51 | Influence of Temperature on the Physiology and Virulence of the Insect Pathogen Serratia sp. Strain SCBI. Applied and Environmental Microbiology, 2012, 78, 8840-8844. | 1.4 | 28 |
| 52 | Genome sequence and comparative analysis of a putative entomopathogenic Serratia isolated from Caenorhabditis briggsae. BMC Genomics, 2015, 16, 531. | 1.2 | 27 |
| 53 | The ins and outs of metal homeostasis by the root nodule actinobacterium Frankia. BMC Genomics, 2014, 15, 1092. | 1.2 | 26 |
| 54 | Antibiotic resistance patterns of <i>Frankia</i> strains. Canadian Journal of Botany, 1999, 77, 1257-1260. | 1,2 | 26 |

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| 55 | Nitrogen Fixation Mutants of the Actinobacterium <i>Frankia Casuarinae</i> Ccl3. Microbes and Environments, 2017, 32, 344-351. | 0.7 | 25 |
| 56 | Nocardia casuarinae sp. nov., an actinobacterial endophyte isolated from root nodules of Casuarina glauca. Antonie Van Leeuwenhoek, 2014, 105, 1099-1106. | 0.7 | 24 |
| 57 | The plant-growth-promoting actinobacteria of the genus Nocardia induces root nodule formation in Casuarina glauca. Antonie Van Leeuwenhoek, 2019, 112, 75-90. | 0.7 | 24 |
| 58 | Draft Genome Sequence of <i>Frankia</i> sp. Strain DC12, an Atypical, Noninfective, Ineffective Isolate from <i>Datisca cannabina</i> . Genome Announcements, 2015, 3, . | 0.8 | 23 |
| 59 | Molecular Characterization of Protease Activity in Serratia sp. Strain SCBI and Its Importance in Cytotoxicity and Virulence. Journal of Bacteriology, 2014, 196, 3923-3936. | 1.0 | 22 |
| 60 | Permanent Draft Genome Sequence for <i>Frankia</i> sp. Strain CeD, a Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Casuarina equistifolia</i> Grown in Senegal. Genome Announcements, 2016, 4, . | 0.8 | 22 |
| 61 | Formation and Regeneration of Protoplasts of the Actinorhizal Nitrogen-Fixing Actinomycete <i>Frankia</i> . Applied and Environmental Microbiology, 1987, 53, 53-56. | 1.4 | 20 |
| 62 | Hemoglobin in five genetically diverseFrankiastrains. Canadian Journal of Microbiology, 2002, 48, 1048-1055. | 0.8 | 18 |
| 63 | On the nature of fur evolution: A phylogenetic approach in Actinobacteria. BMC Evolutionary Biology, 2008, 8, 185. | 3.2 | 18 |
| 64 | Stable Transformation of the Actinobacteria $\mbox{\sc i>Frankia} \mbox{\sc /i> spp.}$ Applied and Environmental Microbiology, 2019, 85, . | 1.4 | 18 |
| 65 | Germination and physiological properties of Frankia spores. Plant and Soil, 2003, 254, 57-67. | 1.8 | 17 |
| 66 | Effect of salt stress on the physiology of Frankia sp strain Ccl6. Journal of Biosciences, 2013, 38, 699-702. | 0.5 | 17 |
| 67 | Diversity of Frankia Strains, Actinobacterial Symbionts of Actinorhizal Plants. Soil Biology, 2013, , 123-148. | 0.6 | 17 |
| 68 | Permanent draft genome sequence of Frankia sp. NRRL B-16219 reveals the presence of canonical nod genes, which are highly homologous to those detected in Candidatus Frankia Dg1 genome. Standards in Genomic Sciences, 2017, 12, 51. | 1.5 | 17 |
| 69 | Detoxification and reduction of selenite to elemental red selenium by Frankia. Antonie Van Leeuwenhoek, 2019, 112, 127-139. | 0.7 | 17 |
| 70 | Molecular responses of (i>Frankia (i>sp. strain QA3 to naphthalene. Canadian Journal of Microbiology, 2015, 61, 281-292. | 0.8 | 16 |
| 71 | Permanent Draft Genome Sequences for Two Variants of <i>Frankia</i> sp. Strain Cpl1, the First <i>Frankia</i> Strain Isolated from Root Nodules of <i>Comptonia peregrina</i> Genome Announcements, 2016, 4, . | 0.8 | 16 |
| 72 | Pb2+ tolerance by Frankia sp. strain EAN1pec involves surface-binding. Microbiology (United Kingdom), 2017, 163, 472-487. | 0.7 | 16 |

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| 73 | A plasmid-encoded anion-translocating ATPase. Biochimica Et Biophysica Acta - Bioenergetics, 1990, 1018, 203-205. | 0.5 | 15 |
