

Kin Ming Tsui

List of Publications by Year in descending order

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121
papers

3,733
citations

159358

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155451

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129
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129
times ranked

4076
citing authors

#	ARTICLE	IF	CITATIONS
1	Genotyping and Drug Resistance Profile of Clinical Isolates of <i>Candida albicans</i> from Vulvovaginal Candidiasis in the Eastern China. <i>Mycopathologia</i> , 2022, 187, 217-224.	1.3	5
2	Cryptic Species Diversity and Phylogenetic Relationship in the Rust Genus <i>Chrysomyxa</i> from China. <i>Journal of Fungi</i> (Basel, Switzerland), 2022, 8, 83.	1.5	6
3	Hit Compounds and Associated Targets in Intracellular <i>Mycobacterium tuberculosis</i> . <i>Molecules</i> , 2022, 27, 4446.	1.7	0
4	Genetic Structure and Asymmetric Migration of Wheat Stripe Rust Pathogen in Western Epidemic Areas of China. <i>Phytopathology</i> , 2021, 111, 1252-1260.	1.1	4
5	Changes in Bacterial and Fungal Microbiomes Associated with Tomatoes of Healthy and Infected by <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> . <i>Microbial Ecology</i> , 2021, 81, 1004-1017.	1.4	39
6	Emerging fungal pathogen: <i>Candida auris</i> . <i>Evolution, Medicine and Public Health</i> , 2021, 9, 246-247.	1.1	3
7	Molecular characterization of clinical carbapenem-resistant Enterobacterales from Qatar. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2021, 40, 1779-1785.	1.3	22
8	Genomic Epidemiology of <i>Candida auris</i> in Qatar Reveals Hospital Transmission Dynamics and a South Asian Origin. <i>Journal of Fungi</i> (Basel, Switzerland), 2021, 7, 240.	1.5	13
9	Draft Genome Sequences of Seven <i>Vibrio cholerae</i> Isolates from Adult Patients in Qatar. <i>Microbiology Resource Announcements</i> , 2021, 10, .	0.3	0
10	Potency of Olorofim (F901318) Compared to Contemporary Antifungal Agents against Clinical <i>Aspergillus fumigatus</i> Isolates and Review of Azole Resistance Phenotype and Genotype Epidemiology in China. <i>Antimicrobial Agents and Chemotherapy</i> , 2021, 65, .	1.4	13
11	Monitoring the effect of environmental conditions on safety of fresh produce sold in Qatar's wholesale market. <i>International Journal of Environmental Health Research</i> , 2021, , 1-19.	1.3	0
12	Whole-Genome Sequencing for Molecular Characterization of Carbapenem-Resistant Enterobacteriaceae Causing Lower Urinary Tract Infection among Pediatric Patients. <i>Antibiotics</i> , 2021, 10, 972.	1.5	14
13	Contribution to rust flora in China I, tremendous diversity from natural reserves and parks. <i>Fungal Diversity</i> , 2021, 110, 1-58.	4.7	12
14	Real-Time SARS-CoV-2 Genotyping by High-Throughput Multiplex PCR Reveals the Epidemiology of the Variants of Concern in Qatar. <i>International Journal of Infectious Diseases</i> , 2021, 112, 52-54.	1.5	59
15	Trends in fecal carriage of carbapenemase-producing Enterobacterales in children before and after the implementation of international travel restrictions in response to COVID-19. <i>Travel Medicine and Infectious Disease</i> , 2021, 43, 102120.	1.5	2
16	Disease-induced changes in plant microbiome assembly and functional adaptation. <i>Microbiome</i> , 2021, 9, 187.	4.9	157
17	Phylogeny and biogeography of the Japanese rhinoceros beetle, <i>Trypoxylus dichotomus</i> (Coleoptera: Tj ETQq1 1 0.784314 rsgBT /Ove	0.8	11
18	Draft Genome Sequence of <i>Rhodotorula mucilaginosa</i> from an Adult Patient in Qatar. <i>Microbiology Resource Announcements</i> , 2021, 10, e0072521.	0.3	1

