## Kim-Kee Tan

List of Publications by Year in descending order

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477173 623574 40 871 14 29 h-index citations g-index papers 42 42 42 1583 docs citations citing authors all docs times ranked

#	Article	IF	CITATIONS
1	Early Detection of Dengue Virus by Use of Reverse Transcription-Recombinase Polymerase Amplification. Journal of Clinical Microbiology, 2015, 53, 830-837.	1.8	87
2	Detection of dengue viruses using reverse transcription-loop-mediated isothermal amplification. BMC Infectious Diseases, 2013, 13, 387.	1.3	84
3	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
4	Full genome SNP-based phylogenetic analysis reveals the origin and global spread of Brucella melitensis. BMC Genomics, 2015, 16, 93.	1.2	72
5	Bacterial community in Haemaphysalis ticks of domesticated animals from the Orang Asli communities in Malaysia. Ticks and Tick-borne Diseases, 2016, 7, 929-937.	1.1	62
6	Colorimetric Detection of Dengue by Single Tube Reverse-Transcription-Loop-Mediated Isothermal Amplification. PLoS ONE, 2015, 10, e0138694.	1.1	61
7	Extract of Scutellaria baicalensis inhibits dengue virus replication. BMC Complementary and Alternative Medicine, 2013, 13, 91.	3.7	60
8	Dengue virus type 1 clade replacement in recurring homotypic outbreaks. BMC Evolutionary Biology, 2013, 13, 213.	3.2	53
9	Resveratrol affects Zika virus replication in vitro. Scientific Reports, 2019, 9, 14336.	1.6	48
10	The Use of NS1 Rapid Diagnostic Test and qRT-PCR to Complement IgM ELISA for Improved Dengue Diagnosis from Single Specimen. Scientific Reports, 2016, 6, 27663.	1.6	31
11	Detection in Malaysia of a Borrelia sp. From Haemaphysalis hystricis (Ixodida: Ixodidae). Journal of Medical Entomology, 2017, 54, 1444-1448.	0.9	23
12	Independent Emergence of the Cosmopolitan Asian Chikungunya Virus, Philippines 2012. Scientific Reports, 2015, 5, 12279.	1.6	22
13	Bacterial communities in Haemaphysalis, Dermacentor and Amblyomma ticks collected from wild boar of an Orang Asli Community in Malaysia. Ticks and Tick-borne Diseases, 2020, 11, 101352.	1.1	19
14	Dengue Outbreak during Ongoing Civil War, Taiz, Yemen. Emerging Infectious Diseases, 2019, 25, 1397-1400.	2.0	17
15	Vector competence of Malaysian <i>Aedes albopictus</i> with and without <i>Wolbachia</i> to four dengue virus serotypes. Tropical Medicine and International Health, 2017, 22, 1154-1165.	1.0	15
16	Disruption of predicted dengue virus type 3 major outbreak cycle coincided with switching of the dominant circulating virus genotype. Infection, Genetics and Evolution, 2017, 54, 271-275.	1.0	14
17	A quantitative reverse transcription-polymerase chain reaction for detection of Getah virus. Scientific Reports, 2018, 8, 17632.	1.6	13
18	Emergence of B.1.524(G) SARS-CoV-2 in Malaysia during the third COVID-19 epidemic wave. Scientific Reports, 2021, 11, 22105.	1.6	13

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19	Emergence of the Asian lineage dengue virus type 3 genotype III in Malaysia. BMC Evolutionary Biology, 2018, 18, 58.	3.2	11
20	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
21	Operational utility of the reverse-transcription recombinase polymerase amplification for detection of dengue virus. BMC Infectious Diseases, 2018, 18, 169.	1.3	10
22	Dengue death with evidence of hemophagocytic syndrome and dengue virus infection in the bone marrow. SpringerPlus, 2015, 4, 665.	1.2	9
23	Identification and characterization of Corynebacterium lactis isolated from Amblyomma testudinarium of Sus scrofa in Malaysia. Systematic and Applied Acarology, 2018, 23, 1838.	0.5	9
24	Draft genome of Bordetella pseudohinzii BH370 isolated from trachea and lung tissues of a laboratory mouse. Genomics Data, 2017, 12, 69-70.	1.3	8
25	Serological evidence of DENV, JEV, and ZIKV among the indigenous people (Orang Asli) of Peninsular Malaysia. Journal of Medical Virology, 2020, 92, 956-962.	2.5	7
26	Draft genome of Paraburkholderia fungorum sequence type 868 recovered from human synovial tissues. Data in Brief, 2019, 25, 104159.	0.5	6
27	Draft genome of the emerging pathogen, Kocuria marina, isolated from a wild urban rat. Memorias Do Instituto Oswaldo Cruz, 2017, 112, 857-859.	0.8	4
28	Sera of patients with systemic lupus erythematosus crossâ€neutralizes dengue viruses. Microbiology and Immunology, 2018, 62, 659-672.	0.7	4
29	Autochthonous spread of DENV-3 genotype III in Malaysia mitigated by pre-existing homotypic and heterotypic immunity. Epidemiology and Infection, 2018, 146, 1635-1641.	1.0	4
30	Geographical distribution of Brucella melitensis inferred from rpoB gene variation. Journal of Infection in Developing Countries, 2017, 11, 420-425.	0.5	4
31	Multiplex sequencing of SARS-Cov-2 genome directly from clinical samples using the Ion Personal Genome Machine (PGM). Tropical Biomedicine, 2021, 38, 283-288.	0.2	3
32	Evaluation of commercial serological assays in Malaysia for detection of anti-Zika virus antibodies. Tropical Biomedicine, 2021, 38, 613-621.	0.2	3
33	A reverse transcription loop-mediated isothermal amplification for broad coverage detection of Asian and African Zika virus lineages. BMC Infectious Diseases, 2020, 20, 947.	1.3	2
34	Strategy for generating tissue-engineered human bone construct. Medical Journal of Malaysia, 2004, 59 Suppl B, 43-4.	0.2	2
35	Emergence of dengue virus genotype 3 subgenotype III in Malaysia. International Journal of Infectious Diseases, 2012, 16, e272.	1.5	1
36	Isolation of Kytococcus schroeteri from the brown rat Rattus norvegicus. Asian Pacific Journal of Tropical Medicine, 2021, 14, 191.	0.4	1

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#	Article	IF	CITATIONS
37	Continuous introduction, dispersal, and replacement of dengue virus type 3 genotype II sustained its transmission in Malaysia. International Journal of Infectious Diseases, 2020, 101, 236.	1.5	1
38	Purifying selection in the evolution of dengue virus type 1 in Malaysia. International Journal of Infectious Diseases, 2012, 16, e273.	1.5	0
39	Isolation of <i>Streptococcus cuniculi</i> from corneal lesion in laboratory-raised mice. Journal of Veterinary Medical Science, 2021, 83, 280-284.	0.3	O
40	Differential heterologous neutralisation profile against strains within DENV-3 genotype II. Epidemiology and Infection, 2022, 150, e33.	1.0	0