Reed S Shabman

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8747248/publications.pdf

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39 papers 2,819 citations

28 h-index 315739 38 g-index

40 all docs

40 docs citations

40 times ranked

4281 citing authors

#	Article	IF	CITATIONS
1	Defining the risk of SARS-CoV-2 variants on immune protection. Nature, 2022, 605, 640-652.	27.8	117
2	Optimization of qRT-PCR assay for zika virus detection in human serum and urine. Virus Research, 2019, 263, 173-178.	2.2	17
3	Ross River virus envelope glycans contribute to disease through activation of the host complement system. Virology, 2018, 515, 250-260.	2.4	13
4	Whole-Genome Sequences of Zika Virus FLR Strains after Passage in Vero or C6/36 Cells. Genome Announcements, $2018, 6, .$	0.8	2
5	Analysis of the Aedes albopictus C6/36 genome provides insight into cell line utility for viral propagation. GigaScience, 2018, 7, 1-13.	6.4	51
6	Whole genome sequencing, variant analysis, phylogenetics, and deep sequencing of Zika virus strains. Scientific Reports, 2018, 8, 15843.	3.3	20
7	Sequences of Zika Virus Genomes from a Pediatric Cohort in Nicaragua. Genome Announcements, 2018, 6, .	0.8	O
8	Differing epidemiological dynamics of Chikungunya virus in the Americas during the 2014-2015 epidemic. PLoS Neglected Tropical Diseases, 2018, 12, e0006670.	3.0	23
9	Conservation of Structure and Immune Antagonist Functions of Filoviral VP35 Homologs Present in Microbat Genomes. Cell Reports, 2018, 24, 861-872.e6.	6.4	16
10	Genetic stability of foot-and-mouth disease virus during long-term infections in natural hosts. PLoS ONE, 2018, 13, e0190977.	2.5	8
11	IFN-λ4 Attenuates Antiviral Responses by Enhancing Negative Regulation of IFN Signaling. Journal of Immunology, 2017, 199, 3808-3820.	0.8	55
12	The Ebola virus VP35 protein binds viral immunostimulatory and host RNAs identified through deep sequencing. PLoS ONE, 2017, 12, e0178717.	2.5	17
13	Zika Virus Antagonizes Type I Interferon Responses during Infection of Human Dendritic Cells. PLoS Pathogens, 2017, 13, e1006164.	4.7	242
14	Isolation and Characterization of a Novel Gammaherpesvirus from a Microbat Cell Line. MSphere, 2016, 1 , .	2.9	16
15	Comprehensive Genome Scale Phylogenetic Study Provides New Insights on the Global Expansion of Chikungunya Virus. Journal of Virology, 2016, 90, 10600-10611.	3.4	72
16	A Universal Next-Generation Sequencing Protocol To Generate Noninfectious Barcoded cDNA Libraries from High-Containment RNA Viruses. MSystems, 2016, 1, .	3.8	28
17	Differential Regulation of Interferon Responses by Ebola and Marburg Virus VP35 Proteins. Cell Reports, 2016, 14, 1632-1640.	6.4	7 5
18	An Intrinsically Disordered Peptide from Ebola Virus VP35 Controls Viral RNA Synthesis by Modulating Nucleoprotein-RNA Interactions. Cell Reports, 2015, 11, 376-389.	6.4	136

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19	Characterization of Uncultivable Bat Influenza Virus Using a Replicative Synthetic Virus. PLoS Pathogens, 2014, 10, e1004420.	4.7	58
20	The Marburg Virus VP24 Protein Interacts with Keap1 to Activate the Cytoprotective Antioxidant Response Pathway. Cell Reports, 2014, 6, 1017-1025.	6.4	95
21	Deep Sequencing Identifies Noncanonical Editing of Ebola and Marburg Virus RNAs in Infected Cells. MBio, 2014, 5, e02011.	4.1	70
22	In Silico Derived Small Molecules Bind the Filovirus VP35 Protein and Inhibit Its Polymerase Cofactor Activity. Journal of Molecular Biology, 2014, 426, 2045-2058.	4.2	75
23	Development of RNA Aptamers Targeting Ebola Virus VP35. Biochemistry, 2013, 52, 8406-8419.	2.5	73
24	Ebola Virus Exploits a Monocyte Differentiation Program To Promote Its Entry. Journal of Virology, 2013, 87, 3801-3814.	3.4	60
25	An Upstream Open Reading Frame Modulates Ebola Virus Polymerase Translation and Virus Replication. PLoS Pathogens, 2013, 9, e1003147.	4.7	66
26	DENV Inhibits Type I IFN Production in Infected Cells by Cleaving Human STING. PLoS Pathogens, 2012, 8, e1002934.	4.7	411
27	Structural basis for Marburg virus VP35–mediated immune evasion mechanisms. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20661-20666.	7.1	90
28	Inclusion Bodies Are a Site of Ebolavirus Replication. Journal of Virology, 2012, 86, 11779-11788.	3.4	183
29	Filoviral Immune Evasion Mechanisms. Viruses, 2011, 3, 1634-1649.	3.3	71
30	The Ebola Virus VP24 Protein Prevents hnRNP C1/C2 Binding to Karyopherin $\hat{l}\pm 1$ and Partially Alters its Nuclear Import. Journal of Infectious Diseases, 2011, 204, S904-S910.	4.0	45
31	DRBP76 Associates With Ebola Virus VP35 and Suppresses Viral Polymerase Function. Journal of Infectious Diseases, 2011, 204, S911-S918.	4.0	40
32	Modulation of type I IFN induction by a virulence determinant within the alphavirus nsP1 protein. Virology, 2010, 399, 1-10.	2.4	42
33	Basic Residues within the Ebolavirus VP35 Protein Are Required for Its Viral Polymerase Cofactor Function. Journal of Virology, 2010, 84, 10581-10591.	3.4	80
34	Structural and Functional Characterization of Reston Ebola Virus VP35 Interferon Inhibitory Domain. Journal of Molecular Biology, 2010, 399, 347-357.	4.2	61
35	Ross River Virus Envelope Glycans Contribute to Type I Interferon Production in Myeloid Dendritic Cells. Journal of Virology, 2008, 82, 12374-12383.	3.4	31
36	Increased Immunogenicity of a DNA-Launched Venezuelan Equine Encephalitis Virus-Based Replicon DNA Vaccine. Journal of Virology, 2007, 81, 13412-13423.	3.4	46

#	Article	IF	CITATIONS
37	Differential Induction of Type I Interferon Responses in Myeloid Dendritic Cells by Mosquito and Mammalian-Cell-Derived Alphaviruses. Journal of Virology, 2007, 81, 237-247.	3.4	85
38	Characterization of Ross River Virus Tropism and Virus-Induced Inflammation in a Mouse Model of Viral Arthritis and Myositis. Journal of Virology, 2006, 80, 737-749.	3.4	185
39	Identification of Adult Mouse Neurovirulence Determinants of the Sindbis Virus Strain AR86. Journal of Virology, 2005, 79, 4219-4228.	3.4	43