Chun Kiat Lee

List of Publications by Year in descending order

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567144 434063 1,035 39 15 31 citations h-index g-index papers 39 39 39 2165 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Covert COVID-19 and false-positive dengue serology in Singapore. Lancet Infectious Diseases, The, 2020, 20, 536.	4.6	256
2	A diagnostic polymerase chain reaction assay for Zika virus. Journal of Medical Virology, 2012, 84, 1501-1505.	2.5	167
3	An engineered CRISPR-Cas12a variant and DNA-RNA hybrid guides enable robust and rapid COVID-19 testing. Nature Communications, 2021, 12, 1739.	5.8	124
4	Differing Symptom Patterns in Early Pandemic vs Seasonal Influenza Infections. Archives of Internal Medicine, 2010, 170, 861.	4.3	43
5	Practical laboratory considerations amidst the COVID-19 outbreak: early experience from Singapore. Journal of Clinical Pathology, 2021, 74, 257-260.	1.0	41
6	Contamination-controlled high-throughput whole genome sequencing for influenza A viruses using the MiSeq sequencer. Scientific Reports, 2016, 6, 33318.	1.6	39
7	Four Patients with COVID-19 and Tuberculosis, Singapore, April–May 2020. Emerging Infectious Diseases, 2020, 26, 2763-2765.	2.0	36
8	Diagnostic Testing for Pandemic Influenza in Singapore. Journal of Molecular Diagnostics, 2010, 12, 636-643.	1.2	30
9	Distinguishing Zika and Dengue Viruses through Simple Clinical Assessment, Singapore. Emerging Infectious Diseases, 2018, 24, 1565-1568.	2.0	30
10	Atypical hand, foot, and mouth disease: eczema coxsackium can also occur in adults. Lancet Infectious Diseases, The, 2014, 14, 1043.	4.6	25
11	The Use of Polymerase Chain Reaction Amplification for the Detection of Viruses and Bacteria in Severe Community-Acquired Pneumonia. Respiration, 2016, 92, 286-294.	1.2	23
12	A Universal Influenza A and B Duplex Realâ€ŧime RTâ€PCR Assay. Journal of Medical Virology, 2012, 84, 1646-1651.	2.5	22
13	Persistent Dengue Infection in an Immunosuppressed Patient Reveals the Roles of Humoral and Cellular Immune Responses in Virus Clearance. Cell Host and Microbe, 2019, 26, 601-605.e3.	5.1	20
14	High-Resolution Melting Approach to Efficient Identification and Quantification of H275Y Mutant Influenza H1N1/2009 Virus in Mixed-Virus-Population Samples. Journal of Clinical Microbiology, 2011, 49, 3555-3559.	1.8	19
15	Correlating indoor and outdoor temperature and humidity in a sample of buildings in tropical climates. Indoor Air, 2021, 31, 2281-2295.	2.0	16
16	Comparison of Luminex NxTAG Respiratory Pathogen Panel and xTAG Respiratory Viral Panel FAST Version 2 for the Detection of Respiratory Viruses. Annals of Laboratory Medicine, 2017, 37, 267-271.	1.2	15
17	<i>CEBPA</i> mutational analysis in acute myeloid leukaemia by a laboratory-developed next-generation sequencing assay. Journal of Clinical Pathology, 2018, 71, 522-531.	1.0	15
18	Cost-benefit analysis of introducing next-generation sequencing (metagenomic) pathogen testing in the setting of pyrexia of unknown origin. PLoS ONE, 2018, 13, e0194648.	1.1	13

