

Guigao Lin

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

117
citations

1684188

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1281871

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docs citations

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times ranked

203
citing authors

#	ARTICLE	IF	CITATIONS
1	Technical Validation of a Next-Generation Sequencing Assay for Detecting Clinically Relevant Levels of Breast Cancer-Related Single-Nucleotide Variants and Copy Number Variants Using Simulated Cell-Free DNA. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 525-536.	2.8	64
2	Hepatitis C Virus RNA Real-Time Quantitative RT-PCR Method Based on a New Primer Design Strategy. <i>Journal of Molecular Diagnostics</i> , 2016, 18, 84-91.	2.8	10
3	A novel cell line generated using the CRISPR/Cas9 technology as universal quality control material for KRAS G12V mutation testing. <i>Journal of Clinical Laboratory Analysis</i> , 2018, 32, e22391.	2.1	9
4	CRISPR/Cas9 Technology-Based Xenograft Tumors as Candidate Reference Materials for Multiple EML4-ALK Rearrangements Testing. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 766-775.	2.8	8
5	Improvements in CYP2C9 Genotyping Accuracy Are Needed. <i>Journal of Cardiovascular Pharmacology</i> , 2015, 66, 129-134.	1.9	6
6	Clustered Regularly Interspaced Short Palindromic Repeat (CRISPR)/CRISPR-Associated Endonuclease Cas9-Mediated Homology-Independent Integration for Generating Quality Control Materials for Clinical Molecular Genetic Testing. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 373-380.	2.8	6
7	Evaluation of tacrolimus-related CYP3A5 genotyping in China: Results from the First External Quality Assessment Exercise. <i>Journal of Clinical Laboratory Analysis</i> , 2018, 32, e22563.	2.1	3
8	A national proficiency scheme for human leucocyte antigen typing by next-generation sequencing. <i>Clinica Chimica Acta</i> , 2022, 533, 85-88.	1.1	3
9	An Initial Survey of the Performances of Exome Variant Analysis and Clinical Reporting Among Diagnostic Laboratories in China. <i>Frontiers in Genetics</i> , 2020, 11, 582637.	2.3	2
10	Preparation of multiplexed control materials for cancer mutation analysis by genome editing in GM12878 cells. <i>Journal of Clinical Laboratory Analysis</i> , 2022, 36, e24139.	2.1	2
11	Adaptation of ACMG-ClinGen Technical Standards for Copy Number Variant Interpretation Concordance. <i>Frontiers in Genetics</i> , 2022, 13, 829728.	2.3	2
12	External quality assessment for laboratory testing of HLA-B*15:02 allele in relation to carbamazepine therapy. <i>Journal of Clinical Laboratory Analysis</i> , 2018, 32, .	2.1	1
13	Development of a Novel Reference Material for Tumor Mutational Burden Measurement Based on CRISPR/Cas9 Technology. <i>Frontiers in Oncology</i> , 2022, 12, 845636.	2.8	1