| 74 | Metagenome Across a Geochemical Gradient of Indian Stone Ruins Found at Historic Sites in Tamil Nadu, India. Microbial Ecology, 2021, 81, 385-395. | 1.4 | 15 |
| 75 | Swarming motility by Photorhabdus temperata is influenced by environmental conditions and uses the same flagella as that used in swimming motilityThis is scientific contribution No. 2431 from the New Hampshire Agricultural Experiment Station Canadian Journal of Microbiology, 2011, 57, 196-203. | 0.8 | 13 |
| 76 | Draft Genome Sequence of <i>Frankia</i> Strain G2, a Nitrogen-Fixing Actinobacterium Isolated from <i>Casuarina equisetifolia</i> and Able To Nodulate Actinorhizal Plants of the Order <i>Rhamnales</i> . Genome Announcements, 2016, 4, . | 0.8 | 13 |
| 77 | Ecology and Physiology of Non-Frankia Actinobacteria from Actinorhizal Plants. , 2014, , 27-42. | | 13 |
| 78 | Antibiotic resistance patterns of <i>Frankia </i> strains. Canadian Journal of Botany, 1999, 77, 1257-1260. | 1.2 | 12 |
| 79 | Effect of electroporation conditions on cell viability of Frankia Eul1c. Plant and Soil, 2003, 254, 83-88. | 1.8 | 12 |
| 80 | Alteration of the exopolysaccharide production and the transcriptional profile of free-living Frankia strain Ccl3 under nitrogen-fixing conditions. Applied Microbiology and Biotechnology, 2013, 97, 10499-10509. | 1.7 | 11 |
| 81 | Permanent Draft Genome Sequence of <i>Frankia</i> sp. Strain ACN1 ^{ag} , a Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Alnus glutinosa</i> . Genome Announcements, 2015, 3, . | 0.8 | 11 |
| 82 | Permanent Draft Genome Sequence of <i>Frankia</i> sp <i>.</i> Strain Allo2, a Salt-Tolerant Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Allocasuarina</i> Genome Announcements, 2016, 4, . | 0.8 | 11 |
| 83 | Effect of growth conditions on the motility of Photorhabdus temperata. Archives of Microbiology, 2003, 180, 17-24. | 1.0 | 10 |
| 84 | Isolation of antibiotic-resistant and antimetabolite-resistant mutants of Frankiastrains Eullc and Cc1.17. Canadian Journal of Microbiology, 2004, 50, 261-267. | 0.8 | 10 |
| 85 | Homology modelling of the Frankia nitrogenase iron protein. Symbiosis, 2010, 50, 37-44. | 1.2 | 10 |
| 86 | Elucidation of the Photorhabdus temperata Genome and Generation of a Transposon Mutant Library To Identify Motility Mutants Altered in Pathogenesis. Journal of Bacteriology, 2015, 197, 2201-2216. | 1.0 | 10 |
| 87 | The Family Frankiaceae. , 2014, , 339-356. | | 10 |
| 88 | Alone Yet Not Alone: Frankia Lives Under the Same Roof With Other Bacteria in Actinorhizal Nodules. Frontiers in Microbiology, 2021, 12, 749760. | 1.5 | 10 |
| 89 | Development of a semi-high-throughput growth assay for the filamentous actinobacteria Frankia. Archives of Microbiology, 2012, 194, 13-20. | 1.0 | 9 |
| 90 | Permanent Draft Genome Sequence of <i>Frankia</i> sp. Strain Avcl1, a Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Alnus viridis</i> subsp. <i>crispa</i> Canada. Genome Announcements, 2015, 3, . | 0.8 | 9 |

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| 91 | Bioinformatic Analysis of Codon Usage Patterns in a Free Living Diazotroph, Azotobacter vinelandii. Biotechnology, 2008, 7, 242-249. | 0.5 | 9 |
| 92 | Identification of TTA codon containing genes in Frankia and exploration of the role of tRNA in regulating these genes. Archives of Microbiology, 2012, 194, 35-45. | 1.0 | 8 |
| 93 | Draft Genome Sequence of the Symbiotic <i>Frankia</i> Sp. Strain KB5 Isolated from Root Nodules of <i>Casuarina equisetifolia</i> Journal of Genomics, 2017, 5, 64-67. | 0.6 | 8 |
| 94 | Simple colony PCR procedure for the filamentous actinobacteria Frankia. Antonie Van Leeuwenhoek, 2019, 112, 109-114. | 0.7 | 8 |
| 95 | Development of a physical map for three Frankia strains and a partial genetic map for Frankia Eul1c. Physiologia Plantarum, 2007, 130, 427-439. | 2.6 | 7 |