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19	Peritoneal Dialysisâ€“Associated Peritonitis Caused by <i>Mycobacterium abscessus</i> in Childrenâ€“A Case Report. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofaa579.	0.4	4
20	Fecal Carriage and Molecular Characterization of Carbapenemase-Producing <i>Enterobacterales</i> in the Pediatric Population in Qatar. <i>Microbiology Spectrum</i> , 2021, 9, e0112221.	1.2	6
21	1247. Molecular Epidemiology of Multi-drug Resistant <i>Klebsiella pneumoniae</i> and <i>K. quasipneumoniae</i> in Qatar. <i>Open Forum Infectious Diseases</i> , 2021, 8, S712-S712.	0.4	0
22	First report of NDM-1-producing <i>Pseudomonas aeruginosa</i> in the Arabian Peninsula. <i>Journal of Global Antimicrobial Resistance</i> , 2021, , .	0.9	0
23	Phylogenetic Relationships, Speciation, and Origin of <i>Armillaria</i> in the Northern Hemisphere: A Lesson Based on rRNA and Elongation Factor 1-Alpha. <i>Journal of Fungi (Basel, Switzerland)</i> , 2021, 7, 1088.	1.5	8
24	Plasmid-mediated colistin resistance encoded by <i>mcr-1</i> gene in <i>Escherichia coli</i> co-carrying <i>bla</i> CTX-M-15 and <i>bla</i> NDM-1 genes in pediatric patients in Qatar. <i>Journal of Global Antimicrobial Resistance</i> , 2020, 22, 662-663.	0.9	11
25	Molecular Characterization of Extended-Spectrum β -Lactamaseâ€“Producing <i>Escherichia coli</i> and <i>Klebsiella pneumoniae</i> Among the Pediatric Population in Qatar. <i>Frontiers in Microbiology</i> , 2020, 11, 581711.	1.5	16
26	Emerging <i>Cryptococcus gattii</i> species complex infections in Guangxi, southern China. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008493.	1.3	12
27	First Case of Rhinocerebral Mucormycosis Caused by <i>Lichtheimia ornata</i> , with a Review of <i>Lichtheimia</i> Infections. <i>Mycopathologia</i> , 2020, 185, 555-567.	1.3	18
28	Draft Genome Sequence of an Extended-Spectrum β -Lactamase-Producing <i>Klebsiella oxytoca</i> Strain Bearing <i>mcr-9</i> from Qatar. <i>Microbiology Resource Announcements</i> , 2020, 9, .	0.3	10
29	Unusual accumulation of a wide array of antimicrobial resistance mechanisms in a patient with cytomegalovirus-associated hemophagocytic lymphohistiocytosis: a case report. <i>BMC Infectious Diseases</i> , 2020, 20, 237.	1.3	7
30	A metagenomics-based diagnostic approach for central nervous system infections in hospital acute care setting. <i>Scientific Reports</i> , 2020, 10, 11194.	1.6	19
31	MymA Bioactivated Thioalkylbenzoxazole Prodrug Family Active against <i>Mycobacterium tuberculosis</i> . <i>Journal of Medicinal Chemistry</i> , 2020, 63, 4732-4748.	2.9	12
32	Intraspecific Diversity and Taxonomy of <i>Emmonsia crescens</i> . <i>Mycopathologia</i> , 2020, 185, 613-627.	1.3	15
33	Emerging <i>Cryptococcus gattii</i> species complex infections in Guangxi, southern China. , 2020, 14, e0008493.		0
34	Emerging <i>Cryptococcus gattii</i> species complex infections in Guangxi, southern China. , 2020, 14, e0008493.		0
35	Emerging <i>Cryptococcus gattii</i> species complex infections in Guangxi, southern China. , 2020, 14, e0008493.		0
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37	Cryptic Speciation in Western North America and Eastern Eurasia of the Pathogens Responsible for Laminated Root Rot. <i>Phytopathology</i> , 2019, 109, 456-468.	1.1	8
38	Draft Genome Sequences of Two <i>Streptococcus pneumoniae</i> Strains Causing Invasive Infections in Children in Qatar. <i>Microbiology Resource Announcements</i> , 2019, 8, .	0.3	1
39	Fine-scale genetic diversity and relatedness in fungi associated with the mountain pine beetle. <i>Canadian Journal of Forest Research</i> , 2019, 49, 933-941.	0.8	4
40	Improved detection and identification of the sudden oak death pathogen <i>Phytophthora ramorum</i> and the Port Orford cedar root pathogen <i>Phytophthora lateralis</i> . <i>Plant Pathology</i> , 2019, 68, 878-888.	1.2	14
41	Evaluation of VITEK MS, Clin-ToF-II MS, Autof MS 1000 and VITEK 2 ANC card for identification of <i>Bacteroides fragilis</i> group isolates and antimicrobial susceptibilities of these isolates in a Chinese university hospital. <i>Journal of Microbiology, Immunology and Infection</i> , 2019, 52, 456-464.	1.5	15
42	Transcriptional profile of the human skin pathogenic fungus <i>Mucor irregularis</i> in response to low oxygen. <i>Medical Mycology</i> , 2018, 56, e2-e2.	0.3	0
43	Protein tyrosine kinase, PtkA, is required for <i>Mycobacterium tuberculosis</i> growth in macrophages. <i>Scientific Reports</i> , 2018, 8, 155.	1.6	30
44	Beaver Fever: Whole-Genome Characterization of Waterborne Outbreak and Sporadic Isolates To Study the Zoonotic Transmission of Giardiasis. <i>MSphere</i> , 2018, 3, .	1.3	34
45	Transcriptional profile of the human skin pathogenic fungus <i>Mucor irregularis</i> in response to low oxygen. <i>Medical Mycology</i> , 2018, 56, 631-644.	0.3	11
46	The world's ten most feared fungi. <i>Fungal Diversity</i> , 2018, 93, 161-194.	4.7	85
47	Genome-Enhanced Detection and Identification (GEDI) of plant pathogens. <i>PeerJ</i> , 2018, 6, e4392.	0.9	24
48	Genetic and genomic evidence of niche partitioning and adaptive radiation in mountain pine beetle fungal symbionts. <i>Molecular Ecology</i> , 2017, 26, 2077-2091.	2.0	52
49	The influence of the mating type on virulence of <i>Mucor irregularis</i> . <i>Scientific Reports</i> , 2017, 7, 10629.	1.6	7
50	Triazole phenotypes and genotypic characterization of clinical <i>Aspergillus fumigatus</i> isolates in China. <i>Emerging Microbes and Infections</i> , 2017, 6, 1-6.	3.0	26
51	Genome Sequences of the <i>Mycobacterium tuberculosis</i> H37Rv- ptkA Deletion Mutant and Its Parental Strain. <i>Genome Announcements</i> , 2017, 5, .	0.8	2
52	<i>Dothiorella magnoliae</i> , a new species associated with dieback of <i>Magnolia grandiflora</i> from China. <i>Mycosphere</i> , 2017, 8, 1031-1041.	1.9	2
53	Phylogenetic evidence places the coraloid jelly fungus <i>Tremellodendropsis tuberosa</i> (Tremellodendropsidales) among early diverging Agaricomycetes. <i>Mycological Progress</i> , 2016, 15, 939-946.	0.5	4
54	Molecular phylogeny, pathogenicity and toxigenicity of <i>Fusarium oxysporum</i> f. sp. <i>lycopersici</i> . <i>Scientific Reports</i> , 2016, 6, 21367.	1.6	89

