

Allison Groseth

List of Publications by Year in descending order

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Version: 2024-02-01

69
papers

3,222
citations

172457

29
h-index

155660

55
g-index

72
all docs

72
docs citations

72
times ranked

3606
citing authors

#	ARTICLE	IF	CITATIONS
1	Immunization with GP1 but Not Core-like Particles Displaying Isolated Receptor-Binding Epitopes Elicits Virus-Neutralizing Antibodies against JunÃn Virus. <i>Vaccines</i> , 2022, 10, 173.	4.4	5
2	Virusâ€Host Cell Interactions. <i>Cells</i> , 2022, 11, 804.	4.1	11
3	Generation of Reporter-Expressing New World Arenaviruses: A Systematic Comparison. <i>Viruses</i> , 2022, 14, 1563.	3.3	3
4	Remdesivir inhibits the polymerases of the novel filoviruses Lloviu and Bombali virus. <i>Antiviral Research</i> , 2021, 192, 105120.	4.1	8
5	CP100356 Hydrochloride, a P-Glycoprotein Inhibitor, Inhibits Lassa Virus Entry: Implication of a Candidate Pan-Mammarenavirus Entry Inhibitor. <i>Viruses</i> , 2021, 13, 1763.	3.3	2
6	Differences in Viral RNA Synthesis but Not Budding or Entry Contribute to the In Vitro Attenuation of Reston Virus Compared to Ebola Virus. <i>Microorganisms</i> , 2020, 8, 1215.	3.6	10
7	Complete genome sequence of Tacaribe virus. <i>Archives of Virology</i> , 2020, 165, 1899-1903.	2.1	6
8	BH3-only sensors Bad, Noxa and Puma are Key Regulators of Tacaribe virus-induced Apoptosis. <i>PLoS Pathogens</i> , 2020, 16, e1008948.	4.7	12
9	Therapeutic strategies to target the Ebola virus life cycle. <i>Nature Reviews Microbiology</i> , 2019, 17, 593-606.	28.6	110
10	High-throughput screening for negative-stranded hemorrhagic fever viruses using reverse genetics. <i>Antiviral Research</i> , 2019, 170, 104569.	4.1	13
11	Assessing cross-reactivity of JunÃn virus-directed neutralizing antibodies. <i>Antiviral Research</i> , 2019, 163, 106-116.	4.1	10
12	Assessment of the function and intergenus-compatibility of Ebola and Lloviu virus proteins. <i>Journal of General Virology</i> , 2019, 100, 760-772.	2.9	10
13	Apoptosis during arenavirus infection: mechanisms and evasion strategies. <i>Microbes and Infection</i> , 2018, 20, 65-80.	1.9	13
14	Serological Evidence for the Circulation of Ebolaviruses in Pigs From Sierra Leone. <i>Journal of Infectious Diseases</i> , 2018, 218, S305-S311.	4.0	20
15	Generation and Optimization of a Green Fluorescent Protein-Expressing Transcription and Replication-Competent Virus-Like Particle System for Ebola Virus. <i>Journal of Infectious Diseases</i> , 2018, 218, S360-S364.	4.0	14
16	Lifecycle modelling systems support inosine monophosphate dehydrogenase (IMPDH) as a pro-viral factor and antiviral target for New World arenaviruses. <i>Antiviral Research</i> , 2018, 157, 140-150.	4.1	21
17	A genome-wide siRNA screen identifies a druggable host pathway essential for the Ebola virus life cycle. <i>Genome Medicine</i> , 2018, 10, 58.	8.2	41
18	Forty Years of Ebolavirus Molecular Biology: Understanding a Novel Disease Agent Through the Development and Application of New Technologies. <i>Methods in Molecular Biology</i> , 2017, 1628, 15-38.	0.9	3

