## Seung-Yoon Oh

List of Publications by Year in descending order

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516561 501076 45 936 16 28 citations g-index h-index papers 45 45 45 1209 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Fungal diversity notes 929–1035: taxonomic and phylogenetic contributions on genera and species of fungi. Fungal Diversity, 2019, 95, 1-273.	4.7	203
2	Taxonomic evaluation of selected <i>Ganoderma</i> species and database sequence validation. PeerJ, 2017, 5, e3596.	0.9	44
3	Delimitation of <i>Russula</i> Subgenus <i>Amoenula</i> in Korea Using Three Molecular Markers. Mycobiology, 2013, 41, 191-201.	0.6	42
4	Identifying airborne fungi in Seoul, Korea using metagenomics. Journal of Microbiology, 2014, 52, 465-472.	1.3	42
5	Distinctive Feature of Microbial Communities and Bacterial Functional Profiles in Tricholoma matsutake Dominant Soil. PLoS ONE, 2016, 11, e0168573.	1.1	39
6	Effect of fruiting body bacteria on the growth of Tricholoma matsutake and its related molds. PLoS ONE, 2018, 13, e0190948.	1.1	36
7	Marine-derived Penicillium in Korea: diversity, enzyme activity, and antifungal properties. Antonie Van Leeuwenhoek, 2014, 106, 331-345.	0.7	34
8	Taxonomic annotation of public fungal ITS sequences from the built environment – a report from an April 10–11, 2017 workshop (Aberdeen, UK). MycoKeys, 2018, 28, 65-82.	0.8	33
9	Linking a Gene Cluster to Atranorin, a Major Cortical Substance of Lichens, through Genetic Dereplication and Heterologous Expression. MBio, 2021, 12, e0111121.	1.8	33
10	Root-associated bacteria influencing mycelial growth of Tricholoma matsutake (pine mushroom). Journal of Microbiology, 2018, 56, 399-407.	1.3	30
11	The diversity and ecological roles of Penicillium in intertidal zones. Scientific Reports, 2019, 9, 13540.	1.6	29
12	A systematic revision of the ectomycorrhizal genus <i>Laccaria</i> from Korea. Mycologia, 2018, 110, 948-961.	0.8	25
13	Species delimitation of three species within the Russula subgenus Compacta in Korea: R. eccentrica, R. nigricans, and R. subnigricans. Journal of Microbiology, 2014, 52, 631-638.	1.3	21
14	Trichoderma songyi sp. nov., a new species associated with the pine mushroom (Tricholoma) Tj ETQq0 0 0 rgBT /	/Overlock 0.7	10 Tf 50 222
15	Diversity and enzyme activity of Penicillium species associated with macroalgae in Jeju Island. Journal of Microbiology, 2016, 54, 646-654.	1.3	18
16	Diversity and effect of Trichoderma isolated from the roots of Pinus densiflora within the fairy ring of pine mushroom (Tricholoma matsutake). PLoS ONE, 2018, 13, e0205900.	1.1	18
17	<i>Penicillium jejuense</i> sp. nov., isolated from the marine environments of Jeju Island, Korea. Mycologia, 2015, 107, 209-216.	0.8	17
18	Diversity and Distribution Patterns of Endolichenic Fungi in Jeju Island, South Korea. Sustainability, 2020, 12, 3769.	1.6	17

