

# Yonghui Zeng

## List of Publications by Year in descending order

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Version: 2024-02-01

58  
papers

1,832  
citations

304743

22  
h-index

276875

41  
g-index

61  
all docs

61  
docs citations

61  
times ranked

2051  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Functional type 2 photosynthetic reaction centers found in the rare bacterial phylum Gemmatimonadetes. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 7795-7800.  | 7.1  | 220       |
| 2  | Distinct distribution pattern of abundance and diversity of aerobic anoxygenic phototrophic bacteria in the global ocean. Environmental Microbiology, 2007, 9, 3091-3099.  | 3.8  | 164       |
| 3  | Ecological anomalies in the East China Sea: Impacts of the Three Gorges Dam?. Water Research, 2007, 41, 1287-1293.   | 11.3 | 138       |
| 4  | Bacterial diversity in the snow over Tibetan Plateau Glaciers. Extremophiles, 2009, 13, 411-423.   | 2.3  | 114       |
| 5  | Characterization of the microaerophilic, bacteriochlorophyll a-containing bacterium Gemmatimonas phototrophica sp. nov., and emended descriptions of the genus Gemmatimonas and Gemmatimonas aurantiaca. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 2410-2419. | 1.7  | 98        |
| 6  | Microbial community structure in moraine lakes and glacial meltwaters, Mount Everest. FEMS Microbiology Letters, 2006, 265, 98-105.  | 1.8  | 72        |
| 7  | Diversity and distribution of pigmented heterotrophic bacteria in marine environments. FEMS Microbiology Ecology, 2006, 57, 92-105.  | 2.7  | 68        |
| 8  | Metagenomic evidence for the presence of phototrophic <i>Gemmatimonadetes</i> bacteria in diverse environments. Environmental Microbiology Reports, 2016, 8, 139-149.  | 2.4  | 66        |
| 9  | Aerobic Anoxygenic Photosynthesis Is Commonly Present within the Genus <i>Limnohabitans</i> . Applied and Environmental Microbiology, 2018, 84, .  | 3.1  | 64        |
| 10 | Bacterial Diversity of Freshwater Alpine Lake Puma Yumco on the Tibetan Plateau. Geomicrobiology Journal, 2009, 26, 131-145.   | 2.0  | 55        |
| 11 | Genome Sequences of Two Freshwater Betaproteobacterial Isolates, <i>Limnohabitans</i> Species Strains Rim28 and Rim47, Indicate Their Capabilities as Both Photoautotrophs and Ammonia Oxidizers. Journal of Bacteriology, 2012, 194, 6302-6303.   | 2.2  | 48        |
| 12 | <i>Gemmatimonas groenlandica</i> sp. nov. Is an Aerobic Anoxygenic Phototroph in the Phylum Gemmatimonadetes. Frontiers in Microbiology, 2020, 11, 606612.   | 3.5  | 48        |
| 13 | Real-time PCR for quantification of aerobic anoxygenic phototrophic bacteria based on <i>pufM</i> gene in marine environment. Journal of Experimental Marine Biology and Ecology, 2006, 329, 113-121.  | 1.5  | 36        |
| 14 | Bacteria variabilities in a Tibetan ice core and their relations with climate change. Global Biogeochemical Cycles, 2008, 22, .  | 4.9  | 34        |
| 15 | Abundant presence of the $\beta$ -like Proteobacterial <i>pufM</i> gene in oxic seawater. FEMS Microbiology Letters, 2006, 263, 200-206.   | 1.8  | 32        |
| 16 | Bacterial Community of the Largest Oligosaline Lake, Namco on the Tibetan Plateau. Geomicrobiology Journal, 2010, 27, 669-682.   | 2.0  | 32        |
| 17 | <i>Mameliella alba</i> gen. nov., sp. nov., a marine bacterium of the Roseobacter clade in the order Rhodobacterales. International Journal of Systematic and Evolutionary Microbiology, 2010, 60, 953-957.  | 1.7  | 31        |
| 18 | Phylogenetic diversity of planktonic archaea in the estuarine region of East China Sea. Microbiological Research, 2007, 162, 26-36.  | 5.3  | 29        |