| 96 | An update on research on Frankia and actinorhizal plants on the occasion of the 18th meeting of the Frankia-actinorhizal plants symbiosis. Symbiosis, 2016, 70, 1-4. | 1.2 | 7 |
| 97 | Comparative genomics of Prauserella sp. Am3, an actinobacterium isolated from root nodules of Alnus nepalensis in India. Symbiosis, 2016, 70, 49-58. | 1.2 | 7 |
| 98 | Permanent Draft Genome Sequence of <i>Rhizobium</i> sp. Strain LCM 4573, a Salt-Tolerant, Nitrogen-Fixing Bacterium Isolated from Senegalese Soils. Genome Announcements, 2017, 5, . | 0.8 | 7 |
| 99 | A novel phylogenetic tree based on the presence of protein domains in selected actinobacteria. Antonie Van Leeuwenhoek, 2019, 112, 101-107. | 0.7 | 7 |
| 100 | Permanent Draft Genome Sequence of <i>Frankia</i> sp. Strain BR, a Nitrogen-Fixing Actinobacterium Isolated from the Root Nodules of <i>Casuarina equisetifolia</i> . Genome Announcements, 2016, 4, . | 0.8 | 6 |
| 101 | Permanent Draft Genome sequence for Frankia sp. strain Ccl49, a Nitrogen-Fixing Bacterium Isolated from Casuarina cunninghamiana that Infects Elaeagnaceae. Journal of Genomics, 2017, 5, 119-123. | 0.6 | 6 |
| 102 | Biosynthetic energy cost of potentially highly expressed proteins vary with niche in selected actinobacteria. Journal of Basic Microbiology, 2018, 58, 154-161. | 1.8 | 6 |
| 103 | Calcium Transport by Frankia sp. Strain EAN1pec. Current Microbiology, 1998, 37, 12-16. | 1.0 | 5 |
| 104 | Characterization of pseudogenes in members of the order Frankineae. Journal of Biosciences, 2013, 38, 727-732. | 0.5 | 5 |
| 105 | Draft Genome Sequence of Photorhabdus luminescens Strain BA1, an Entomopathogenic Bacterium Isolated from Nematodes Found in Egypt. Genome Announcements, 2014, 2, . | 0.8 | 5 |
| 106 | Permanent Improved High-Quality Draft Genome Sequence of Nocardia casuarinae Strain BMG51109, an Endophyte of Actinorhizal Root Nodules of Casuarina glauca. Genome Announcements, 2016, 4, . | 0.8 | 5 |
| 107 | Molecular Methods for Research on Actinorhiza. Rhizosphere Biology, 2019, , 35-59. | 0.4 | 5 |
| 108 | Contrasted evolutionary constraints on carbohydrate active enzymes (CAZymes) in selected Frankia strains. Antonie Van Leeuwenhoek, 2019, 112, 115-125. | 0.7 | 5 |

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| 109 | Germination and physiological properties of Frankia spores. , 2003, , 57-67. | | 5 |
| 110 | Characterization of haemoglobin from Actinorhizal plants – An in silico approach. Journal of Biosciences, 2013, 38, 777-787. | 0.5 | 4 |
| 111 | Contrasted Reactivity to Oxygen Tensions inFrankiasp. Strain Ccl3 throughout Nitrogen Fixation and Assimilation. BioMed Research International, 2014, 2014, 1-8. | 0.9 | 4 |
| 112 | Draft Genome Sequence of Photorhabdus temperata Strain Meg1, an Entomopathogenic Bacterium Isolated from Heterorhabditis megidis Nematodes. Genome Announcements, 2014, 2, . | 0.8 | 4 |
| 113 | Frankia as a Biodegrading Agent. , 2016, , . | | 4 |
| 114 | Characterization of PAS domains in Frankia and selected Actinobacteria and their possible interaction with other co-domains for environmental adaptation. Symbiosis, 2016, 70, 69-78. | 1.2 | 4 |
| 115 | Permanent Draft Genome Sequence for <i>Frankia</i> sp <i>.</i> Strain El5c, a Single-Spore Isolate of a Nitrogen-Fixing Actinobacterium, Isolated from the Root Nodules of <i>Elaeagnus angustifolia</i> Genome Announcements, 2016, 4, . | 0.8 | 4 |
| 116 | Permanent Draft Genome Sequences of Three <i>Frankia</i> sp. Strains That Are Atypical, Noninfective, Ineffective Isolates. Genome Announcements, 2017, 5, . | 0.8 | 4 |
| 117 | Draft Genome Sequences of 10 Bacterial Strains Isolated from Root Nodules of Alnus Trees in New Hampshire. Microbiology Resource Announcements, 2020, 9, . | 0.3 | 4 |
| 118 | In silico Analysis of Chlorobium Genomes Divulge Insights into the Lifestyle of the Bacteria. Research Journal of Microbiology, 2008, 3, 600-613. | 0.2 | 4 |