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19	Sequence Validation for the Identification of the White-Rot Fungi Bjerkandera in Public Sequence Databases. Journal of Microbiology and Biotechnology, 2014, 24, 1301-1307.	0.9	17
20	Effect of fairy ring bacteria on the growth of Tricholoma matsutake in vitro culture. Mycorrhiza, 2018, 28, 411-419.	1.3	16
21	Fungal diversity and enzyme activity associated with sailfin sandfish egg masses in Korea. Fungal Ecology, 2018, 34, 1-9.	0.7	14
22	Effect of Isolation Conditions on Diversity of Endolichenic Fungal Communities from a Foliose Lichen, Parmotrema tinctorum. Journal of Fungi (Basel, Switzerland), 2021, 7, 335.	1.5	13
23	Morphometrics of the final instar exuviae of five cicada species occurring in urban areas of central Korea. Journal of Asia-Pacific Entomology, 2012, 15, 627-630.	0.4	12
24	Lactarius cucurbitoides (Russulales, Basidiomycota), a new species from South Korea supported by molecular and morphological data. Phytotaxa, 2015, 205, 168.	0.1	12
25	Reâ€evaluation of <i>Armillaria</i> and <i>Desarmillaria</i> in South Korea based on <scp>ITS</scp> / <i>tef</i> 1 sequences and morphological characteristics. Forest Pathology, 2018, 48, e12447.	0.5	11
26	Influence of Season and Soil Properties on Fungal Communities of Neighboring Climax Forests (Carpinus cordata and Fraxinus rhynchophylla). Frontiers in Microbiology, 2020, 11, 572706.	1.5	11
27	Investigating Wood Decaying Fungi Diversity in Central Siberia, Russia Using ITS Sequence Analysis and Interaction with Host Trees. Sustainability, 2020, 12, 2535.	1.6	11
28	Successional Variation in the Soil Microbial Community in Odaesan National Park, Korea. Sustainability, 2020, 12, 4795.	1.6	11
29	Host availability hypothesis: complex interactions with abiotic factors and predators may best explain population densities of cicada species. Animal Cells and Systems, 2014, 18, 143-153.	0.8	10
30	Four New Species of <i>Amanita</i> in Inje County, Korea. Mycobiology, 2015, 43, 408-414.	0.6	10
31	Successional Change of the Fungal Microbiome Pine Seedling Roots Inoculated With Tricholoma matsutake. Frontiers in Microbiology, 2020, 11, 574146.	1.5	10
32	Co-occurrence patterns of wood-decaying fungi and ants in dead pines of South Korea. Journal of Asia-Pacific Entomology, 2019, 22, 1154-1160.	0.4	8
33	The Influence of Microfungi on the Mycelial Growth of Ectomycorrhizal Fungus Tricholoma matsutake. Microorganisms, 2019, 7, 169.	1.6	8
34	Endolichenic Fungal Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of <i>Parmotrema tinctorum </i> i>Name of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Metabarcoding: A Case Study of Community Analysis by Pure Culture Isolation and Community Analysis by Pure Culture Isolation Analysis by Pure Culture Isol	0.6	8
35	A Checklist of the Basidiomycetous Macrofungi and a Record of Five New Species from Mt. Oseo in Korea. Mycobiology, 2014, 42, 132-139.	0.6	7
36	Diversity of fungi associated with roots of Calanthe orchid species in Korea. Journal of Microbiology, 2018, 56, 49-55.	1.3	7

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37	<strong>Two new foliicolous species of <em>Strigula </em>(Strigulaceae, Strigulales) in Korea offer insight in phorophyte-dependent variation of thallus morphology</strong> . Phytotaxa, 2020, 443, 1-12.	0.1	7
38	Biodiversity and Community Structure of Mesozooplankton in the Marine and Coastal National Park Areas of Korea. Diversity, 2020, 12, 233.	0.7	7
39	Re-evaluation of the Genus <i>Antrodia</i> (Polyporales, Basidiomycota) in Korea. Mycobiology, 2014, 42, 114-119.	0.6	6
40	Guild Patterns of Basidiomycetes Community Associated With Quercus mongolica in Mt. Jeombong, Republic of Korea. Mycobiology, 2018, 46, 13-23.	0.6	6
41	Cellulosic Nanomaterial Production Via Fermentation by Komagataeibacter sp. SFCB22-18 Isolated from Ripened Persimmons. Journal of Microbiology and Biotechnology, 2019, 29, 617-624.	0.9	4
42	High mortality in Bufo gargarizans eggs associated with an undescribed Saprolegnia ferax strain in the Republic of Korea. Diseases of Aquatic Organisms, 2019, 137, 89-99.	0.5	4
43	Distribution of Foliicolous Lichen Strigula and Genetic Structure of S. multiformis on Jeju Island, South Korea. Microorganisms, 2019, 7, 430.	1.6	3
44	Fungal diversity living in the root and sporophore of the endemic Korean fern Mankyua chejuense. Fungal Ecology, 2021, 50, 101038.	0.7	3
45	Determination of Diversity, Distribution and Host Specificity of Korean <i>Laccaria</i> Using Four Approaches. Mycobiology, 2021, 49, 461-468.	0.6	O