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19	Clinical performance of Roche cobas 6800, Luminex ARIES, MiRXES Fortitude Kit 2.1, Altona RealStar, and Applied Biosystems TaqPath for SARSâ€CoVâ€2 detection in nasopharyngeal swabs. Journal of Medical Virology, 2021, 93, 4603-4607.	2.5	13
20	False negative results caused by erroneous automated result interpretation algorithm on the FilmArray 2.0 instrument. Clinical Chemistry and Laboratory Medicine, 2018, 56, e43-e45.	1.4	10
21	Epidemiology of GB virus type C among patients infected with HIV in Singapore. Journal of Medical Virology, 2014, 86, 737-744.	2.5	8
22	Evaluation of the Luminex ARIES HSV 1&2 Assay and Comparison with the FTD Neuro 9 and In-house Real-Time PCR Assays for Detecting Herpes Simplex Viruses. Annals of Laboratory Medicine, 2018, 38, 440-445.	1.2	8
23	Detection of a Novel Single Nucleotide Polymorphism of the Human Thiopurine S-Methyltransferase Gene in a Chinese Individual. Drug Metabolism and Pharmacokinetics, 2012, 27, 559-561.	1.1	7
24	Range of Varicella Zoster Co-Infections with COVID-19, Singapore. Infection and Chemotherapy, 2021, 53, 391.	1.0	7
25	Design and Multicenter Clinical Validation of a 3-Dimensionally Printed Nasopharyngeal Swab for SARS-CoV-2 Testing. JAMA Otolaryngology - Head and Neck Surgery, 2021, 147, 418.	1.2	7
26	An inâ€house HIV genotyping assay for the detection of drug resistance mutations in Southeast Asian patients infected with HIVâ€1. Journal of Medical Virology, 2012, 84, 394-401.	2.5	5
27	Comparison of cobas HCV GT against Versant HCV Genotype 2.0 (LiPA) with confirmation by Sanger sequencing. Journal of Virological Methods, 2018, 255, 8-13.	1.0	5
28	Next generation sequencing identifies multi-drug resistant herpes simplex virus- associated scrotal ulceration. Journal of Infection, 2020, 80, 232-254.	1.7	5
29	Clinical Diagnostic Study of a Novel Injection Molded Swab for SARS-Cov-2 Testing. Infectious Diseases and Therapy, 2021, 10, 1015-1022.	1.8	5
30	Simple "Rule-of-6―Predicts Severe Coronavirus Disease 2019 (COVID-19). Clinical Infectious Diseases, 2021, 72, 1861-1862.	2.9	4
31	Parallel testing of 241 clinical nasopharyngeal swabs for the detection of SARS-CoV-2 virus on the Cepheid Xpert Xpress SARS-CoV-2 and the Roche cobas SARS-CoV-2 assays. Clinical Chemistry and Laboratory Medicine, 2021, 59, e45-e48.	1.4	4
32	Uncovering the hidden villain within the human respiratory microbiome. Diagnosis, 2014, 1, 203-212.	1.2	3
33	Complete Genome Sequence of an ON1 Human Respiratory Syncytial Virus Strain Isolated in Lebanon in 2015. Genome Announcements, 2018, 6, .	0.8	2
34	Performance evaluation of Cepheid Xpert Norovirus kit with a userâ€modified protocol. Journal of Medical Virology, 2018, 90, 485-489.	2.5	2
35	Can we do better? A guide to pandemics – some Dos and Don'ts for the next one. Journal of Infection, 2021, 83, 119-145.	1.7	2
36	Effects of acute SARS-CoV-2 infection on male hormone profile, ACE2 and TMPRSS2 expression and potential for transmission of SARS-CoV-2 in semen of Asian men. F&S Science, 2021, , .	0.5	2

#	Article	IF	CITATIONS
37	Missed detection of an avian influenza A (H7N9) virus by the Luminex xTAG respiratory viral panel FAST version 2. Pathology, 2017, 49, 330-332.	0.3	1
38	Validation of a multi-gene qPCR-based pharmacogenomics panel across major ethnic groups in Singapore and Indonesia. Pharmacogenomics, 2021, 22, 1041-1056.	0.6	1
39	Lack of oseltamivir-resistance in A/H1N1p-infected patients, Singapore. Journal of Infection, 2012, 65, 93-95.	1.7	O