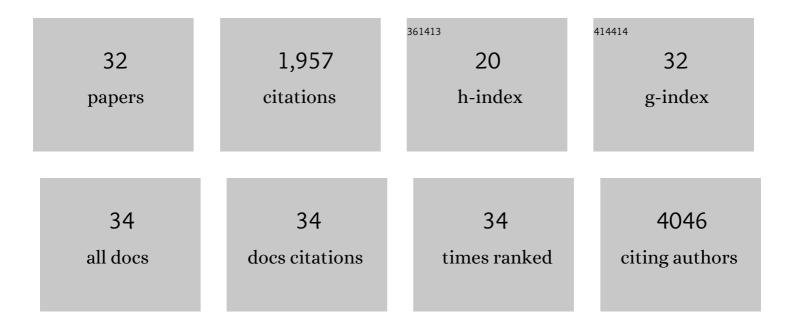
Wang-Ngai Chow

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7791365/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Induction of amphotericin B resistance in susceptible <i>Candida auris</i> by extracellular vesicles. Emerging Microbes and Infections, 2022, 11, 1900-1909.	6.5	12
2	A Sensitive and Specific Competitive Enzyme-Linked Immunosorbent Assay for Serodiagnosis of COVID-19 in Animals. Microorganisms, 2021, 9, 1019.	3.6	3
3	Disentangling sRNA-Seq data to study RNA communication between species. Nucleic Acids Research, 2020, 48, e21-e21.	14.5	8
4	Extracellular vesicles from Heligmosomoides bakeri and Trichuris muris contain distinct microRNA families and small RNAs that could underpin different functions in the host. International Journal for Parasitology, 2020, 50, 719-729.	3.1	16
5	A Rapid, Simple, Inexpensive, and Mobile Colorimetric Assay COVID-19-LAMP for Mass On-Site Screening of COVID-19. International Journal of Molecular Sciences, 2020, 21, 5380.	4.1	85
6	Identification of a Novel Betacoronavirus (Merbecovirus) in Amur Hedgehogs from China. Viruses, 2019, 11, 980.	3.3	42
7	Secretion of an Argonaute protein by a parasitic nematode and the evolution of its siRNA guides. Nucleic Acids Research, 2019, 47, 3594-3606.	14.5	75
8	Comparative analysis of small RNAs released by the filarial nematode Litomosoides sigmodontis in vitro and in vivo. PLoS Neglected Tropical Diseases, 2019, 13, e0007811.	3.0	19
9	<i>Daphnia magna</i> micro <scp>RNA</scp> s respond to nutritional stress and ageing but are not transgenerational. Molecular Ecology, 2018, 27, 1402-1412.	3.9	21
10	Immune stimuli shape the small non-coding transcriptome of extracellular vesicles released by dendritic cells. Cellular and Molecular Life Sciences, 2018, 75, 3857-3875.	5.4	57
11	Novel Partitivirus Enhances Virulence of and Causes Aberrant Gene Expression in Talaromyces marneffei. MBio, 2018, 9, .	4.1	54
12	Obstacles and opportunities in the functional analysis of extracellular vesicle RNA – an ISEV position paper. Journal of Extracellular Vesicles, 2017, 6, 1286095.	12.2	561
13	<i>Coronavirus HKU15</i> in respiratory tract of pigs and first discovery of coronavirus quasispecies in 5′-untranslated region. Emerging Microbes and Infections, 2017, 6, 1-7.	6.5	29
14	<i>In Vitro</i> Activity of Posaconazole against Talaromyces marneffei by Broth Microdilution and Etest Methods and Comparison to Itraconazole, Voriconazole, and Anidulafungin. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	20
15	Two novel dromedary camel bocaparvoviruses from dromedaries in the Middle East with unique genomic features. Journal of General Virology, 2017, 98, 1349-1359.	2.9	13
16	Metabolomic Profiling of Plasma from Melioidosis Patients Using UHPLC-QTOF MS Reveals Novel Biomarkers for Diagnosis. International Journal of Molecular Sciences, 2016, 17, 307.	4.1	15
17	Elizabethkingia anophelis bacteremia is associated with clinically significant infections and high mortality. Scientific Reports, 2016, 6, 26045.	3.3	146
18	Enterovirus D68 Infections Associated with Severe Respiratory Illness in Elderly Patients and Emergence of a Novel Clade in Hong Kong. Scientific Reports, 2016, 6, 25147.	3.3	44

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19	Matrix-assisted laser desorption ionization time-of-flight mass spectrometry for rapid identification of mold and yeast cultures of Penicillium marneffei. BMC Microbiology, 2016, 16, 36.	3.3	32
20	Metabolomic profiling of Burkholderia pseudomallei using UHPLC-ESI-Q-TOF-MS reveals specific biomarkers including 4-methyl-5-thiazoleethanol and unique thiamine degradation pathway. Cell and Bioscience, 2015, 5, 26.	4.8	12
21	Laboratory diagnosis of melioidosis: Past, present and future. Experimental Biology and Medicine, 2015, 240, 742-751.	2.4	79
22	Identification of specific metabolites in culture supernatant of <i>Mycobacterium tuberculosis</i> using metabolomics: exploration of potential biomarkers. Emerging Microbes and Infections, 2015, 4, 1-10.	6.5	55
23	Severe Acute Respiratory Syndrome (SARS) Coronavirus ORF8 Protein Is Acquired from SARS-Related Coronavirus from Greater Horseshoe Bats through Recombination. Journal of Virology, 2015, 89, 10532-10547.	3.4	172
24	Metabolomic Profiling of Plasma from Patients with Tuberculosis by Use of Untargeted Mass Spectrometry Reveals Novel Biomarkers for Diagnosis. Journal of Clinical Microbiology, 2015, 53, 3750-3759.	3.9	44
25	Signature Gene Expression Reveals Novel Clues to the Molecular Mechanisms of Dimorphic Transition in Penicillium marneffei. PLoS Genetics, 2014, 10, e1004662.	3.5	38
26	Identification of MicroRNA-Like RNAs in Mycelial and Yeast Phases of the Thermal Dimorphic Fungus Penicillium marneffei. PLoS Neglected Tropical Diseases, 2013, 7, e2398.	3.0	121
27	Unraveling the Molecular Basis of Temperature-Dependent Genetic Regulation in Penicillium marneffei. Eukaryotic Cell, 2013, 12, 1214-1224.	3.4	32
28	Draft Genome Sequence of Penicillium marneffei Strain PM1. Eukaryotic Cell, 2011, 10, 1740-1741.	3.4	42
29	Regulation of complementâ€3 protein expression in human and mouse oviducts. Molecular Reproduction and Development, 2009, 76, 301-308.	2.0	18
30	Complement 3 deficiency impairs early pregnancy in mice. Molecular Reproduction and Development, 2009, 76, 647-655.	2.0	36
31	Endocrine gland–derived vascular endothelial growth factor stimulates proliferation and tube formation in human uterine microvascular endothelial cell through the mitogen-activated protein kinase but not through the Akt pathway. Fertility and Sterility, 2009, 91, 2163-2171.	1.0	10
32	Preimplantation Embryos Cooperate with Oviductal Cells to Produce Embryotrophic Inactivated Complement-3b. Endocrinology, 2008, 149, 1268-1276.	2.8	45