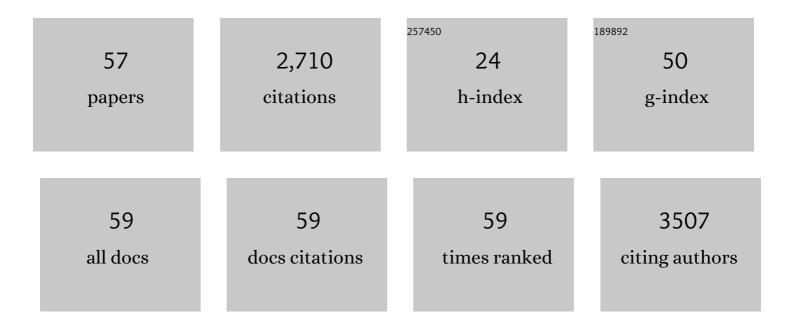
Hui-min Neoh

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Correlation between Reduced Daptomycin Susceptibility and Vancomycin Resistance in Vancomycin-Intermediate <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2006, 50, 1079-1082.	3.2	400
2	Targeting Staphylococcus aureus Toxins: A Potential form of Anti-Virulence Therapy. Toxins, 2016, 8, 72.	3.4	214
3	Novel Mechanism of Antibiotic Resistance Originating in Vancomycin-Intermediate <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2006, 50, 428-438.	3.2	213
4	An RpoB Mutation Confers Dual Heteroresistance to Daptomycin and Vancomycin in <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2010, 54, 5222-5233.	3.2	188
5	DNA Microarray-Based Identification of Genes Associated with Glycopeptide Resistance in <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2005, 49, 3404-3413.	3.2	150
6	Mutated Response Regulator graR Is Responsible for Phenotypic Conversion of Staphylococcus aureus from Heterogeneous Vancomycin-Intermediate Resistance to Vancomycin-Intermediate Resistance. Antimicrobial Agents and Chemotherapy, 2008, 52, 45-53.	3.2	147
7	Antibacterial performance of Ag nanoparticles and AgGO nanocomposites prepared via rapid microwave-assisted synthesis method. Nanoscale Research Letters, 2012, 7, 541.	5.7	144
8	<i>walK</i> and <i>clpP</i> Mutations Confer Reduced Vancomycin Susceptibility in <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2011, 55, 3870-3881.	3.2	138
9	Contribution of <i>vraSR</i> and <i>graSR</i> Point Mutations to Vancomycin Resistance in Vancomycin-Intermediate <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2009, 53, 1231-1234.	3.2	122
10	Serial Daptomycin Selection Generates Daptomycin-Nonsusceptible <i>Staphylococcus aureus</i> Strains with a Heterogeneous Vancomycin-Intermediate Phenotype. Antimicrobial Agents and Chemotherapy, 2008, 52, 4289-4299.	3.2	109
11	Pulsed-field gel electrophoresis (PFGE): A review of the "gold standard―for bacteria typing and current alternatives. Infection, Genetics and Evolution, 2019, 74, 103935.	2.3	100
12	Parvimonas micra, Peptostreptococcus stomatis, Fusobacterium nucleatum and Akkermansia muciniphila as a four-bacteria biomarker panel of colorectal cancer. Scientific Reports, 2021, 11, 2925.	3.3	83
13	Dengue epidemic in Malaysia: Not a predominantly urban disease anymore. BMC Research Notes, 2011, 4, 216.	1.4	69
14	Coordinated phenotype switching with large-scale chromosome flip-flop inversion observed in bacteria. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, E1647-56.	7.1	69
15	Impact of reduced vancomycin susceptibility on the therapeutic outcome of MRSA bloodstream infections. Annals of Clinical Microbiology and Antimicrobials, 2007, 6, 13.	3.8	63
16	Antibacterial hybrid cellulose–graphene oxide nanocomposite immobilized with silver nanoparticles. RSC Advances, 2015, 5, 26263-26268.	3.6	41
17	16S rRNA Gene Sequencing for Deciphering the Colorectal Cancer Gut Microbiome: Current Protocols and Workflows. Frontiers in Microbiology, 2018, 9, 767.	3.5	39
18	Advantages and Limitations of 16S rRNA Next-Generation Sequencing for Pathogen Identification in the Diagnostic Microbiology Laboratory: Perspectives from a Middle-Income Country. Diagnostics, 2020, 10. 816.	2.6	39

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#	Article	IF	CITATIONS
19	Metaproteomic analysis of human gut microbiota: where are we heading?. Journal of Biomedical Science, 2017, 24, 36.	7.0	36
