

Durrell D Kapan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4841011/publications.pdf>

Version: 2024-02-01

8
papers

499
citations

1163117
8
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1588992
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8
all docs

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docs citations

8
times ranked

870
citing authors

#	ARTICLE	IF	CITATIONS
1	Metagenomic shotgun sequencing reveals host species as an important driver of virome composition in mosquitoes. <i>Scientific Reports</i> , 2021, 11, 8448.	3.3	26
2	Human-Mosquito Contact: A Missing Link in Our Understanding of Mosquito-Borne Disease Transmission Dynamics. <i>Annals of the Entomological Society of America</i> , 2021, 114, 397-414.	2.5	30
3	Patterns, Drivers, and Challenges of Vector-Borne Disease Emergence. <i>Vector-Borne and Zoonotic Diseases</i> , 2020, 20, 159-170.	1.5	74
4	Estimates of introgression as a function of pairwise distances. <i>BMC Bioinformatics</i> , 2019, 20, 207.	2.6	21
5	Metagenomic shotgun sequencing of a Bunyavirus in wild-caught <i>Aedes aegypti</i> from Thailand informs the evolutionary and genomic history of the Phleboviruses. <i>Virology</i> , 2014, 464-465, 312-319.	2.4	46
6	<i>Flavobacterium akiainvivens</i> sp. nov., from decaying wood of <i>Wikstroemia oahuensis</i> , Hawaii, and emended description of the genus <i>Flavobacterium</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 3280-3286.	1.7	90
7	Hybridization Reveals the Evolving Genomic Architecture of Speciation. <i>Cell Reports</i> , 2013, 5, 666-677.	6.4	118
8	Mosquito Vector Diversity across Habitats in Central Thailand Endemic for Dengue and Other Arthropod-Borne Diseases. <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2507.	3.0	94