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55	Biology and Ecology of Freshwater Fungi. <i>Fungal Biology</i> , 2016, , 285-313.	0.3	22
56	Friends or foes? Emerging insights from fungal interactions with plants. <i>FEMS Microbiology Reviews</i> , 2016, 40, 182-207.	3.9	238
57	Bioprospecting Fungi and the Labyrinthulomycetes at the Ocean-Land Interface. , 2015, , 379-392.		0
58	Global Spread of Human Chromoblastomycosis Is Driven by Recombinant <i>Cladophialophora carrionii</i> and Predominantly Clonal <i>Fonsecaea</i> Species. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0004004.	1.3	21
59	Colonization History, Host Distribution, Anthropogenic Influence and Landscape Features Shape Populations of White Pine Blister Rust, an Invasive Alien Tree Pathogen. <i>PLoS ONE</i> , 2015, 10, e0127916.	1.1	19
60	<i>Giardia</i> spp. Are Commonly Found in Mixed Assemblages in Surface Water, as Revealed by Molecular and Whole-Genome Characterization. <i>Applied and Environmental Microbiology</i> , 2015, 81, 4827-4834.	1.4	38
61	The Faces of Fungi database: fungal names linked with morphology, phylogeny and human impacts. <i>Fungal Diversity</i> , 2015, 74, 3-18.	4.7	471
62	<i>Helicocentralis hyalina</i> gen. et sp. nov., an aero-aquatic helicosporous fungus (Leotiomyces), <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462</i>	0.5	4
63	Asexual Propagation of a Virulent Clone Complex in a Human and Feline Outbreak of Sporotrichosis. <i>Eukaryotic Cell</i> , 2015, 14, 158-169.	3.4	47
64	The MAT1-1:MAT1-2 Ratio of <i>Sporothrix globosa</i> Isolates in Japan. <i>Mycopathologia</i> , 2015, 179, 81-86.	1.3	12
65	Single-nucleotide polymorphism discovery in <i>Leptographium longiclavatum</i> , a mountain pine beetle-associated symbiotic fungus, using whole-genome resequencing. <i>Molecular Ecology Resources</i> , 2014, 14, 401-410.	2.2	13
66	Population Structure of Mountain Pine Beetle Symbiont <i>Leptographium longiclavatum</i> and the Implication on the Multipartite Beetle-Fungi Relationships. <i>PLoS ONE</i> , 2014, 9, e105455.	1.1	30
67	The molecular phylogeny of aquatic hyphomycetes with affinity to the Leotiomyces. <i>Fungal Biology</i> , 2013, 117, 660-672.	1.1	75
68	Unequal Recombination and Evolution of the Mating-Type (MAT) Loci in the Pathogenic Fungus <i>Grosmannia clavigera</i> and Relatives. <i>G3: Genes, Genomes, Genetics</i> , 2013, 3, 465-480.	0.8	49
69	Methods for Sampling and Analyzing Wetland Fungi. , 2013, , 93-121.		6
70	First Report of Pitch Canker Disease Caused by <i>Rhizosphaera kalkhoffii</i> on <i>Pinus sylvestris</i> in China. <i>Plant Disease</i> , 2013, 97, 283-283.	0.7	5
71	Population structure and migration pattern of a conifer pathogen, <i>Grosmannia clavigera</i> , as influenced by its symbiont, the mountain pine beetle. <i>Molecular Ecology</i> , 2012, 21, 71-86.	2.0	46
72	Zoospore production and motility of mangrove thraustochytrids from Hong Kong under various salinities. <i>Mycoscience</i> , 2012, 53, 1-9.	0.3	12

