

# Kim-Kee Tan

## List of Publications by Year in descending order

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Version: 2024-02-01

40  
papers

871  
citations

623574

14  
h-index

477173

29  
g-index

42  
all docs

42  
docs citations

42  
times ranked

1583  
citing authors

#	ARTICLE	IF	CITATIONS
1	Differential heterologous neutralisation profile against strains within DENV-3 genotype II. <i>Epidemiology and Infection</i> , 2022, 150, e33.	1.0	0
2	Isolation of <i>Streptococcus cuniculi</i> from corneal lesion in laboratory-raised mice. <i>Journal of Veterinary Medical Science</i> , 2021, 83, 280-284.	0.3	0
3	Isolation of <i>Kytococcus schroeteri</i> from the brown rat <i>Rattus norvegicus</i> . <i>Asian Pacific Journal of Tropical Medicine</i> , 2021, 14, 191.	0.4	1
4	Multiplex sequencing of SARS-Cov-2 genome directly from clinical samples using the Ion Personal Genome Machine (PGM). <i>Tropical Biomedicine</i> , 2021, 38, 283-288.	0.2	3
5	Emergence of B.1.524(G) SARS-CoV-2 in Malaysia during the third COVID-19 epidemic wave. <i>Scientific Reports</i> , 2021, 11, 22105.	1.6	13
6	Evaluation of commercial serological assays in Malaysia for detection of anti-Zika virus antibodies. <i>Tropical Biomedicine</i> , 2021, 38, 613-621.	0.2	3
7	Serological evidence of DENV, JEV, and ZIKV among the indigenous people (Orang Asli) of Peninsular Malaysia. <i>Journal of Medical Virology</i> , 2020, 92, 956-962.	2.5	7
8	Bacterial communities in <i>Haemaphysalis</i> , <i>Dermacentor</i> and <i>Amblyomma</i> ticks collected from wild boar of an Orang Asli Community in Malaysia. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101352.	1.1	19
9	A reverse transcription loop-mediated isothermal amplification for broad coverage detection of Asian and African Zika virus lineages. <i>BMC Infectious Diseases</i> , 2020, 20, 947.	1.3	2
10	Continuous introduction, dispersal, and replacement of dengue virus type 3 genotype II sustained its transmission in Malaysia. <i>International Journal of Infectious Diseases</i> , 2020, 101, 236.	1.5	1
11	Draft genome of <i>Paraburkholderia fungorum</i> sequence type 868 recovered from human synovial tissues. <i>Data in Brief</i> , 2019, 25, 104159.	0.5	6
12	Resveratrol affects Zika virus replication in vitro. <i>Scientific Reports</i> , 2019, 9, 14336.	1.6	48
13	Dengue Outbreak during Ongoing Civil War, Taiz, Yemen. <i>Emerging Infectious Diseases</i> , 2019, 25, 1397-1400.	2.0	17
14	A quantitative reverse transcription-polymerase chain reaction for detection of Getah virus. <i>Scientific Reports</i> , 2018, 8, 17632.	1.6	13
15	Sera of patients with systemic lupus erythematosus cross-neutralizes dengue viruses. <i>Microbiology and Immunology</i> , 2018, 62, 659-672.	0.7	4
16	Identification and characterization of <i>Corynebacterium lactis</i> isolated from <i>Amblyomma testudinarium</i> of <i>Sus scrofa</i> in Malaysia. <i>Systematic and Applied Acarology</i> , 2018, 23, 1838.	0.5	9
17	Operational utility of the reverse-transcription recombinase polymerase amplification for detection of dengue virus. <i>BMC Infectious Diseases</i> , 2018, 18, 169.	1.3	10
18	Emergence of the Asian lineage dengue virus type 3 genotype III in Malaysia. <i>BMC Evolutionary Biology</i> , 2018, 18, 58.	3.2	11

