Timothy Devin Minogue

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparison of Illumina and Oxford Nanopore Sequencing Technologies for Pathogen Detection from Clinical Matrices Using Molecular Inversion Probes. Journal of Molecular Diagnostics, 2022, 24, 395-405.	2.8	11
2	Development of a coronavirus disease 2019 nonhuman primate model using airborne exposure. PLoS ONE, 2021, 16, e0246366.	2.5	52
3	Transcriptomic Analysis Reveals Host miRNAs Correlated with Immune Gene Dysregulation during Fatal Disease Progression in the Ebola Virus Cynomolgus Macaque Disease Model. Microorganisms, 2021, 9, 665.	3.6	4
4	Comparison of transcriptional responses between pathogenic and nonpathogenic hantavirus infections in Syrian hamsters using NanoString. PLoS Neglected Tropical Diseases, 2021, 15, e0009592.	3.0	4
5	Host response transcriptomic analysis of Crimean-Congo hemorrhagic fever pathogenesis in the cynomolgus macaque model. Scientific Reports, 2021, 11, 19807.	3.3	6
6	Modeling the stability of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) on skin, currency, and clothing. PLoS Neglected Tropical Diseases, 2020, 14, e0008831.	3.0	109
7	FDA-ARGOS is a database with public quality-controlled reference genomes for diagnostic use and regulatory science. Nature Communications, 2019, 10, 3313.	12.8	101
8	2018 Ebola virus disease outbreak in Équateur Province, Democratic Republic of the Congo: a retrospective genomic characterisation. Lancet Infectious Diseases, The, 2019, 19, 641-647.	9.1	27
9	Next-Generation Sequencing for Biodefense: Biothreat Detection, Forensics, and the Clinic. Clinical Chemistry, 2019, 65, 383-392.	3.2	23
10	Diagnostic targETEd seQuencing adjudicaTion (DETEQT). Journal of Molecular Diagnostics, 2019, 21, 99-110.	2.8	5
11	Magnetic Nanotrap Particles Preserve the Stability of Venezuelan Equine Encephalitis Virus in Blood for Laboratory Detection. Frontiers in Veterinary Science, 2019, 6, 509.	2.2	12
12	Detection of 16S rRNA and KPC Genes from Complex Matrix Utilizing a Molecular Inversion Probe Assay for Next-Generation Sequencing. Scientific Reports, 2018, 8, 2028.	3.3	9
13	Virus-encoded miRNAs in Ebola virus disease. Scientific Reports, 2018, 8, 6480.	3.3	34
14	A conserved transcriptional response to intranasal Ebola virus exposure in nonhuman primates prior to onset of fever. Science Translational Medicine, 2018, 10, .	12.4	25
15	A highly multiplexed broad pathogen detection assay for infectious disease diagnostics. PLoS Neglected Tropical Diseases, 2018, 12, e0006889.	3.0	23
16	Rapid antibiotic susceptibility testing from blood culture bottles with species agnostic real-time polymerase chain reaction. PLoS ONE, 2018, 13, e0209042.	2.5	10
17	Sequence Optimized Real-Time Reverse Transcription Polymerase Chain Reaction Assay for Detection of Crimean-Congo Hemorrhagic Fever Virus. American Journal of Tropical Medicine and Hygiene, 2018, 98, 211-215.	1.4	18
18	African and Asian Zika Virus Isolates Display Phenotypic Differences Both In Vitro and In Vivo. American Journal of Tropical Medicine and Hygiene, 2018, 98, 432-444.	1.4	65

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19	Targeted Next-Generation Sequencing for Diagnostics and Forensics. Clinical Chemistry, 2017, 63, 450-452.	3.2	6
20	Inactivation of West Nile virus in serum with heat, ionic detergent, and reducing agent for proteomic applications. Journal of Virological Methods, 2017, 248, 1-6.	2.1	3
21	Comparison of Transcriptomic Platforms for Analysis of Whole Blood from Ebola-Infected Cynomolgus Macaques. Scientific Reports, 2017, 7, 14756.	3.3	32
22	Semi-quantitative MALDI-TOF for antimicrobial susceptibility testing in Staphylococcus aureus. PLoS ONE, 2017, 12, e0183899.	2.5	26
23	Circulating microRNA profiles of Ebola virus infection. Scientific Reports, 2016, 6, 24496.	3.3	50
24	Targeted next-generation sequencing for the detection of ciprofloxacin resistance markers using molecular inversion probes. Scientific Reports, 2016, 6, 25904.	3.3	32
25	Development of realâ€ŧime PCR assays for the detection of Moraxella macacae associated with bloody nose syndrome in rhesus (Macaca mulatta) and cynomolgus (Macaca fascicularis) macaques. Journal of Medical Primatology, 2015, 44, 364-372.	0.6	4
26	Evaluation of Signature Erosion in Ebola Virus Due to Genomic Drift and Its Impact on the Performance of Diagnostic Assays. Viruses, 2015, 7, 3130-3154.	3.3	35
27	Optimized microRNA purification from TRIzol-treated plasma. BMC Genomics, 2015, 16, 95.	2.8	43
28	Demonstration of the Pre–Emergency Use Authorization Path Using 3 Minor Groove Binder–Hydrolysis Probe Assays to Detect Escherichia coli O104:H4. Clinical Chemistry, 2015, 61, 1391-1398.	3.2	2
29	Real-time reverse transcriptase polymerase chain reaction assays for Middle East Respiratory Syndrome. Molecular and Cellular Probes, 2015, 29, 511-513.	2.1	6
30	Development and Evaluation of a Panel of Filovirus Sequence Capture Probes for Pathogen Detection by Next-Generation Sequencing. PLoS ONE, 2014, 9, e107007.	2.5	28
31	Development of real-time PCR assays for specific detection of hmsH, hmsF, hmsR, and irp2 located within the 102-kb pgm locus of Yersinia pestis. Molecular and Cellular Probes, 2014, 28, 288-295.	2.1	7
32	Cross-Institute Evaluations of Inhibitor-Resistant PCR Reagents for Direct Testing of Aerosol and Blood Samples Containing Biological Warfare Agent DNA. Applied and Environmental Microbiology, 2014, 80, 1322-1329.	3.1	16
33	Stabilization of biothreat diagnostic samples through vitrification matrices. Journal of Microbiological Methods, 2014, 101, 81-85.	1.6	2