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|----|---|------|-----------|
| 19 | Seasonal variation of snow microbial community structure in the East Rongbuk glacier, Mt. Everest. <i>Science Bulletin</i> , 2006, 51, 1476-1486.   | 9.0  | 28        |
| 20 | Genomic Analysis of the Evolution of Phototrophy among Haloalkaliphilic Rhodobacterales. <i>Genome Biology and Evolution</i> , 2017, 9, 1950-1962.  | 2.5  | 25        |
| 21 | Genetic diversity assessment of anoxygenic photosynthetic bacteria by distance-based grouping analysis of <i>pufM</i> sequences. <i>Letters in Applied Microbiology</i> , 2007, 45, 639-645.                                      | 2.2  | 24        |
| 22 | Bacterial diversity in various coastal mariculture ponds in Southeast China and in diseased eels as revealed by culture and culture-independent molecular techniques. <i>Aquaculture Research</i> , 2010, 41, e172-e186.          | 1.8  | 24        |
| 23 | Microbial community structure in major habitats above 6000 m on Mount Everest. <i>Science Bulletin</i> , 2007, 52, 2350-2357.   | 1.7  | 23        |
| 24 | Potential Rhodopsin- and Bacteriochlorophyll-Based Dual Phototrophy in a High Arctic Glacier. <i>MBio</i> , 2020, 11, .   | 4.1  | 23        |
| 25 | Natural community structure of cyanobacteria in the South China Sea as revealed by <i>rpoC1</i> gene sequence analysis. <i>Letters in Applied Microbiology</i> , 2004, 39, 353-358.   | 2.2  | 21        |
| 26 | Genomics of Aerobic Photoheterotrophs in Wheat Phyllosphere Reveals Divergent Evolutionary Patterns of Photosynthetic Genes in <i>Methylobacterium</i> spp.. <i>Genome Biology and Evolution</i> , 2019, 11, 2895-2908.           | 2.5  | 19        |
| 27 | Vertical distribution and phylogenetic composition of bacteria in the Eastern Tropical North Pacific Ocean. <i>Microbiological Research</i> , 2009, 164, 624-633.   | 5.3  | 18        |
| 28 | Regressive Evolution of Photosynthesis in the Roseobacter Clade. <i>Advances in Botanical Research</i> , 2013, 66, 385-405.   | 1.1  | 18        |
| 29 | Long PCR-RFLP of 16S-ITS-23S rRNA genes: a high-resolution molecular tool for bacterial genotyping. <i>Journal of Applied Microbiology</i> , 2013, 114, 433-447.  | 3.1  | 18        |
| 30 | Novel <i>acsF</i> Gene Primers Revealed a Diverse Phototrophic Bacterial Population, Including Gemmatimonadetes, in Lake Taihu (China). <i>Applied and Environmental Microbiology</i> , 2016, 82, 5587-5594.                      | 3.1  | 18        |
| 31 | <i>Sandarakinorhabdus cyanobacteriorum</i> sp. nov., a novel bacterium isolated from cyanobacterial aggregates in a eutrophic lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 730-735. | 1.7  | 18        |
| 32 | Aerobic Anoxygenic Phototrophic Bacteria Promote the Development of Biological Soil Crusts. <i>Frontiers in Microbiology</i> , 2018, 9, 2715.   | 3.5  | 17        |
| 33 | <i>Elstera cyanobacteriorum</i> sp. nov., a novel bacterium isolated from cyanobacterial aggregates in a eutrophic lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2017, 67, 4272-4275.          | 1.7  | 17        |
| 34 | Bacteria in the lakes of the Tibetan Plateau and polar regions. <i>Science of the Total Environment</i> , 2021, 754, 142248.  | 8.0  | 16        |
| 35 | 2.4-Å... structure of the double-ring <i>Gemmatimonas phototrophica</i> photosystem. <i>Science Advances</i> , 2022, 8, eabk3139.   | 10.3 | 16        |
| 36 | Genomic Insights of <i>Cryobacterium</i> Isolated From Ice Core Reveal Genome Dynamics for Adaptation in Glacier. <i>Frontiers in Microbiology</i> , 2020, 11, 1530.  | 3.5  | 12        |

