

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
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8
docs citations

8
times ranked

870
citing authors

| # | ARTICLE | IF | CITATIONS |
|---|--|-----|-----------|
| 1 | Hybridization Reveals the Evolving Genomic Architecture of Speciation. <i>Cell Reports</i> , 2013, 5, 666-677. | 6.4 | 118 |
| 2 | 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 |
| 3 | <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 |
| 4 | Patterns, Drivers, and Challenges of Vector-Borne Disease Emergence. <i>Vector-Borne and Zoonotic Diseases</i> , 2020, 20, 159-170. | 1.5 | 74 |
| 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 | 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 |
| 7 | 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 |
| 8 | Estimates of introgression as a function of pairwise distances. <i>BMC Bioinformatics</i> , 2019, 20, 207. | 2.6 | 21 |