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73	Revision of lignicolous Tubeufiaceae based on morphological reexamination and phylogenetic analysis. <i>Fungal Diversity</i> , 2011, 51, 63-102.	4.7	95
74	Misidentification of OLGA-PH/J/92, believed to be the only crustacean cell line. <i>In Vitro Cellular and Developmental Biology - Animal</i> , 2011, 47, 665-674.	0.7	9
75	Genome and transcriptome analyses of the mountain pine beetle-fungal symbiont <i>Grosmannia clavigera</i> , a lodgepole pine pathogen. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 2504-2509.	3.3	218
76	Molecular techniques for pathogen identification and fungus detection in the environment. <i>IMA Fungus</i> , 2011, 2, 177-189.	1.7	81
77	Transfer of two <i>Helicoma</i> species to <i>Tropospora</i> based on molecular and morphological data. <i>Mycoscience</i> , 2010, 51, 144-148.	0.3	9
78	Rapid identification and detection of pine pathogenic fungi associated with mountain pine beetles by padlock probes. <i>Journal of Microbiological Methods</i> , 2010, 83, 26-33.	0.7	24
79	Multigene phylogeny of filamentous ambrosia fungi associated with ambrosia and bark beetles. <i>Mycological Research</i> , 2009, 113, 822-835.	2.5	88
80	Labyrinthulomycetes phylogeny and its implications for the evolutionary loss of chloroplasts and gain of ectoplasmic gliding. <i>Molecular Phylogenetics and Evolution</i> , 2009, 50, 129-140.	1.2	104
81	Characterization of microsatellite loci in the fungus, <i>Grosmannia clavigera</i> , a pine pathogen associated with the mountain pine beetle. <i>Molecular Ecology Resources</i> , 2009, 9, 1500-1503.	2.2	9
82	Genetic Diversity of the <i>Cryptococcus</i> Species Complex Suggests that <i>Cryptococcus gattii</i> Deserves to Have Varieties. <i>PLoS ONE</i> , 2009, 4, e5862.	1.1	144
83	Docosahexaenoic acid production and ultrastructure of the thraustochytrid <i>Aurantiochytrium mangrovei</i> MP2 under high glucose concentrations. <i>Mycoscience</i> , 2008, 49, 266-270.	0.3	30
84	Hyperbranched rolling circle amplification as a rapid and sensitive method for species identification within the <i>Cryptococcus</i> species complex. <i>Electrophoresis</i> , 2008, 29, 3183-3191.	1.3	59
85	Re-examining the phylogeny of clinically relevant <i>Candida</i> species and allied genera based on multigene analyses. <i>FEMS Yeast Research</i> , 2008, 8, 651-659.	1.1	54
86	<i>Tubeufia asiana</i> , the teleomorph of <i>Aquaphila albicans</i> in the Tubeufiaceae, Pleosporales, based on cultural and molecular data. <i>Mycologia</i> , 2007, 99, 884-894.	0.8	11
87	Phylogenetic relationships and convergence of helicosporous fungi inferred from ribosomal DNA sequences. <i>Molecular Phylogenetics and Evolution</i> , 2006, 39, 587-597.	1.2	62
88	Molecular systematics of <i>Helicoma</i> , <i>Helicomycetes</i> and <i>Helicosporium</i> and their teleomorphs inferred from rDNA sequences. <i>Mycologia</i> , 2006, 98, 94-104.	0.8	47
89	The halotolerant fungus <i>Glomerobolus gelineus</i> is a member of the Ostropales. <i>Mycological Research</i> , 2006, 110, 257-263.	2.5	35
90	Fungi on <i>Juncus roemerianus</i> . 17. New ascomycetes and the hyphomycete genus <i>Kolletes</i> gen. nov.. <i>Botanica Marina</i> , 2005, 48, .	0.6	6

