Jee-Soo Lee

List of Publications by Year
in descending order

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23 papers	283 citations	1307594 7 h-index	996975 15 g-index
24 all docs	24 docs citations	24 times ranked	729 citing authors

#	Article	IF	Citations
1	In-depth blood proteome profiling analysis revealed distinct functional characteristics of plasma proteins between severe and non-severe COVID-19 patients. Scientific Reports, 2020, 10, 22418.	3.3	80
2	Evidence of Severe Acute Respiratory Syndrome Coronavirus 2 Reinfection After Recovery from Mild Coronavirus Disease 2019. Clinical Infectious Diseases, 2021, 73, e3002-e3008.	5.8	68
3	Longitudinal proteomic profiling provides insights into host response and proteome dynamics in COVIDâ€19 progression. Proteomics, 2021, 21, e2000278.	2.2	26
4	Pitfalls of Multiple Ligation-Dependent Probe Amplifications in Detecting DMD Exon Deletions or Duplications. Journal of Molecular Diagnostics, 2016, 18, 253-259.	2.8	23
5	Molecular basis and diagnosis of thalassemia. Blood Research, 2021, 56, S39-S43.	1.3	20
6	Large Deletions of TSPAN12 Cause Familial Exudative Vitreoretinopathy (FEVR). , 2016, 57, 6902.		11
7	Evaluation of a Real-Time Reverse Transcription-PCR (RT-PCR) Assay for Detection of Middle East Respiratory Syndrome Coronavirus (MERS-CoV) in Clinical Samples from an Outbreak in South Korea in 2015. Journal of Clinical Microbiology, 2017, 55, 2554-2555.	3.9	9
8	SnackVar. Journal of Molecular Diagnostics, 2021, 23, 140-148.	2.8	7
9	Rates of Coinfection Between SARS-CoV-2 and Other Respiratory Viruses in Korea. Annals of Laboratory Medicine, 2022, 42, 110-112.	2.5	7
10	Noninvasive prenatal test of single-gene disorders by linked-read direct haplotyping: application in various diseases. European Journal of Human Genetics, 2021, 29, 463-470.	2.8	5
11	Consistent count regionâ€“copy number variation (CCR-CNV): an expandable and robust tool for clinical diagnosis of copy number variation at the exon level using next-generation sequencing data. Genetics in Medicine, 2022, 24, 663-672.	2.4	5
12	Epidemiologic Trends of Thalassemia, 2006â€“2018: A Nationwide Population-Based Study. Journal of Clinical Medicine, 2022, 11, 2289.	2.4	5
13	Evaluation of the new Abbott Real-Time EBV assay: fully automated quantification of EBV in whole blood by targeting BLLF1. Diagnostic Microbiology and Infectious Disease, 2019, 94, 135-139.	1.8	3
14	Determination of Clinical Characteristics of Mycobacterium kansasii-Derived Species by Reanalysis of Isolates Formerly Reported as M. kansasii. Annals of Laboratory Medicine, 2021, 41, 463-468.	2.5	3
15	SnackNTM: An Open-Source Software for Sanger Sequencing-based Identification of Nontuberculous Mycobacterial Species. Annals of Laboratory Medicine, 2022, 42, 213-248.	2.5	3
16	Intronic LINE-1 insertion in SLCO1B3 as a highly prevalent cause of rotor syndrome in East Asian population. Journal of Human Genetics, 2022, 67, 71-77.	2.3	2
17	Evaluation of the AccuPowerÂ® RV1 Real-Time RT-PCR Kit and the AccuPowerÂ® RV1 Multiplex Kit for SARS-CoV-2 and Influenza Virus Detection. Laboratory Medicine Online, 2021, 11, 290-296.	0.2	2
18	Immunohistochemical Staining to Identify Concomitant Systemic Mastocytosis in Acute Myeloid Leukemia with RUNX1::RUNX1T1. Annals of Laboratory Medicine, 2022, 42, 678-682.	2.5	2

#	Article	IF	Citations
19	No association between POU4F1, POU4F2, ISL1 polymorphisms and normal-tension glaucoma. Ophthalmic Genetics, 2020, 41, 427-431.	1.2	1
20	FMS-like Tyrosine Kinase 3-Internal Tandem Duplication Allele Concentrations Should Be Determined in a Mutation-Type-Specific Manner. Clinical Chemistry, 2021, 67, 691-693.	3.2	0
21	The First Korean Case of Griscelli Syndrome Type 2 With Hemophagocytic Lymphohistiocytosis and Partial Albinism. Annals of Laboratory Medicine, 2022, 42, 384-388.	2.5	0
22	Comparison of Respiratory Specimens for the Detection of SARS-CoV-2. Annals of Clinical and Laboratory Science, 2021, 51, 140-144.	0.2	0
23	Report on the External Quality Assessment Scheme for Genetic Disorders and Other Human Genetics Molecular Diagnostics in Korea (2018â€“2021). , 2022, 44, 61-75.		0