Jihoon G Yoon

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4503080/publications.pdf

Version: 2024-02-01

		1163117	996975
16	521	8	15
papers	citations	h-index	g-index
16	16	16	796
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Machine Learning–Based Model for Prediction of Outcomes in Acute Stroke. Stroke, 2019, 50, 1263-1265.	2.0	323
2	Molecular Characterization of Biliary Tract Cancer Predicts Chemotherapy and Programmed Death 1/Programmed Deathâ€Ligand 1 Blockade Responses. Hepatology, 2021, 74, 1914-1931.	7.3	48
3	Comparison of MALDI-TOF MS, Housekeeping Gene Sequencing, and 16S rRNA Gene Sequencing for Identification of <i>Aeromonas </i> Clinical Isolates. Yonsei Medical Journal, 2015, 56, 550.	2.2	39
4	Regulation of CFTR Bicarbonate Channel Activity by WNK1: Implications for Pancreatitis and CFTR-Related Disorders. Cellular and Molecular Gastroenterology and Hepatology, 2020, 9, 79-103.	4.5	27
5	Molecular Diagnosis of Craniosynostosis Using Targeted Next-Generation Sequencing. Neurosurgery, 2020, 87, 294-302.	1.1	20
6	Comparison of Six Automated Treponema-Specific Antibody Assays. Journal of Clinical Microbiology, 2016, 54, 163-167.	3.9	16
7	Machine learning-based diagnosis for disseminated intravascular coagulation (DIC): Development, external validation, and comparison to scoring systems. PLoS ONE, 2018, 13, e0195861.	2.5	11
8	A pilot study to investigate the utility of NAT2 genotype-guided isoniazid monotherapy regimens in NAT2 slow acetylators. Pharmacogenetics and Genomics, 2021, 31, 68-73.	1.5	10
9	Magnetic bead-based nucleic acid purification kit: Clinical application and performance evaluation in stool specimens. Journal of Microbiological Methods, 2016, 124, 62-68.	1.6	7
10	Unraveling the Genomic Architecture of the CYP3A Locus and ADME Genes for Personalized Tacrolimus Dosing. Transplantation, 2021, 105, 2213-2225.	1.0	6
11	Functional Characterization of Pharmcogenetic Variants of Human Cytochrome P450 2C9 in Korean Populations. Biomolecules and Therapeutics, 2019, 27, 577-583.	2.4	4
12	An 18.3-Mb Duplication on Chromosome 14q With Multiple Cardiac Anomalies and Clubfoot Was Identified by Microarray Analysis. Annals of Laboratory Medicine, 2016, 36, 194-196.	2.5	3
13	Automated CH50 liposome-based immunoassay: consideration in dilution and validation of reference interval. Clinical Chemistry and Laboratory Medicine, 2016, 54, e309-12.	2.3	3
14	Adopting automated image analysis tool for fibrin network: Can we obtain clot properties for practical application?. International Journal of Laboratory Hematology, 2017, 39, e121-e123.	1.3	3
15	Bacteroides nordii and Bacteroides salyersiae Isolated from Post-operative Peritonitis Patients. Laboratory Medicine Online, 2016, 6, 111.	0.2	1
16	Targeted next-generation sequencing to identify genetic polymorphism associated with levetiracetam-induced psychosis. Proceedings for Annual Meeting of the Japanese Pharmacological Society, 2018, WCP2018, OR11-2.	0.0	0