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91	Fungi on submerged wood in the Koito River, Japan. <i>Mycoscience</i> , 2003, 44, 55-59.	0.3	14
92	Three new species of <i>Aquaticola</i> (Ascomycetes) from tropical freshwater habitats. <i>Nova Hedwigia</i> , 2003, 77, 161-168.	0.2	11
93	Reflections on the Genus <i>Vanakripa</i> , and a Description of <i>V. ellipsoidea</i> sp. nov.. <i>Mycologia</i> , 2003, 95, 124.	0.8	2
94	Reflections on the genus <i>Vanakripa</i> , and a description of <i>V. ellipsoidea</i> sp. nov. <i>Mycologia</i> , 2003, 95, 124-127.	0.8	4
95	Three new species of <i>Annulatascus</i> (Ascomycetes) from Hong Kong freshwater habitats. <i>Mycoscience</i> , 2002, 43, 383-389.	0.3	13
96	<i>Brunneosporella aquatica</i> gen. et sp. nov., <i>Aqualignicola hyalina</i> gen. et sp. nov., <i>Jobellisia viridifusca</i> sp. nov. and <i>Porosphaerellopsis bipolaris</i> sp. nov. (ascomycetes) from submerged wood in freshwater habitats. <i>Mycological Research</i> , 2001, 105, 625-633.	2.5	23
97	New Species or Records of <i>Cacumisporium</i> , <i>Helicosporium</i> , <i>Monotosporella</i> and <i>Bahusutrabeeja</i> on Submerged Wood in Hong Kong Streams. <i>Mycologia</i> , 2001, 93, 389.	0.8	12
98	New species or records of <i>Cacumisporium</i> , <i>Helicosporium</i> , <i>Monotosporella</i> and <i>Bahusutrabeeja</i> on submerged wood in Hong Kong streams. <i>Mycologia</i> , 2001, 93, 389-397.	0.8	25
99	<i>Paraniesslia tuberculata</i> gen. et sp. nov., and new records or species of <i>Clypeosphaeria</i> , <i>Leptosphaeria</i> and <i>Astrosphaeriella</i> in Hong Kong freshwater habitats. <i>Mycologia</i> , 2001, 93, 1002-1009.	0.8	13
100	<i>Paraniesslia tuberculata</i> gen. et sp. nov., and New Records or Species of <i>Clypeosphaeria</i> , <i>Leptosphaeria</i> and <i>Astrosphaeriella</i> in Hong Kong Freshwater Habitats. <i>Mycologia</i> , 2001, 93, 1002.	0.8	12
101	Three new <i>Ophioceras</i> species (Ascomycetes) from the tropics. <i>Mycoscience</i> , 2001, 42, 321-326.	0.3	9
102	Longitudinal and temporal distribution of freshwater ascomycetes and dematiaceous hyphomycetes on submerged wood in the Lam Tsuen River, Hong Kong. <i>Journal of the North American Benthological Society</i> , 2001, 20, 533-549.	3.0	36
103	Colonization patterns of wood-inhabiting fungi on baits in Hong Kong rivers, with reference to the effects of organic pollution. <i>Antonie Van Leeuwenhoek</i> , 2001, 79, 33-38.	0.7	23
104	New records or species of <i>Dictyochaeta</i> , <i>Endophragmiella</i> and <i>Ramichloridium</i> from submerged wood in Hong Kong freshwater streams. <i>Cryptogamie, Mycologie</i> , 2001, 22, 139-145.	0.2	5
105	<i>Verticicola caudatus</i> gen. et sp. nov., and a new species of <i>Rivulicola</i> from submerged wood in freshwater habitats. <i>Mycologia</i> , 2000, 92, 1019-1026.	0.8	14
106	<i>Verticicola caudatus</i> gen. et sp. nov., and a New Species of <i>Rivulicola</i> from Submerged Wood in Freshwater Habitats. <i>Mycologia</i> , 2000, 92, 1019.	0.8	8
107	<i>Torrentispora fibrosa</i> gen. sp. nov. (Annulatascaceae) from freshwater habitats. <i>Mycological Research</i> , 2000, 104, 1399-1403.	2.5	12
108	Biodiversity of fungi on submerged wood in Hong Kong streams. <i>Aquatic Microbial Ecology</i> , 2000, 21, 289-298.	0.9	81

