Chun Wie Chong

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5689284/publications.pdf

Version: 2024-02-01

75 papers 1,928 citations

³⁶¹³⁸⁸
20
h-index

289230 40 g-index

78 all docs 78 docs citations

78 times ranked 2824 citing authors

#	Article	IF	CITATIONS
1	Deciphering chicken gut microbial dynamics based on high-throughput 16S rRNA metagenomics analyses. Gut Pathogens, 2015, 7, 4.	3.4	230
2	Terpene Derivatives as a Potential Agent against Antimicrobial Resistance (AMR) Pathogens. Molecules, 2019, 24, 2631.	3.8	220
3	Probiotics for Constipation in Parkinson Disease. Neurology, 2021, 96, e772-e782.	1.1	100
4	Gut Microbial Ecosystem in Parkinson Disease: New Clinicobiological Insights from Multiâ€Omics. Annals of Neurology, 2021, 89, 546-559.	5. 3	99
5	High levels of spatial heterogeneity in the biodiversity of soil prokaryotes on Signy Island, Antarctica. Soil Biology and Biochemistry, 2010, 42, 601-610.	8.8	86
6	Ethnicity influences the gut microbiota of individuals sharing a geographical location: a cross-sectional study from a middle-income country. Scientific Reports, 2021, 11, 2618.	3.3	73
7	Effect of ethnicity and socioeconomic variation to the gut microbiota composition among pre-adolescent in Malaysia. Scientific Reports, 2015, 5, 13338.	3.3	68
8	Environmental influences on bacterial diversity of soils on Signy Island, maritime Antarctic. Polar Biology, 2009, 32, 1571-1582.	1.2	63
9	Emerging spatial patterns in Antarctic prokaryotes. Frontiers in Microbiology, 2015, 6, 1058.	3.5	58
10	DGGE fingerprinting of bacteria in soils from eight ecologically different sites around Casey Station, Antarctica. Polar Biology, 2009, 32, 853-860.	1.2	57
11	Gut microbiome in Parkinson's disease: New insights from meta-analysis. Parkinsonism and Related Disorders, 2022, 94, 1-9.	2.2	55
12	Patterns in the distribution of soil bacterial 16S rRNA gene sequences from different regions of Antarctica. Geoderma, 2012, 181-182, 45-55.	5.1	50
13	Effect of short term variation in temperature and water content on the bacterial community in a tropical soil. Applied Soil Ecology, 2016, 107, 279-289.	4.3	48
14	Assessment of soil bacterial communities on Alexander Island (in the maritime and continental) Tj ETQq0 0 0 rgB	Γ/Overloc	k 10 Tf 50 22
15	Altered gut microbiome and metabolome in patients with multiple system atrophy. Movement Disorders, 2018, 33, 174-176.	3.9	45
16	Probiotics for Parkinson's disease: Current evidence and future directions. JGH Open, 2021, 5, 414-419.	1.6	41
17	The emergence of carbapenem resistant Klebsiella pneumoniae in Malaysia: correlation between microbiological trends with host characteristics and clinical factors. Antimicrobial Resistance and Infection Control, 2017, 6, 5.	4.1	33
18	Differences in the bacteriome of swab, saliva, and tissue biopsies in oral cancer. Scientific Reports, 2021, 11, 1181.	3.3	30