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19	Autochthonous spread of DENV-3 genotype III in Malaysia mitigated by pre-existing homotypic and heterotypic immunity. <i>Epidemiology and Infection</i> , 2018, 146, 1635-1641.	1.0	4
20	Detection in Malaysia of a <i>Borrelia</i> sp. From <i>Haemaphysalis hystricis</i> (Ixodida: Ixodidae). <i>Journal of Medical Entomology</i> , 2017, 54, 1444-1448.	0.9	23
21	Disruption of predicted dengue virus type 3 major outbreak cycle coincided with switching of the dominant circulating virus genotype. <i>Infection, Genetics and Evolution</i> , 2017, 54, 271-275.	1.0	14
22	Draft genome of <i>Bordetella pseudohinzii</i> BH370 isolated from trachea and lung tissues of a laboratory mouse. <i>Genomics Data</i> , 2017, 12, 69-70.	1.3	8
23	Vector competence of Malaysian <i>Aedes albopictus</i> with and without <i>Wolbachia</i> to four dengue virus serotypes. <i>Tropical Medicine and International Health</i> , 2017, 22, 1154-1165.	1.0	15
24	Draft genome of the emerging pathogen, <i>Kocuria marina</i> , isolated from a wild urban rat. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2017, 112, 857-859.	0.8	4
25	Geographical distribution of <i>Brucella melitensis</i> inferred from <i>rpoB</i> gene variation. <i>Journal of Infection in Developing Countries</i> , 2017, 11, 420-425.	0.5	4
26	The Use of NS1 Rapid Diagnostic Test and qRT-PCR to Complement IgM ELISA for Improved Dengue Diagnosis from Single Specimen. <i>Scientific Reports</i> , 2016, 6, 27663.	1.6	31
27	Bacterial community in <i>Haemaphysalis</i> ticks of domesticated animals from the Orang Asli communities in Malaysia. <i>Ticks and Tick-borne Diseases</i> , 2016, 7, 929-937.	1.1	62
28	Independent Emergence of the Cosmopolitan Asian Chikungunya Virus, Philippines 2012. <i>Scientific Reports</i> , 2015, 5, 12279.	1.6	22
29	Dengue death with evidence of hemophagocytic syndrome and dengue virus infection in the bone marrow. <i>SpringerPlus</i> , 2015, 4, 665.	1.2	9
30	Full genome SNP-based phylogenetic analysis reveals the origin and global spread of <i>Brucella melitensis</i> . <i>BMC Genomics</i> , 2015, 16, 93.	1.2	72
31	Early Detection of Dengue Virus by Use of Reverse Transcription-Recombinase Polymerase Amplification. <i>Journal of Clinical Microbiology</i> , 2015, 53, 830-837.	1.8	87
32	Colorimetric Detection of Dengue by Single Tube Reverse-Transcription-Loop-Mediated Isothermal Amplification. <i>PLoS ONE</i> , 2015, 10, e0138694.	1.1	61
33	Extract of <i>Scutellaria baicalensis</i> inhibits dengue virus replication. <i>BMC Complementary and Alternative Medicine</i> , 2013, 13, 91.	3.7	60
34	Detection of dengue viruses using reverse transcription-loop-mediated isothermal amplification. <i>BMC Infectious Diseases</i> , 2013, 13, 387.	1.3	84
35	Dengue virus type 1 clade replacement in recurring homotypic outbreaks. <i>BMC Evolutionary Biology</i> , 2013, 13, 213.	3.2	53
36	Emergence of dengue virus genotype 3 subgenotype III in Malaysia. <i>International Journal of Infectious Diseases</i> , 2012, 16, e272.	1.5	1

#	ARTICLE	IF	CITATIONS
37	Purifying selection in the evolution of dengue virus type 1 in Malaysia. International Journal of Infectious Diseases, 2012, 16, e273.	1.5	0
38	Differential osteogenic activity of osteoprogenitor cells on HA and TCP/HA scaffold of tissue engineered bone. Journal of Biomedical Materials Research - Part A, 2008, 85A, 301-312.	2.1	77
39	Bone graft substitute using hydroxyapatite scaffold seeded with tissue engineered autologous osteoprogenitor cells in spinal fusion: early result in a sheep model. Medical Journal of Malaysia, 2005, 60 Suppl C, 53-8.	0.2	11
40	Strategy for generating tissue-engineered human bone construct. Medical Journal of Malaysia, 2004, 59 Suppl B, 43-4.	0.2	2