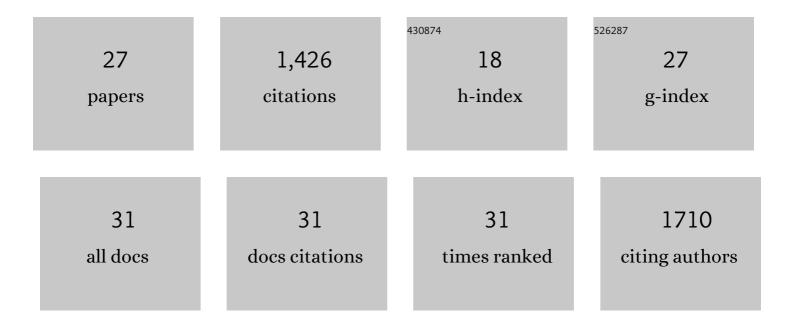
## Shingo Miyauchi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2691794/publications.pdf Version: 2024-02-01



| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Large-scale genome sequencing of mycorrhizal fungi provides insights into the early evolution of symbiotic traits. Nature Communications, 2020, 11, 5125.  | 12.8 | 258       |
| 2  | Dominant bacteria in soils of Marble Point and Wright Valley, Victoria Land, Antarctica. Soil Biology<br>and Biochemistry, 2006, 38, 3041-3056.  | 8.8  | 229       |
| 3  | Comparative genomics of <i>Rhizophagus irregularis</i> , <i> R.Âcerebriforme</i> , <i> R.Âdiaphanus</i><br>and <i>Gigaspora rosea</i> highlights specific genetic features in Glomeromycotina. New Phytologist,<br>2019, 222, 1584-1598. | 7.3  | 133       |
| 4  | Pezizomycetes genomes reveal the molecular basis of ectomycorrhizal truffle lifestyle. Nature Ecology and Evolution, 2018, 2, 1956-1965.   | 7.8  | 95        |
| 5  | Genomic Analysis Enlightens Agaricales Lifestyle Evolution and Increasing Peroxidase Diversity.<br>Molecular Biology and Evolution, 2021, 38, 1428-1446.   | 8.9  | 72        |
| 6  | The integrative omics of white-rot fungus Pycnoporus coccineus reveals co-regulated CAZymes for orchestrated lignocellulose breakdown. PLoS ONE, 2017, 12, e0175528.   | 2.5  | 64        |
| 7  | Genetic determinants of endophytism in the Arabidopsis root mycobiome. Nature Communications, 2021, 12, 7227.  | 12.8 | 58        |
| 8  | Human Papilloma Viruses and Breast Cancer. Frontiers in Oncology, 2015, 5, 277.  | 2.8  | 51        |
| 9  | Visual Comparative Omics of Fungi for Plant Biomass Deconstruction. Frontiers in Microbiology, 2016, 7, 1335.  | 3.5  | 46        |
| 10 | Integrative visual omics of the white-rot fungus Polyporus brumalis exposes the biotechnological potential of its oxidative enzymes for delignifying raw plant biomass. Biotechnology for Biofuels, 2018, 11, 201.                       | 6.2  | 45        |
| 11 | Gene family expansions and transcriptome signatures uncover fungal adaptations to wood decay.<br>Environmental Microbiology, 2021, 23, 5716-5732.  | 3.8  | 44        |
| 12 | Conserved white-rot enzymatic mechanism for wood decay in the Basidiomycota genus<br><i>Pycnoporus</i> . DNA Research, 2020, 27, .   | 3.4  | 32        |
| 13 | Insights into an unusual Auxiliary Activity 9 family member lacking the histidine brace motif of lytic polysaccharide monooxygenases. Journal of Biological Chemistry, 2019, 294, 17117-17130.   | 3.4  | 30        |
| 14 | An ectomycorrhizal fungus alters sensitivity to jasmonate, salicylate, gibberellin, and ethylene in host<br>roots. Plant, Cell and Environment, 2020, 43, 1047-1068.   | 5.7  | 30        |
| 15 | Human Papilloma Virus Identification in Breast Cancer Patients with Previous Cervical Neoplasia.<br>Frontiers in Oncology, 2015, 5, 298.   | 2.8  | 29        |
| 16 | Expression of a bacterial xylanase in Trichoderma reesei under the egl2 and cbh2 glycosyl hydrolase<br>gene promoters. New Biotechnology, 2013, 30, 523-530.   | 4.4  | 26        |
| 17 | Evolution of the Mode of Nutrition in Symbiotic and Saprotrophic Fungi in Forest Ecosystems. Annual<br>Review of Ecology, Evolution, and Systematics, 2021, 52, 385-404.   | 8.3  | 26        |
| 18 | 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        |

**Shingo Miyauchi** 

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|----|---|-----|-----------|
| 19 | Evolutionary transition to the ectomycorrhizal habit in the genomes of a hyperdiverse lineage of mushroomâ€forming fungi. New Phytologist, 2022, 233, 2294-2309.                  | 7.3 | 21        |
| 20 | Desert truffle genomes reveal their reproductive modes and new insights into plant–fungal<br>interaction and ectendomycorrhizal lifestyle. New Phytologist, 2021, 229, 2917-2932. | 7.3 | 19        |
| 21 | Evolutionary innovations through gain and loss of genes in the ectomycorrhizal Boletales. New Phytologist, 2022, 233, 1383-1400.  | 7.3 | 19        |
| 22 | Dynamics of the Phanerochaete carnosa transcriptome during growth on aspen and spruce. BMC Genomics, 2018, 19, 815.   | 2.8 | 15        |
| 23 | Comparative genomics reveals a dynamic genome evolution in the ectomycorrhizal milkâ€cap<br>( <i>Lactarius</i> ) mushrooms. New Phytologist, 2022, 235, 306-319.                  | 7.3 | 14        |
| 24 | Simultaneous expression of the bacterial Dictyoglomus thermophilum xynB gene under three different Trichoderma reesei promoters. New Biotechnology, 2014, 31, 98-103.             | 4.4 | 11        |
| 25 | A Transcriptomic Atlas of the Ectomycorrhizal Fungus Laccaria bicolor. Microorganisms, 2021, 9, 2612.   | 3.6 | 11        |
| 26 | Phylogenomics and Comparative Genomics Highlight Specific Genetic Features in Ganoderma Species.<br>Journal of Fungi (Basel, Switzerland), 2022, 8, 311.                          | 3.5 | 10        |
| 27 | Autism Susceptibility Genes and the Transcriptional Landscape of the Human Brain. International Review of Neurobiology, 2013, 113, 303-318.                                       | 2.0 | 7         |