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19	Ebola virus VP24 interacts with NP to facilitate nucleocapsid assembly and genome packaging. <i>Scientific Reports</i> , 2017, 7, 7698.	3.3	55
20	Maguari Virus Associated with Human Disease. <i>Emerging Infectious Diseases</i> , 2017, 23, 1325-1331.	4.3	19
21	Generation of Recombinant Ebola Viruses Using Reverse Genetics. <i>Methods in Molecular Biology</i> , 2017, 1628, 177-188.	0.9	3
22	The Merits of Malaria Diagnostics during an Ebola Virus Disease Outbreak. <i>Emerging Infectious Diseases</i> , 2016, 22, 323-6.	4.3	25
23	Nanopore Sequencing as a Rapidly Deployable Ebola Outbreak Tool. <i>Emerging Infectious Diseases</i> , 2016, 22, 331-4.	4.3	175
24	Clinical Chemistry of Patients With Ebola in Monrovia, Liberia. <i>Journal of Infectious Diseases</i> , 2016, 214, S303-S307.	4.0	7
25	Response to Comment on "Mutation rate and genotype variation of Ebola virus from Mali case sequences". <i>Science</i> , 2016, 353, 658-658.	12.6	4
26	Plasmodium Parasitemia Associated With Increased Survival in Ebola Virus-Infected Patients. <i>Clinical Infectious Diseases</i> , 2016, 63, 1026-1033.	5.8	42
27	Ebola Laboratory Response at the Eternal Love Winning Africa Campus, Monrovia, Liberia, 2014-2015. <i>Journal of Infectious Diseases</i> , 2016, 214, S169-S176.	4.0	24
28	The New World arenavirus Tacaribe virus induces caspase-dependent apoptosis in infected cells. <i>Journal of General Virology</i> , 2016, 97, 855-866.	2.9	12
29	Complete genome sequence of trivittatus virus. <i>Archives of Virology</i> , 2015, 160, 2637-2639.	2.1	4
30	Mutation rate and genotype variation of Ebola virus from Mali case sequences. <i>Science</i> , 2015, 348, 117-119.	12.6	127
31	Spatiotemporal Analysis of Guaroa Virus Diversity, Evolution, and Spread in South America. <i>Emerging Infectious Diseases</i> , 2015, 21, 460-463.	4.3	4
32	Assessing the contribution of interferon antagonism to the virulence of West African Ebola viruses. <i>Nature Communications</i> , 2015, 6, 8000.	12.8	19
33	Molecular Characterization of Human Pathogenic Bunyaviruses of the Nyando and Bwamba/Pongola Virus Groups Leads to the Genetic Identification of Moju-dos Campos and Kaeng Khoi Virus. <i>PLoS Neglected Tropical Diseases</i> , 2014, 8, e3147.	3.0	23
34	Complete Genome Sequences of Three Ebola Virus Isolates from the 2014 Outbreak in West Africa. <i>Genome Announcements</i> , 2014, 2, .	0.8	28
35	A Novel Life Cycle Modeling System for Ebola Virus Shows a Genome Length-Dependent Role of VP24 in Virus Infectivity. <i>Journal of Virology</i> , 2014, 88, 10511-10524.	3.4	134
36	Development and application of reporter-expressing mononegaviruses: Current challenges and perspectives. <i>Antiviral Research</i> , 2014, 103, 78-87.	4.1	22