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109	Ultrastructural Studies of <i>Massarina ingoldiana</i> and <i>M. purpurascens</i> . <i>Mycologia</i> , 1999, 91, 721.	0.8	1
110	The genera <i>Aniptodera</i> , <i>Halosarpheia</i> , <i>Nais</i> and <i>Phaeonectriella</i> from freshwater habitats. <i>Mycoscience</i> , 1999, 40, 165-183.	0.3	34
111	Reflections on <i>Menisporopsis</i> , with the addition of <i>M. multisetulata</i> sp. nov. from submerged wood in Hong Kong. <i>Mycological Research</i> , 1999, 103, 148-152.	2.5	12
112	<i>Massarina proprietunicata</i> sp. nov., from submerged wood in streams in Hong Kong. <i>Mycological Research</i> , 1999, 103, 1575-1578.	2.5	6
113	Role of fungi in freshwater ecosystems. <i>Biodiversity and Conservation</i> , 1998, 7, 1187-1206.	1.2	180
114	<i>Elegantimyces sporidesmiopsis</i> gen. et sp. nov. on submerged wood from Hong Kong. <i>Mycological Research</i> , 1998, 102, 239-242.	2.5	7
115	The hyphomycete genus <i>Acrogenospora</i> , with two new species and two new combinations. <i>Mycological Research</i> , 1998, 102, 1309-1315.	2.5	17
116	A new species of <i>Clohiesia</i> from Hong Kong. <i>Mycoscience</i> , 1998, 39, 257-259.	0.3	9
117	<i>Yinmingella mitriformis</i> gen. et sp.nov., a new sporodochial hyphomycete from submerged wood in Hong Kong. <i>Canadian Journal of Botany</i> , 1998, 76, 1693-1697.	1.2	2
118	A New Freshwater Species of <i>Saccardoella</i> from Hong Kong and South Africa. <i>Mycologia</i> , 1998, 90, 701.	0.8	5
119	A new freshwater species of <i>Saccardoella</i> from Hong Kong and South Africa. <i>Mycologia</i> , 1998, 90, 701-704.	0.8	6
120	Four new species of <i>Xylomyces</i> from submerged wood. <i>Mycological Research</i> , 1997, 101, 1323-1328.	2.5	26
121	A Re-Visit to the Evolution and Ecophysiology of the Labyrinthulomycetes. , 0, , .		0