

Darrell O Ricke

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

312
citations

1651377

6
h-index

1427216

11
g-index

47
all docs

47
docs citations

47
times ranked

349
citing authors

#	ARTICLE	IF	CITATIONS
1	Two Different Antibody-Dependent Enhancement (ADE) Risks for SARS-CoV-2 Antibodies. <i>Frontiers in Immunology</i> , 2021, 12, 640093.	2.2	93
2	COVID-19: Famotidine, Histamine, Mast Cells, and Mechanisms. <i>Frontiers in Pharmacology</i> , 2021, 12, 633680.	1.6	64
3	Hyperactivated Mast Cells Pathogenesis Hypothesis for COVID-19 Cutaneous Manifestations. <i>JID Innovations</i> , 2021, 1, 100052.	1.2	3
4	Models for COVID-19 Early Cardiac Pathology Following SARS-CoV-2 Infection. <i>International Journal of Infectious Diseases</i> , 2021, 113, 331-335.	1.5	2
5	Models for SARS-CoV-2 associated thrombocytopenia associated with hemophagocytic histiocytes. <i>Medical Hypotheses</i> , 2021, 157, 110700.	0.8	1
6	Analysis of complex DNA mixtures using massively parallel sequencing of SNPs with low minor allele frequencies. <i>Forensic Science International: Genetics</i> , 2020, 46, 102234.	1.6	15
7	Kawasaki Disease, Multisystem Inflammatory Syndrome in Children: Antibody-Induced Mast Cell Activation Hypothesis. <i>Journal of Pediatrics and Pediatric Medicine</i> , 2020, 4, 1-7.	0.9	9
8	Letter to the Editor "Detecting HTS Barcode Contamination. <i>Journal of Forensic Sciences</i> , 2019, 64, 961-962.	0.9	0
9	Estimating Individual Contributions to Complex DNA SNP Mixtures. <i>Journal of Forensic Sciences</i> , 2019, 64, 1468-1474.	0.9	8
10	IdPrism: Rapid Analysis of Forensic DNA Samples Using MPS SNP Profiles. , 2019, , .		1
11	Grigora<scp>SNP</scp>s: Optimized Analysis of <scp>SNP</scp>s for <scp>DNA</scp> Forensics. <i>Journal of Forensic Sciences</i> , 2018, 63, 1841-1845.	0.9	6
12	FastID: Extremely fast forensic DNA comparisons. , 2017, , .		5
13	KinLinks: Software Toolkit for kinship analysis and pedigree generation from HTS datasets. , 2016, , .		9
14	Human CODIS STR loci profiling from HTS data. , 2016, , .		4
15	Dawn: Rapid large-scale protein multiple sequence alignment and conservation analysis. , 2015, , .		2
16	Robust detection of individual forensic profiles in DNA mixtures. <i>Forensic Science International: Genetics</i> , 2015, 14, 31-37.	1.6	40
17	Nonrandom patterns of simple and cryptic triplet repeats in coding and noncoding sequences. <i>Genomics</i> , 1995, 26, 510-520.	1.3	19
18	TranslucentID: analysis of complex DNA SNP mixtures with large numbers of donors. <i>Australian Journal of Forensic Sciences</i> , 0, , 1-12.	0.7	3

#	ARTICLE	IF	CITATIONS
19	Fast P(RMNE): Fast forensic DNA probability of random man not excluded calculation. F1000Research, 0, 6, 2154.	0.8	4
20	Vasomotor symptoms monitoring with a commercial activity tracking watch. F1000Research, 0, 6, 2155.	0.8	0
21	Integrated Biomedical System. F1000Research, 0, 7, 162.	0.8	0
22	Fast P(RMNE): Fast forensic DNA probability of random man not excluded calculation. F1000Research, 0, 6, 2154.	0.8	1
23	Etiology Models of Antibody Triggered Histamine Intolerance Inducing Kawasaki Disease and Multisystem Inflammatory Syndromes Diseases. Qeios, 0, , .	0.0	0