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37	A novel Ebola virus expressing luciferase allows for rapid and quantitative testing of antivirals. <i>Antiviral Research</i> , 2013, 99, 207-213.	4.1	55
38	The role of oligomerization for the biological functions of the arenavirus nucleoprotein. <i>Archives of Virology</i> , 2013, 158, 1895-1905.	2.1	11
39	Arenavirus Budding: A Common Pathway with Mechanistic Differences. <i>Viruses</i> , 2013, 5, 528-549.	3.3	29
40	An Upstream Open Reading Frame Modulates Ebola Virus Polymerase Translation and Virus Replication. <i>PLoS Pathogens</i> , 2013, 9, e1003147.	4.7	66
41	Cleavage of the Junin Virus Nucleoprotein Serves a Decoy Function To Inhibit the Induction of Apoptosis during Infection. <i>Journal of Virology</i> , 2013, 87, 224-233.	3.4	24
42	Profile and Persistence of the Virus-Specific Neutralizing Humoral Immune Response in Human Survivors of Sudan Ebolavirus (Gulu). <i>Journal of Infectious Diseases</i> , 2013, 208, 299-309.	4.0	47
43	The Ebola Virus Glycoprotein Contributes to but Is Not Sufficient for Virulence In Vivo. <i>PLoS Pathogens</i> , 2012, 8, e1002847.	4.7	88
44	Complete Genome Sequencing of Four Geographically Diverse Strains of Batai Virus. <i>Journal of Virology</i> , 2012, 86, 13844-13845.	3.4	14
45	Profiling the Native Specific Human Humoral Immune Response to Sudan Ebola Virus Strain Gulu by Chemiluminescence Enzyme-Linked Immunosorbent Assay. <i>Vaccine Journal</i> , 2012, 19, 1844-1852.	3.1	26
46	Complete Genome Sequencing of Mosquito and Human Isolates of Ngari Virus. <i>Journal of Virology</i> , 2012, 86, 13846-13847.	3.4	18
47	Inclusion Bodies Are a Site of Ebolavirus Replication. <i>Journal of Virology</i> , 2012, 86, 11779-11788.	3.4	183
48	Current ebola vaccines. <i>Expert Opinion on Biological Therapy</i> , 2012, 12, 859-872.	3.1	76
49	Minigenomes, transcription and replication competent virus-like particles and beyond: Reverse genetics systems for filoviruses and other negative stranded hemorrhagic fever viruses. <i>Antiviral Research</i> , 2011, 91, 195-208.	4.1	103
50	Clinical aspects of Marburg hemorrhagic fever. <i>Future Virology</i> , 2011, 6, 1091-1106.	1.8	102
51	Vesicular Stomatitis Virus-Based Ebola Vaccines With Improved Cross-Protective Efficacy. <i>Journal of Infectious Diseases</i> , 2011, 204, S1066-S1074.	4.0	102
52	Tacaribe Virus but Not Junin Virus Infection Induces Cytokine Release from Primary Human Monocytes and Macrophages. <i>PLoS Neglected Tropical Diseases</i> , 2011, 5, e1137.	3.0	51
53	Both matrix proteins of Ebola virus contribute to the regulation of viral genome replication and transcription. <i>Virology</i> , 2010, 403, 56-66.	2.4	131
54	Oligomerization of Ebola Virus VP40 Is Essential for Particle Morphogenesis and Regulation of Viral Transcription. <i>Journal of Virology</i> , 2010, 84, 7053-7063.	3.4	109

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55	Viral Protein Determinants of Lassa Virus Entry and Release from Polarized Epithelial Cells. <i>Journal of Virology</i> , 2010, 84, 3178-3188.	3.4	56
56	Efficient Budding of the Tacaribe Virus Matrix Protein Z Requires the Nucleoprotein. <i>Journal of Virology</i> , 2010, 84, 3603-3611.	3.4	59
57	Detection of Sudan ebolavirus (strain Gulu) epitopes that are targets of the humoral immune response in survivors. <i>International Journal of Infectious Diseases</i> , 2010, 14, e461-e462.	3.3	2
58	The Ebola virus ribonucleoprotein complex: A novel VP30â€“L interaction identified. <i>Virus Research</i> , 2009, 140, 8-14.	2.2	84
59	In Vitro Evaluation of Antisense RNA Efficacy against Filovirus Infection, by Use of Reverse Genetics. <i>Journal of Infectious Diseases</i> , 2007, 196, S382-S389.	4.0	13
60	In Vitro and In Vivo Characterization of Recombinant Ebola Viruses Expressing Enhanced Green Fluorescent Protein. <i>Journal of Infectious Diseases</i> , 2007, 196, S313-S322.	4.0	74
61	The ecology of Ebola virus. <i>Trends in Microbiology</i> , 2007, 15, 408-416.	7.7	201
62	Ebola virus: unravelling pathogenesis to combat a deadly disease. <i>Trends in Molecular Medicine</i> , 2006, 12, 206-215.	6.7	152
63	Infection of Naïve Target Cells with Virus-Like Particles: Implications for the Function of Ebola Virus VP24. <i>Journal of Virology</i> , 2006, 80, 7260-7264.	3.4	123
64	Hemorrhagic Fever Viruses as Biological Weapons. , 2005, , 169-191.		4
65	The role of reverse genetics systems in studying viral hemorrhagic fevers. <i>Thrombosis and Haemostasis</i> , 2005, 94, 240-53.	3.4	20
66	RNA Polymerase I-Driven Minigenome System for Ebola Viruses. <i>Journal of Virology</i> , 2005, 79, 4425-4433.	3.4	78
67	The role of reverse genetics systems in determining filovirus pathogenicity. , 2005, , 157-177.		5
68	Rescue of Ebola virus from cDNA using heterologous support proteins. <i>Virus Research</i> , 2004, 106, 43-50.	2.2	33
69	Molecular characterization of an isolate from the 1989/90 epizootic of Ebola virus Reston among macaques imported into the United States. <i>Virus Research</i> , 2002, 87, 155-163.	2.2	40