| 119 | Permanent Draft Genome Sequence of Nocardia sp. BMG111209, an Actinobacterium Isolated from Nodules of Casuarina glauca. Genome Announcements, 2016, 4, . | 0.8 | 3 |
| 120 | Permanent Draft Genome Sequence for Frankia sp. Strain Cc1.17, a Nitrogen-Fixing Actinobacterium Isolated from Root Nodules of Colletia cruciata. Genome Announcements, 2017, 5, . | 0.8 | 3 |
| 121 | Advances in Frankia genome studies and molecular aspects of tolerance to environmental stresses. , 2020, , 381-389. | | 3 |
| 122 | Draft Genome Sequence of the Symbiotic <i>Frankia</i> sp. strain B2 isolated from root nodules of <i>Casuarina cunninghamiana</i> found in Algeria. Journal of Genomics, 2020, 8, 11-15. | 0.6 | 3 |
| 123 | Construction and purification of His-tagged staphylococcal ArsB protein, an integral membrane protein that is involved in arsenical salt resistance. Indian Journal of Microbiology, 2009, 49, 212-218. | 1.5 | 2 |
| 124 | Microbial Processes in Fractured Rock Environments. Geophysical Monograph Series, 2013, , 183-193. | 0.1 | 2 |
| 125 | Draft Genome Sequence of <i>Photorhabdus luminescens</i> subsp. <i>laumondii</i> HP88, an Entomopathogenic Bacterium Isolated from Nematodes. Genome Announcements, 2016, 4, . | 0.8 | 2 |
| 126 | Permanent Draft Genome Sequence of the French Bean Symbiont Rhizobium sp. Strain RSm-3 Isolated from the Eastern Himalayan Region of India. Genome Announcements, 2017, 5, . | 0.8 | 2 |

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| 127 | Draft Genome Sequences for the Frankia sp. strains CgS1, Ccl156 and CgMl4, Nitrogen-Fixing Bacteria Isolated from Casuarina sp. in Egypt. Journal of Genomics, 2020, 8, 84-88. | 0.6 | 2 |
| 128 | Draft Genome Sequences of 6 Actinobacterial Strains Isolated from Rock Surfaces Obtained from Indian Stone Ruins in Tamil Nadu, India, and Rocks from New England, United States. Microbiology Resource Announcements, 2022, 11, e0002422. | 0.3 | 2 |
| 129 | Interaction of ${\rm \tilde{A}\hat{A}}$ %-conotoxin and the membrane calcium transport system of Escherichia coli. FEMS Microbiology Letters, 2000, 188, 97-101. | 0.7 | 1 |
| 130 | Microfracture Surface Geochemistry and Adherent Microbial Population Metabolism in TCE-Contaminated Competent Bedrock. Geomicrobiology Journal, 2007, 24, 307-330. | 1.0 | 1 |
| 131 | Permanent Draft Genome Sequence of Photorhabdus temperata Strain Hm, an Entomopathogenic Bacterium Isolated from Nematodes. Genome Announcements, 2017, 5, . | 0.8 | 1 |
| 132 | Inactivation of the Major Hemolysin Gene Influences Expression of the Nonribosomal Peptide Synthetase Gene swrA in the Insect Pathogen Serratia sp. Strain SCBI. Journal of Bacteriology, 2017, 199, . | 1.0 | 1 |
| 133 | Permanent Draft Genome Sequence of <i>Ensifer</i> sp. Strain LCM 4579, a Salt-Tolerant, Nitrogen-Fixing Bacterium Isolated from Senegalese Soil. Genome Announcements, 2017, 5, . | 0.8 | 1 |
| 134 | Draft Genome Sequence for Frankia sp. Strain BMG5.11, a Nitrogen-Fixing Bacterium Isolated from Elaeagnus angustifolia. Microbiology Resource Announcements, 2020, 9, . | 0.3 | 1 |
| 135 | Identification and characterization of Photorhabdus temperata mutants altered in hemolysis and virulence. Canadian Journal of Microbiology, 2016, 62, 657-667. | 0.8 | 0 |
| 136 | Permanent Draft Genome Sequences for Mesorhizobium sp. Strains LCM 4576, LCM 4577, and ORS3428, Salt-Tolerant, Nitrogen-Fixing Bacteria Isolated from Senegalese Soils. Genome Announcements, 2017, 5, . | 0.8 | 0 |
| 137 | Quantitative Analysis of Gene Expression During Calcium Homeostasis in E. coli. Bangladesh Journal of Microbiology, 2019, 34, 47-54. | 0.2 | 0 |
| 138 | Calcium Homeostasis in Escherichia coli: Characterization of Mutants and Genome Expression of MG1655. Bangladesh Journal of Microbiology, 0, , 1-8. | 0.2 | 0 |