

# Ya Zhang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3231626/publications.pdf>

Version: 2024-02-01

22  
papers

2,753  
citations

471371

17  
h-index

642610

23  
g-index

24  
all docs

24  
docs citations

24  
times ranked

2438  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Climate warming enhances microbial network complexity and stability. <i>Nature Climate Change</i> , 2021, 11, 343-348.  | 8.1 | 672       |
| 2  | Global diversity and biogeography of bacterial communities in wastewater treatment plants. <i>Nature Microbiology</i> , 2019, 4, 1183-1195.   | 5.9 | 491       |
| 3  | A quantitative framework reveals ecological drivers of grassland microbial community assembly in response to warming. <i>Nature Communications</i> , 2020, 11, 4717.  | 5.8 | 417       |
| 4  | Potential impacts of changing supply-water quality on drinking water distribution: A review. <i>Water Research</i> , 2017, 116, 135-148.  | 5.3 | 211       |
| 5  | Small and mighty: adaptation of superphylum Patescibacteria to groundwater environment drives their genome simplicity. <i>Microbiome</i> , 2020, 8, 51.   | 4.9 | 205       |
| 6  | Assessing the origin of bacteria in tap water and distribution system in an unchlorinated drinking water system by SourceTracker using microbial community fingerprints. <i>Water Research</i> , 2018, 138, 86-96.              | 5.3 | 110       |
| 7  | Stepwise pH control to promote synergy of chemical and biological processes for augmenting short-chain fatty acid production from anaerobic sludge fermentation. <i>Water Research</i> , 2019, 155, 193-203.                    | 5.3 | 92        |
| 8  | Reduction of microbial diversity in grassland soil is driven by long-term climate warming. <i>Nature Microbiology</i> , 2022, 7, 1054-1062.   | 5.9 | 86        |
| 9  | Seasonal dynamics of the microbial community in two full-scale wastewater treatment plants: Diversity, composition, phylogenetic group based assembly and co-occurrence pattern. <i>Water Research</i> , 2021, 200, 117295.     | 5.3 | 83        |
| 10 | Hotspots for selected metal elements and microbes accumulation and the corresponding water quality deterioration potential in an unchlorinated drinking water distribution system. <i>Water Research</i> , 2017, 124, 435-445.  | 5.3 | 77        |
| 11 | Impact of drinking water treatment and distribution on the microbiome continuum: an ecological disturbance's perspective. <i>Environmental Microbiology</i> , 2017, 19, 3163-3174.  | 1.8 | 56        |
| 12 | Tracing fecal pollution sources in karst groundwater by Bacteroidales genetic biomarkers, bacterial indicators, and environmental variables. <i>Science of the Total Environment</i> , 2014, 490, 1082-1090.                    | 3.9 | 55        |
| 13 | The application of molecular tools to study the drinking water microbiome – Current understanding and future needs. <i>Critical Reviews in Environmental Science and Technology</i> , 2019, 49, 1188-1235.                      | 6.6 | 38        |
| 14 | 360-Degree Distribution of Biofilm Quantity and Community in an Operational Unchlorinated Drinking Water Distribution Pipe. <i>Environmental Science &amp; Technology</i> , 2020, 54, 5619-5628.                                | 4.6 | 33        |
| 15 | Ugly ducklings—the dark side of plastic materials in contact with potable water. <i>Npj Biofilms and Microbiomes</i> , 2018, 4, 7.  | 2.9 | 28        |
| 16 | Alkaline environments benefit microbial K-strategists to efficiently utilize protein substrate and promote valorization of protein waste into short-chain fatty acids. <i>Chemical Engineering Journal</i> , 2021, 404, 127147. | 6.6 | 24        |
| 17 | Phenotypic and Phylogenetic Identification of Coliform Bacteria Obtained Using 12 Coliform Methods Approved by the U.S. Environmental Protection Agency. <i>Applied and Environmental Microbiology</i> , 2015, 81, 6012-6023.   | 1.4 | 21        |
| 18 | Benefits of Genomic Insights and CRISPR-Cas Signatures to Monitor Potential Pathogens across Drinking Water Production and Distribution Systems. <i>Frontiers in Microbiology</i> , 2017, 8, 2036.                              | 1.5 | 15        |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Temporal Changes of Virus-Like Particle Abundance and Metagenomic Comparison of Viral Communities in Cropland and Prairie Soils. <i>MSphere</i> , 2021, 6, e0116020.  | 1.3 | 12        |
| 20 | Assessing the contribution of biofilm to bacterial growth during stagnation in shower hoses. <i>Water Science and Technology: Water Supply</i> , 2020, 20, 2564-2576. | 1.0 | 5         |
| 21 | Temporal Dynamics of Bacterial Communities along a Gradient of Disturbance in a U.S. Southern Plains Agroecosystem. <i>MBio</i> , 2022, 13, e0382921.                 | 1.8 | 4         |
| 22 | Nontuberculous Mycobacteria Infection: Source and Treatment. <i>Current Pulmonology Reports</i> , 2019, 8, 151-159.   | 0.5 | 1         |