20	Emergence of chikungunya seropositivity in healthy Malaysian adults residing in outbreak-free locations: Chikungunya seroprevalence results from the Malaysian Cohort. BMC Infectious Diseases, 2013, 13, 67.	2.9	34
21	Pathogens and Carcinogenesis: A Review. Biology, 2021, 10, 533.	2.8	32
22	Spatial Analysis of Colorectal Cancer Cases in Kuala Lumpur. Asian Pacific Journal of Cancer Prevention, 2014, 15, 1149-1154.	1.2	31
23	A graphene oxide facilitated a highly porous and effective antibacterial regenerated cellulose membrane containing stabilized silver nanoparticles. Cellulose, 2014, 21, 4261-4270.	4.9	26
24	Effective immobilization of silver nanoparticles on a regenerated cellulose–chitosan composite membrane and its antibacterial activity. New Journal of Chemistry, 2017, 41, 5061-5065.	2.8	26
25	Seropositivity of Dengue Antibodies during Pregnancy. Scientific World Journal, The, 2014, 2014, 1-4.	2.1	18
26	Activated ADI pathway: the initiator of intermediate vancomycin resistance in <i>Staphylococcus aureus</i> . Canadian Journal of Microbiology, 2017, 63, 260-264.	1.7	17
27	Comparison of Prognostic Accuracy of the quick Sepsis-Related Organ Failure Assessment between Short- & Long-term Mortality in Patients Presenting Outside of the Intensive Care Unit – A Systematic Review & Meta-analysis. Scientific Reports, 2018, 8, 16698.	3.3	17
28	Effects of simulated microgravity on gene expression and biological phenotypes of a single generation Caenorhabditis elegans cultured on 2 different media. Life Sciences in Space Research, 2017, 15, 11-17.	2.3	15
29	Improved Antimicrobial Activity of Linezolid against Vancomycin-Intermediate <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2008, 52, 4207-4208.	3.2	13
30	Epidemiological Study on Staphylococcus aureus Isolates Reveals Inverse Relationship between Antibiotic Resistance and Virulence Repertoire. Indian Journal of Microbiology, 2013, 53, 321-322.	2.7	13
31	Antimicrobial resistance profiling and molecular typing of methicillin-resistant Staphylococcus aureus isolated from a Malaysian teaching hospital. Journal of Medical Microbiology, 2016, 65, 1476-1481.	1.8	11
32	First report on the molecular epidemiology of Malaysian Staphylococcus epidermidis isolated from a University Teaching Hospital. BMC Research Notes, 2014, 7, 597.	1.4	10
33	Identification of Schizosaccharomyces pombe in the guts of healthy individuals and patients with colorectal cancer: preliminary evidence from a gut microbiome secretome study. Gut Pathogens, 2018, 10, 29.	3.4	9
34	Dengue epidemic in Malaysia: urban versus rural comparison of dengue immunoglobulin G seroprevalence among Malaysian adults aged 35–74 years. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2020, 114, 798-811.	1.8	8
35	Methicillin-Resistant Staphylococcus aureus (MRSA) Clonal Replacement in a Malaysian Teaching Hospital: Findings from an Eight-Year Interval Molecular Surveillance. Antibiotics, 2021, 10, 320.	3.7	8
36	Comparison of sPLA2IIA performance with high-sensitive CRP neutrophil percentage PCT and lactate to identify bacterial infection. Scientific Reports, 2021, 11, 11369.	3.3	8

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#	Article	IF	CITATIONS
37	Silver Nanoparticles - Graphene Oxide Nanocomposite for Antibacterial Purpose. Advanced Materials Research, 0, 364, 439-443.	0.3	7