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19	Prevalence and risk factors of intestinal parasitism among two indigenous sub-ethnic groups in Peninsular Malaysia. Infectious Diseases of Poverty, 2016, 5, 77.	3.7	28
20	Pneumococcal conjugate vaccine implementation in middle-income countries. Pneumonia (Nathan Qld) Tj ETQo	10 0 0 rgB	Γ/Oygrlock 10
21	The identification of environmental parameters which could influence soil bacterial community composition on the Antarctic Peninsula - a statistical approach. Antarctic Science, 2012, 24, 249-258.	0.9	24
22	Salivary bacterial shifts in oral leukoplakia resemble the dysbiotic oral cancer bacteriome. Journal of Oral Microbiology, 2021, 13, 1857998.	2.7	24
23	Effect of temperature on bacterial community in petroleum hydrocarbon-contaminated and uncontaminated Antarctic soil. Polar Biology, 2018, 41, 1763-1775.	1.2	21
24	Occurrence of virulent multidrug-resistantEnterococcus faecalisandEnterococcus faeciumin the pigs, farmers and farm environments in Malaysia. PeerJ, 2018, 6, e5353.	2.0	21
25	Correlation between antibiotic consumption and the occurrence of multidrug-resistant organisms in a Malaysian tertiary hospital: a 3-year observational study. Scientific Reports, 2022, 12, 3106.	3.3	20
26	The Missing Piece: Recent Approaches Investigating the Antimicrobial Mode of Action of Essential Oils. Evolutionary Bioinformatics, 2021, 17, 117693432093839.	1.2	19
27	Spatial pattern in Antarctica: what can we learn from Antarctic bacterial isolates?. Extremophiles, 2013, 17, 733-745.	2.3	17
28	Intestinal carriage of multidrug-resistant gram-negative bacteria in preterm-infants during hospitalization in neonatal intensive care unit (NICU). Pathogens and Global Health, 2016, 110, 238-246.	2.3	17
29	Diversity of endocervical microbiota associated with genital Chlamydia trachomatis infection and infertility among women visiting obstetrics and gynecology clinics in Malaysia. PLoS ONE, 2019, 14, e0224658.	2.5	17
30	Comparative assessment of faecal microbial composition and metabonome of swine, farmers and human control. Scientific Reports, 2020, 10, 8997.	3.3	14
31	Compositional profile of mucosal bacteriome of smokers and smokeless tobacco users. Clinical Oral Investigations, 2022, 26, 1647-1656.	3.0	13
32	Comparative effectiveness of cardiovascular, renal and safety outcomes of secondâ€line antidiabetic drugs use in people with type 2 diabetes: A systematic review and network metaâ€analysis of randomised controlled trials. Diabetic Medicine, 2022, 39, e14780.	2.3	13
33	Fecal Calprotectin in Parkinson's Disease and Multiple System Atrophy. Journal of Movement Disorders, 2022, 15, 106-114.	1.3	13
34	Elucidating the survival and response of carbapenem resistant <i>Klebsiella pneumoniae</i> after exposure to imipenem at sub-lethal concentrations. Pathogens and Global Health, 2018, 112, 378-386.	2.3	11
35	LC-QTOF-MS analysis of xanthone content in different parts of <i>Garcinia mangostana</i> and its influence on cholinesterase inhibition. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 1433-1441.	5.2	11
36	Neonatal intensive care unit (NICU) exposures exert a sustained influence on the progression of gut microbiota and metabolome in the first year of life. Scientific Reports, 2021, 11, 1353.	3.3	11

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37	Molecular Characterization of Carbapenem Resistant Klebsiella pneumoniae in Malaysia Hospital. Pathogens, 2021, 10, 279.	2.8	11
38	Geographical separation and ethnic origin influence the human gut microbial composition: a meta-analysis from a Malaysian perspective. Microbial Genomics, 2021, 7, .	2.0	11
39	Molecular evidence of <i>Burkholderia pseudomallei </i> genotypes based on geographical distribution. Peerl, 2016, 4, e1802.	2.0	11
40	Clonal relatedness in the acquisition of intestinal carriage and transmission of multidrug resistant (MDR) Klebsiella pneumoniae and Escherichia coli and its risk factors among preterm infants admitted to the neonatal intensive care unit (NICU). Pediatrics and Neonatology, 2021, 62, 129-137.	0.9	10
41	One-step differential detection of Salmonella enterica serovar Typhi, serovar Paratyphi A and other Salmonella spp. by using a quadruplex real-time PCR assay. Journal of Microbiological Methods, 2021, 183, 106184.	1.6	10
42	Evaluation of Salmonella Typhi antigen YncE alongside HlyE for the detection of typhoid fever and its carriers. Medical Microbiology and Immunology, 2020, 209, 593-601.	4.8	9
43	In Vitro Efficacy of Flomoxef against Extended-Spectrum Beta-Lactamase-Producing Escherichia coli and Klebsiella pneumoniae Associated with Urinary Tract Infections in Malaysia. Antibiotics, 2021, 10, 181.	3.7	9
44	Understanding rat occurrences in oil palm plantation using high-resolution satellite image and GIS data. Precision Agriculture, 2018, 19, 42-54.	6.0	8
45	Association of antimicrobial resistance and gut microbiota composition in human and non-human primates at an urban ecotourism site. Gut Pathogens, 2020, 12, 14.	3.4	8
46	Whole genome analysis of multidrug-resistant Citrobacter freundii B9-C2 isolated from preterm neonate's stool in the first week. Journal of Global Antimicrobial Resistance, 2020, 21, 246-251.	2.2	8
47	Field evaluation of a rapid diagnostic test to detect antibodies in human toxocariasis. Acta Tropica, 2015, 148, 32-37.	2.0	7
48	Role of coaggregation in the pathogenicity and prolonged colonisation of Vibrio cholerae. Medical Microbiology and Immunology, 2019, 208, 793-809.	4.8	7
49	Genetic mechanisms and correlated risk factors of antimicrobial-resistant ESKAPEE pathogens isolated in a tertiary hospital in Malaysia. Antimicrobial Resistance and Infection Control, 2021, 10, 70.	4.1	7
50	Serotypes & Serotypes amp; penicillin susceptibility of Streptococcus pneumoniae isolated from children admitted to a tertiary teaching hospital in Malaysia. Indian Journal of Medical Research, 2018, 148, 225.	1.0	7
51	Pan-genome and resistome analysis of extended-spectrum ĀŸ-lactamase-producing Escherichia coli: A multi-setting epidemiological surveillance study from Malaysia. PLoS ONE, 2022, 17, e0265142.	2.5	7
52	Acclimatisation-induced stress influenced host metabolic and gut microbial composition change. Molecular BioSystems, 2015, 11, 297-306.	2.9	6
53	Coffee consumption revealed sex differences in host endogenous metabolism and gut microbiota in healthy adults. Journal of Food Biochemistry, 2020, 44, e13535.	2.9	6
54	Amelioration of high-fat diet-induced obesity and its associated complications by a myricetin derivative-rich fraction from <i>Syzygium malaccense</i> in C57BL/6J mice. Food and Function, 2021, 12, 5876-5891.	4.6	6