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|----|--|-----|-----------|
| 37 | Genetic diversity of aerobic anoxygenic photosynthetic bacteria in open ocean surface waters and upper twilight zones. <i>Marine Biology</i> , 2009, 156, 425-437.   | 1.5 | 11        |
| 38 | Whole-Genome Sequences of an Aerobic Anoxygenic Phototroph, <i>Blastomonas</i> sp. Strain AAP53, Isolated from a Freshwater Desert Lake in Inner Mongolia, China. <i>Genome Announcements</i> , 2013, 1, e0007113.                               | 0.8 | 11        |
| 39 | Phototrophic Gemmatimonadetes: A New "Purple" Branch on the Bacterial Tree of Life. , 2017, , 163-192.   |     | 11        |
| 40 | Simultaneous Presence of Bacteriochlorophyll and Xanthorhodopsin Genes in a Freshwater Bacterium. <i>MSystems</i> , 2020, 5, .   | 3.8 | 11        |
| 41 | Development and evaluation of specific 16S rDNA primers for marine Cytophaga-Flavobacteria cluster. <i>Molecular Ecology Notes</i> , 2006, 6, 1278-1281.   | 1.7 | 10        |
| 42 | Draft Genome Sequence of the Cellulolytic Bacterium <i>Clavibacter</i> sp. CF11, a Strain Producing Cold-Active Cellulase. <i>Genome Announcements</i> , 2015, 3, .  | 0.8 | 10        |
| 43 | Characterization of the Aerobic Anoxygenic Phototrophic Bacterium <i>Sphingomonas</i> sp. AAP5. <i>Microorganisms</i> , 2021, 9, 768.  | 3.6 | 10        |
| 44 | <i>Niveispirillum lacus</i> sp. nov., isolated from cyanobacterial aggregates in a eutrophic lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 507-512.   | 1.7 | 10        |
| 45 | Comparison of bacterioplankton communities in three mariculture ponds farming different commercial animals in subtropical Chinese coast. <i>Hydrobiologia</i> , 2009, 632, 107-126.  | 2.0 | 8         |
| 46 | <i>Flavobacterium cyanobacteriorum</i> sp. nov., isolated from cyanobacterial aggregates in a eutrophic lake. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1279-1284.                                    | 1.7 | 8         |
| 47 | Source environment feature related phylogenetic distribution pattern of anoxygenic photosynthetic bacteria as revealed by <i>pufM</i> analysis. <i>Journal of Microbiology</i> , 2007, 45, 205-12.   | 2.8 | 6         |
| 48 | Abundance and diversity of snow bacteria in two glaciers at the Tibetan Plateau. <i>Frontiers of Earth Science</i> , 2009, 3, 80-90.   | 0.5 | 5         |
| 49 | Whole-Genome Sequence of a Freshwater Aerobic Anoxygenic Phototroph, <i>Porphyrobacter</i> sp. Strain AAP82, Isolated from the Huguangyan Maar Lake in Southern China. <i>Genome Announcements</i> , 2013, 1, e0007213.                          | 0.8 | 5         |
| 50 | Genome Sequences and Photosynthesis Gene Cluster Composition of a Freshwater Aerobic Anoxygenic Phototroph, <i>Sandarakinorhabdus</i> sp. Strain AAP62, Isolated from the Shahu Lake in Ningxia, China. <i>Genome Announcements</i> , 2013, 1, . | 0.8 | 4         |
| 51 | A NOVEL METHOD FOR ASSESSMENT OF 16S RRNA GENE COPY NUMBER IN BACTERIAL GENOMES BY PULSED-FIELD GEL ELECTROPHORESIS AND PCR AMPLIFICATION. <i>Journal of Rapid Methods and Automation in Microbiology</i> , 2009, 17, 274-279.                   | 0.4 | 1         |
| 52 | Phylogenetic analysis of aerobic anoxygenic phototrophic bacteria and their relatives based on farnesyl pyrophosphate synthase gene. <i>Acta Oceanologica Sinica</i> , 2010, 29, 82-89.  | 1.0 | 1         |
| 53 | Whole genome sequences of a free-living <i>Pseudomonas</i> sp. strain ML96 isolated from a freshwater Maar Lake. <i>Marine Genomics</i> , 2015, 24, 219-221.   | 1.1 | 1         |
| 54 | High-quality draft genome sequence of <i>Aquidulcibacter paucihalophilus</i> TH1-2T isolated from cyanobacterial aggregates in a eutrophic lake. <i>Standards in Genomic Sciences</i> , 2017, 12, 69.  | 1.5 | 1         |

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|----|--|-----|-----------|
| 55 | Draft Genome Sequence of <i>Aquicola tertiaricarbonis</i> MIMt <sub>k</sub> pLc11, an Aerobic Anoxygenic Phototrophic Bacterial Strain Isolated from Biological Soil Crusts. <i>Microbiology Resource Announcements</i> , 2018, 7, . | 0.6 | 1         |
| 56 | Genome of Betaproteobacterium <i>Caenimonas</i> sp. Strain SL110 Contains a Coenzyme F420 Biosynthesis Gene Cluster. <i>Journal of Microbiology and Biotechnology</i> , 2014, 24, 1490-1494.   | 2.1 | 1         |
| 57 | Contrasting diversity pattern of <i>Cytophaga</i> "Flavobacteria" in the estuarine and open ocean regions of the East China Sea. <i>Marine Biology Research</i> , 2007, 3, 428-437.  | 0.7 | 0         |
| 58 | Draft genome sequence of <i>Elstera cyanobacteriorum</i> , a novel facultative aerobic bacterium isolated from cyanobacterial aggregates in a eutrophic lake. <i>Gene Reports</i> , 2017, 9, 136-138.                                | 0.8 | 0         |