Andrii Slonchak

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8740363/publications.pdf

Version: 2024-02-01

932766 887659 20 567 10 17 citations h-index g-index papers 23 23 23 872 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Functional non-coding RNAs derived from the flavivirus $3\hat{a}\in^2$ untranslated region. Virus Research, 2015, 206, 53-61.	1.1	97
2	Expression of Mosquito MicroRNA Aae-miR-2940-5p Is Downregulated in Response to West Nile Virus Infection To Restrict Viral Replication. Journal of Virology, 2014, 88, 8457-8467.	1.5	90
3	Subgenomic flaviviral RNAs: What do we know after the first decade of research. Antiviral Research, 2018, 159, 13-25.	1.9	82
4	Human MicroRNA miR-532-5p Exhibits Antiviral Activity against West Nile Virus via Suppression of Host Genes SESTD1 and TAB3 Required for Virus Replication. Journal of Virology, 2016, 90, 2388-2402.	1.5	60
5	Determinants of Zika virus host tropism uncovered by deep mutational scanning. Nature Microbiology, 2019, 4, 876-887.	5.9	50
6	Zika virus noncoding RNA suppresses apoptosis and is required for virus transmission by mosquitoes. Nature Communications, 2020, 11, 2205.	5.8	50
7	Lower temperatures reduce type I interferon activity and promote alphaviral arthritis. PLoS Pathogens, 2017, 13, e1006788.	2.1	37
8	West Nile virus infection and interferon alpha treatment alter the spectrum and the levels of coding and noncoding host RNAs secreted in extracellular vesicles. BMC Genomics, 2019, 20, 474.	1.2	23
9	Helicase Domain of West Nile Virus NS3 Protein Plays a Role in Inhibition of Type I Interferon Signalling. Viruses, 2017, 9, 326.	1.5	13
10	Injection site vaccinology of a recombinant vaccinia-based vector reveals diverse innate immune signatures. PLoS Pathogens, 2021, 17, e1009215.	2.1	13
11	Structural analysis of 3'UTRs in insect flaviviruses reveals novel determinants of sfRNA biogenesis and provides new insights into flavivirus evolution. Nature Communications, 2022, 13, 1279.	5 . 8	13
12	Full genome sequence of Rocio virus reveal substantial variations from the prototype Rocio virus SPH 34675 sequence. Archives of Virology, 2018, 163, 255-258.	0.9	10
13	Sequencing of Historical Isolates, K-mer Mining and High Serological Cross-Reactivity with Ross River Virus Argue against the Presence of Getah Virus in Australia. Pathogens, 2020, 9, 848.	1.2	10
14	Reporter Flaviviruses as Tools to Demonstrate Homologous and Heterologous Superinfection Exclusion. Viruses, 2022, 14, 1501.	1.5	7
15	Neural Epidermal Growth Factor-Like Like Protein 2 Is Expressed in Human Oligodendroglial Cell Types. Frontiers in Cell and Developmental Biology, 2022, 10, 803061.	1.8	4
16	The distinguishing NS5-M114V mutation in American Zika virus isolates has negligible impacts on virus replication and transmission potential. PLoS Neglected Tropical Diseases, 2022, 16, e0010426.	1.3	4
17	Crosstalk between transcription factors in regulation of the human glutathione S-transferase P1 gene expression in Me45 melanoma cells. Biopolymers and Cell, 2009, 25, 210-217.	0.1	2
18	Regulation of the Glutathione S-Transferase P1 Expression in Melanoma Cells. , 0, , .		0

Andrii Slonchak

#	Article	IF	CITATIONS
19	Zika Virus sfRNA Plays an Essential Role in the Infection of Insects and Mammals. Proceedings (mdpi), 2020, 50, .	0.2	0
20	West Nile Virus Infection and Interferon Alpha Treatment Induce Incorporation of Common and Unique Host Antiviral RNAs Into Extracellular Vesicles. SSRN Electronic Journal, 0, , .	0.4	0