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55	Genotypic and metabolic approaches towards the segregation of Klebsiella pneumoniae strains producing different antibiotic resistant enzymes. Metabolomics, 2017, 13, 1.	3.0	5
56	Genomic Insights into Two Colistin-Resistant <i>Klebsiella pneumoniae</i> Strains Isolated from the Stool of Preterm Neonate During the First Week of Life. Microbial Drug Resistance, 2020, 26, 190-203.	2.0	5
57	Carriage of upper respiratory tract pathogens in rural communities of Sarawak, Malaysian Borneo. Pneumonia (Nathan Qld), 2021, 13, 6.	6.1	5
58	The Detection of Hypermucoviscous Carbapenem-Resistant <i>Klebsiella pneumoniae</i> from a Tertiary Teaching Hospital in Malaysia and Assessment of Hypermucoviscous as Marker of Hypervirulence. Microbial Drug Resistance, 2021, 27, 1319-1327.	2.0	5
59	Molecular Characterization of Carbapenem-Resistant Acinetobacter baumannii Isolated from the Intensive Care Unit in a Tertiary Teaching Hospital in Malaysia. Antibiotics, 2021, 10, 1340.	3.7	5
60	\hat{l}^2 -Lactam Resistance in Upper Respiratory Tract Pathogens Isolated from a Tertiary Hospital in Malaysia. Pathogens, 2021, 10, 1602.	2.8	5
61	Significant Clinical Presentation of Leptospirosis in Relation to Sociodemographic and Risk Factors in a Tertiary Hospital, Malaysia. Vector-Borne and Zoonotic Diseases, 2020, 20, 268-274.	1.5	4
62	The Prebiotic Potential of Geraniin and Geraniin-Enriched Extract against High-Fat-Diet-Induced Metabolic Syndrome in Sprague Dawley Rats. Antioxidants, 2022, 11, 632.	5.1	4
63	lsolation and Quantification of Mimivirus-Like and Marseillevirus-Like Viruses from Soil Samples in An Aboriginal <i>(</i> Orang asli <i>)</i> Village in Peninsular Malaysia. Intervirology, 2018, 61, 92-95.	2.8	3
64	Informing pneumococcal conjugate vaccine policy in middle-income countries: The case of Malaysia. Vaccine, 2017, 35, 2288-2290.	3.8	2
65	Effects of supplementing freezeâ€dried Mitsuokella jalaludinii phytase on the growth performance and gut microbial diversity of broiler chickens. Journal of Animal Physiology and Animal Nutrition, 2020, 104, 116-125.	2.2	2
66	First detection of Klebsiella quasipneumoniae producing OXA-181 carbapenemase in Malaysia. Journal of Infection and Chemotherapy, 2020, 26, 1058-1061.	1.7	2
67	A retrospective study on molecular epidemiology trends of carbapenem resistant <i>Enterobacteriaceae</i> in a teaching hospital in Malaysia. PeerJ, 2022, 10, e12830.	2.0	2
68	Impact of IsaA Gene Disruption: Decreasing Staphylococcal Biofilm and Alteration of Transcriptomic and Proteomic Profiles. Microorganisms, 2022, 10, 1119.	3.6	2
69	The Antibacterial Agent Identified from Acidocella spp. in the Fluid of Nepenthes gracilis Against Multidrug-Resistant Klebsiella pneumoniae: A Functional Metagenomic Approach. Microbial Drug Resistance, 2020, 27, 1018-1028.	2.0	1
70	Mitsuokella Jalaludinii Supplementation Improved Nutrient Utilization of Broilers Fed Low-Available Phosphorus Diet. Brazilian Journal of Poultry Science, 2021, 23, .	0.7	1
71	Genomic comparison and phenotypic profiling of small colony variants of Burkholderia pseudomallei. PLoS ONE, 2021, 16, e0261382.	2.5	1
72	Systems biology analyses of the dynamic host response to Toxoplasma gondii infection in a murine model. Parasitology Open, 2016, 2, .	0.9	0

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73	One-Step Differential Detection of OXA-48-Like Variants Using High-Resolution Melting (HRM) Analysis. Antibiotics, 2020, 9, 256.	3.7	O
74	Overview of the distribution of Burkholderia pseudomallei sequence types and the Emergence of sequence type 1342 in Malaysia. Transboundary and Emerging Diseases, 2021, , .	3.0	0
75	IDDF2021-ABS-0140â€Gut microbiota significantly correlate with body constitution in traditional chinese medicine. , 2021, , .		0