## Rui Guan

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7593914/publications.pdf

Version: 2024-02-01

567281 794594 1,806 19 15 19 citations h-index g-index papers 21 21 21 3056 citing authors all docs docs citations times ranked

| #  | Article  | IF          | CITATIONS |
|----|--|-------------|-----------|
| 1  | The fungal root endophyte <i>Serendipita vermifera</i> displays inter-kingdom synergistic beneficial effects with the microbiota in <i>Arabidopsis thaliana</i> and barley. ISME Journal, 2022, 16, 876-889.         | 9.8         | 22        |
| 2  | Shared features and reciprocal complementation of the Chlamydomonas and Arabidopsis microbiota. Nature Communications, 2022, 13, 406.  | 12.8        | 28        |
| 3  | Maize Field Study Reveals Covaried Microbiota and Metabolic Changes in Roots over Plant Growth.<br>MBio, 2022, 13, e0258421.   | 4.1         | 15        |
| 4  | Host preference and invasiveness of commensal bacteria in the Lotus and Arabidopsis root microbiota. Nature Microbiology, 2021, 6, 1150-1162.  | 13.3        | 89        |
| 5  | Root-Secreted Coumarins and the Microbiota Interact to Improve Iron Nutrition in Arabidopsis. Cell<br>Host and Microbe, 2020, 28, 825-837.e6.  | 11.0        | 199       |
| 6  | On the origin of vertebrate body plan: Insights from the endoderm using the hourglass model. Gene Expression Patterns, 2020, 37, 119125.   | 0.8         | 2         |
| 7  | Alternate succession of aggregate-forming cyanobacterial genera correlated with their attached bacteria by co-pathways. Science of the Total Environment, 2019, 688, 867-879.  | 8.0         | 32        |
| 8  | Chromosome level comparative analysis of Brassica genomes. Plant Molecular Biology, 2019, 99, 237-249.   | 3.9         | 14        |
| 9  | The genetic architecture of floral traits in the woody plant Prunus mume. Nature Communications, 2018, 9, 1702.  | 12.8        | 73        |
| 10 | Evaluation of different 16S rRNA gene V regions for exploring bacterial diversity in a eutrophic freshwater lake. Science of the Total Environment, 2018, 618, 1254-1267.  | 8.0         | 115       |
| 11 | Phagocytic intracellular digestion in amphioxus ( <i>Branchiostoma</i> ). Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20180438.  | 2.6         | 11        |
| 12 | Draft genome sequence of the Tibetan medicinal herb Rhodiola crenulata. GigaScience, 2017, 6, 1-5.   | 6.4         | 33        |
| 13 | Microbial profiles of a drinking water resource based on different 16S rRNA V regions during a heavy cyanobacterial bloom in Lake Taihu, China. Environmental Science and Pollution Research, 2017, 24, 12796-12808. | <b>5.</b> 3 | 26        |
| 14 | Draft genome of the living fossil Ginkgo biloba. GigaScience, 2016, 5, 49.   | 6.4         | 232       |
| 15 | Complete genome sequence and genomic characterization of Microcystis panniformis FACHB 1757 by third-generation sequencing. Standards in Genomic Sciences, 2016, 11, 11.   | 1.5         | 19        |
| 16 | A genome draft of the legless anguid lizard, Ophisaurus gracilis. GigaScience, 2015, 4, 17.  | 6.4         | 23        |
| 17 | Reference genome of wild goat (capra aegagrus) and sequencing of goat breeds provide insight into genic basis of goat domestication. BMC Genomics, 2015, 16, 431.  | 2.8         | 103       |
| 18 | Identification of a novel salt tolerance gene in wild soybean by whole-genome sequencing. Nature Communications, 2014, 5, 4340.  | 12.8        | 332       |

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 19 | The sheep genome illuminates biology of the rumen and lipid metabolism. Science, 2014, 344, 1168-1173. | 12.6 | 436       |