38	Screening and Detection of Heterogenous Vancomycin Intermediate Staphylococcus aureus in Hospital Kuala Lumpur Malaysia, Using the Glycopeptide Resistance Detection Etest and Population Analysis Profiling. Infectious Disease Reports, 2012, 4, 71-72.	3.1	7
39	Screening and detection of heterogenous vancomycin intermediate Staphylococcus aureus in Hospital Kuala Lumpur Malaysia, using the glycopeptide resistance detection etest and population analysis profiling. Gastroenterology Insights, 2012, 4, 20.	1.2	6
40	Draft Genome Sequences of Four Nosocomial Methicillin-Resistant Staphylococcus aureus (MRSA) Strains (PPUKM-261-2009, PPUKM-332-2009, PPUKM-377-2009, and PPUKM-775-2009) Representative of Dominant MRSA Pulsotypes Circulating in a Malaysian University Teaching Hospital. Genome Announcements, 2013, 1, .	0.8	4
41	Establishment of an immunofluorescence assay to detect IgM antibodies to Nipah virus using HeLa cells expressing recombinant nucleoprotein. Journal of Virological Methods, 2019, 269, 83-87.	2.1	4
42	Analyzing the Secretome of Gut Microbiota as the Next Strategy For Early Detection of Colorectal Cancer. Proteomics, 2019, 19, 1800176.	2.2	4
43	Rapid Detection of Sepsis using CESDA: the Caenorhabditis elegans Sepsis Detection Assay. Revista Da Sociedade Brasileira De Medicina Tropical, 2019, 52, e20180300.	0.9	3
44	Molecular surveillance of methicillin-susceptible Staphylococcus aureus (MSSA) isolated over a one-year period from a Malaysian Teaching Hospital. Germs, 2020, 10, 104-111.	1.3	3
45	Clonal distribution and possible microevolution of methicillin-resistant Staphylococcus aureus strains in a teaching hospital in Malaysia. Asian Pacific Journal of Tropical Biomedicine, 2013, 3, 224-228.	1.2	2
46	The heterogeneic distribution of Helicobacter pylori cag pathogenicity island reflects different pathologies in multiracial Malaysian population. Brazilian Journal of Infectious Diseases, 2020, 24, 545-551.	0.6	2
47	Clonal Diversity of Methicillin-resistant Staphylococcus aureus in UKM Medical Centre: Characterisation by Multilocus Sequence Typing of Different SCCmec Type Representatives. Sains Malaysiana, 2015, 44, 1315-1323.	0.5	2
48	IDDF2018-ABS-0199â€Gut microbiome profiling of malaysians: a snapshot. , 2018, , .		1
49	Oxidative stress resistance and fitness-compensatory response in vancomycin-intermediate <i>Staphylococcus aureus</i> (VISA). Canadian Journal of Microbiology, 2019, 65, 623-628.	1.7	1
50	Prerequisite Evaluation of Anaerobic Settings for Gut Microbiome Functional Studies. Journal of Biomedicine and Translational Research, 2021, 7, 48-50.	0.2	1
51	Simplified protocol of gel-based multiple locus variable number of tandem repeats analysis (MLVA) for Staphylococcus aureus typing. Germs, 2017, 7, 98-100.	1.3	1
52	Seroprevalence of Hepatitis B and C in Healthy Malaysian Adults: A Preliminary Report. Journal of Biomedicine and Translational Research, 2019, 5, 23.	0.2	1
53	IDDF2018-ABS-0227â€Saccharomyces pombe may be a double-edge sword in the guts of malaysian adults: an evidence from the gut microbiome secretome study. , 2018, , .		0
54	IDDF2018-ABS-0228â€Human gut secretome profiling in malaysian adults: a case study on colorectal cancer. , 2018, , .		0

#	Article	IF	CITATIONS
55	IDDF2018-ABS-0191â€Aberrant gut mucosal microbiome signatures of malaysian colorectal cancer patients. , 2018, , .		0
56	IDDF2019-ABS-0300â€Dysbiosis of gut fungal microbiota in colorectal cancer: an assessment from faecal microbiome secretome. , 2019, , .		0
57	IDDF2019-ABS-0301â€Mapping of gut microbiome secretome in colorectal cancer: a malaysian data. , 2019, ,		0