Kenneth S Plante

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9578492/publications.pdf

Version: 2024-02-01

39 papers 5,860 citations

304602 22 h-index 289141 40 g-index

54 all docs

54 docs citations

54 times ranked 10959 citing authors

#	Article	IF	CITATIONS
1	Spike mutation D614G alters SARS-CoV-2 fitness. Nature, 2021, 592, 116-121.	13.7	1,380
2	An Infectious cDNA Clone of SARS-CoV-2. Cell Host and Microbe, 2020, 27, 841-848.e3.	5.1	617
3	Loss of furin cleavage site attenuates SARS-CoV-2 pathogenesis. Nature, 2021, 591, 293-299.	13.7	579
4	Severe Acute Respiratory Syndrome Coronavirus 2 from Patient with Coronavirus Disease, United States. Emerging Infectious Diseases, 2020, 26, 1266-1273.	2.0	523
5	The N501Y spike substitution enhances SARS-CoV-2 infection and transmission. Nature, 2022, 602, 294-299.	13.7	364
6	The variant gambit: COVID-19's next move. Cell Host and Microbe, 2021, 29, 508-515.	5.1	305
7	Adjuvanting a subunit COVID-19 vaccine to induce protective immunity. Nature, 2021, 594, 253-258.	13.7	253
8	Multi-peaked adaptive landscape for chikungunya virus evolution predicts continued fitness optimization in Aedes albopictus mosquitoes. Nature Communications, 2014, 5, 4084.	5.8	179
9	Novel Chikungunya Vaccine Candidate with an IRES-Based Attenuation and Host Range Alteration Mechanism. PLoS Pathogens, 2011, 7, e1002142.	2.1	148
10	Attenuation of Chikungunya Virus Vaccine Strain 181/Clone 25 Is Determined by Two Amino Acid Substitutions in the E2 Envelope Glycoprotein. Journal of Virology, 2012, 86, 6084-6096.	1.5	142
11	Acute Respiratory Distress in Aged, SARS-CoV-2–Infected African Green Monkeys but Not Rhesus Macaques. American Journal of Pathology, 2021, 191, 274-282.	1.9	123
12	Chikungunya Vaccine Candidate Is Highly Attenuated and Protects Nonhuman Primates Against Telemetrically Monitored Disease Following a Single Dose. Journal of Infectious Diseases, 2014, 209, 1891-1899.	1.9	86
13	Nucleocapsid mutations in SARS-CoV-2 augment replication and pathogenesis. PLoS Pathogens, 2022, 18, e1010627.	2.1	85
14	Mouse-adapted SARS-CoV-2 protects animals from lethal SARS-CoV challenge. PLoS Biology, 2021, 19, e3001284.	2.6	54
15	A trans-complementation system for SARS-CoV-2 recapitulates authentic viral replication without virulence. Cell, 2021, 184, 2229-2238.e13.	13.5	51
16	VLDLR and ApoER2 are receptors for multiple alphaviruses. Nature, 2022, 602, 475-480.	13.7	49
17	Structural divergence creates new functional features in alphavirus genomes. Nucleic Acids Research, 2018, 46, 3657-3670.	6.5	45
18	Extended Preclinical Safety, Efficacy and Stability Testing of a Live-attenuated Chikungunya Vaccine Candidate. PLoS Neglected Tropical Diseases, 2015, 9, e0004007.	1.3	39

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19	The Role of Innate versus Adaptive Immune Responses in a Mouse Model of O'Nyong-Nyong Virus Infection. American Journal of Tropical Medicine and Hygiene, 2013, 88, 1170-1179.	0.6	37
20	Effects of Chikungunya virus immunity on Mayaro virus disease and epidemic potential. Scientific Reports, 2019, 9, 20399.	1.6	35
21	Complex Genetic Architecture Underlies Regulation of Influenza-A-Virus-Specific Antibody Responses in the Collaborative Cross. Cell Reports, 2020, 31, 107587.	2.9	31
22	Role of mutational reversions and fitness restoration in Zika virus spread to the Americas. Nature Communications, 2021, 12, 595.	5.8	29
23	Bayesian Diallel Analysis Reveals <i>Mx1</i> -Dependent and <i>Mx1</i> -Independent Effects on Response to Influenza A Virus in Mice. G3: Genes, Genomes, Genetics, 2018, 8, 427-445.	0.8	27
24	Antiviral activity of oleandrin and a defined extract of Nerium oleander against SARS-CoV-2. Biomedicine and Pharmacotherapy, 2021, 138, 111457.	2.5	23
25	Using SHAPE-MaP To Model RNA Secondary Structure and Identify 3′UTR Variation in Chikungunya Virus. Journal of Virology, 2020, 94, .	1.5	22
26	Tiled-ClickSeq for targeted sequencing of complete coronavirus genomes with simultaneous capture of RNA recombination and minority variants. ELife, 2021, 10, .	2.8	22
27	Stability of Yellow Fever Virus under Recombinatory Pressure as Compared with Chikungunya Virus. PLoS ONE, 2011, 6, e23247.	1.1	21
28	Bunyavirus Taxonomy: Limitations and Misconceptions Associated with the Current ICTV Criteria Used for Species Demarcation. American Journal of Tropical Medicine and Hygiene, 2018, 99, 11-16.	0.6	21
29	High Seroprevalence of Dengue Virus Infection in Sudan: Systematic Review and Meta-Analysis. Tropical Medicine and Infectious Disease, 2020, 5, 120.	0.9	20
30	Venezuelan equine encephalitis vaccine with rearranged genome resists reversion and protects non-human primates from viremia after aerosol challenge. Vaccine, 2020, 38, 3378-3386.	1.7	18
31	The pigtail macaque (Macaca nemestrina) model of COVID-19 reproduces diverse clinical outcomes and reveals new and complex signatures of disease. PLoS Pathogens, 2021, 17, e1010162.	2.1	11
32	Immune predictors of mortality following RNA virus infection. Journal of Infectious Diseases, 2020, 221, 882-889.	1.9	10
33	Optimized production and immunogenicity of an insect virus-based chikungunya virus candidate vaccine in cell culture and animal models. Emerging Microbes and Infections, 2021, 10, 305-316.	3.0	9
34	Lineage Divergence and Vector-Specific Adaptation Have Driven Chikungunya Virus onto Multiple Adaptive Landscapes. MBio, 2021, 12, e0273821.	1.8	8
35	Rationally Attenuated Vaccines for Venezuelan Equine Encephalitis Protect Against Epidemic Strains with a Single Dose. Vaccines, 2020, 8, 497.	2.1	6
36	Designing multivalent immunogens for alphavirus vaccine optimization. Virology, 2021, 561, 117-124.	1.1	3

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37	Outbreak of coronavirus disease 2019 (COVID-19) among operating room staff of a tertiary referral center: An epidemiologic and environmental investigation. Infection Control and Hospital Epidemiology, 2022, 43, 319-325.	1.0	2
38	Preclinical Evaluation of a Live Attenuated Chikungunya Vaccine. Procedia in Vaccinology, 2012, 6, 141-149.	0.4	1
39	Characterization of a Dengue Virus Serotype 1 Isolated from a Patient in Ciudad Juarez, Mexico. Pathogens, 2021, 10, 872.	1.2	1