Agnieszka M Szemiel

List of Publications by Year in descending order

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840776 1281871 11 652 11 11 citations h-index g-index papers 14 14 14 1360 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A plasmid DNA-launched SARS-CoV-2 reverse genetics system and coronavirus toolkit for COVID-19 research. PLoS Biology, 2021, 19, e3001091.	5.6	163
2	Human Rhinovirus Infection Blocks Severe Acute Respiratory Syndrome Coronavirus 2 Replication Within the Respiratory Epithelium: Implications for COVID-19 Epidemiology. Journal of Infectious Diseases, 2021, 224, 31-38.	4.0	119
3	In vitro selection of Remdesivir resistance suggests evolutionary predictability of SARS-CoV-2. PLoS Pathogens, 2021, 17, e1009929.	4.7	108
4	Establishment of a reverse genetics system for Schmallenberg virus, a newly emerged orthobunyavirus in Europe. Journal of General Virology, 2013, 94, 851-859.	2.9	57
5	Bunyamwera orthobunyavirus glycoprotein precursor is processed by cellular signal peptidase and signal peptidase. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 8825-8830.	7.1	40
6	Role of Bunyamwera Orthobunyavirus NSs Protein in Infection of Mosquito Cells. PLoS Neglected Tropical Diseases, 2012, 6, e1823.	3.0	35
7	A role for glycolipid biosynthesis in severe fever with thrombocytopenia syndrome virus entry. PLoS Pathogens, 2017, 13, e1006316.	4.7	33
8	Elevated temperature inhibits SARS-CoV-2 replication in respiratory epithelium independently of IFN-mediated innate immune defenses. PLoS Biology, 2021, 19, e3001065.	5.6	26
9	Sugar feeding protects against arboviral infection by enhancing gut immunity in the mosquito vector Aedes aegypti. PLoS Pathogens, 2021, 17, e1009870.	4.7	23
10	Severe Acute Respiratory Syndrome Coronavirus 2 Serosurveillance in a Patient Population Reveals Differences in Virus Exposure and Antibody-Mediated Immunity According to Host Demography and Healthcare Setting. Journal of Infectious Diseases, 2021, 223, 971-980.	4.0	20
11	Development of reverse genetics systems and investigation of host response antagonism and reassortment potential for Cache Valley and Kairi viruses, two emerging orthobunyaviruses of the Americas. PLoS Neglected Tropical Diseases, 2018, 12, e0006884.	3.0	12