Population bottlenecks and founder effects: implication emergence

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Citation Report

#	Article	IF	CITATIONS
1	Did Zika virus attenuation or increased virulence lead to the emergence of congenital Zika syndrome?. Journal of Travel Medicine, 2021, 28, .	1.4	8
2	Current Status of Chikungunya in India. Frontiers in Microbiology, 2021, 12, 695173.	1.5	24
3	Multilocus Genotyping of â€~Candidatus Phytoplasma solani' Associated with Rubbery Taproot Disease of Sugar Beet in the Pannonian Plain. Microorganisms, 2021, 9, 1950.	1.6	11
5	Lineage Divergence and Vector-Specific Adaptation Have Driven Chikungunya Virus onto Multiple Adaptive Landscapes. MBio, 2021, 12, e0273821.	1.8	8
6	Investigation of Biological Factors Contributing to Individual Variation in Viral Titer after Oral Infection of Aedes aegypti Mosquitoes by Sindbis Virus. Viruses, 2022, 14, 131.	1.5	7
7	Genetic Drift and Purifying Selection Shaped Mitochondrial Genome Variation in the High Royal Jelly-Producing Honeybee Strain (Apis mellifera ligustica). Frontiers in Genetics, 2022, 13, 835967.	1.1	2
8	Glycosaminoglycan binding by arboviruses: a cautionary tale. Journal of General Virology, 2022, 103, .	1.3	5
9	Impact of structural dynamics on biological functions of flaviviruses. FEBS Journal, 2023, 290, 1973-1985.	2.2	5
12	Peptide and Protein Alphavirus Antigens for Broad Spectrum Vaccine Design. SSRN Electronic Journal, 0, , .	0.4	0
13	Vector-Borne Viral Diseases as a Current Threat for Human and Animal Health—One Health Perspective. Journal of Clinical Medicine, 2022, 11, 3026.	1.0	22
14	Mitochondrial Genome Contributes to the Thermal Adaptation of the Oomycete Phytophthora infestans. Frontiers in Microbiology, 0, 13, .	1.5	0
15	Heterogeneity of Rift Valley fever virus transmission potential across livestock hosts, quantified through a model-based analysis of host viral load and vector infection. PLoS Computational Biology, 2022, 18, e1010314.	1.5	10
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19	Is monkeypox a new, emerging sexually transmitted disease? A rapid review of the literature. Journal of Medical Virology, 2023, 95, .	2.5	12
20	Neighboring mutationâ€mediated enhancement of dengue virus infectivity and spread. EMBO Reports, 2022, 23, .	2.0	5
21	Potential Nosocomial Infections by the Zika and Chikungunya Viruses in Public Health Facilities in the Metropolitan Area of Recife, Brazil. Tropical Medicine and Infectious Disease, 2022, 7, 351.	0.9	1
22	Incomplete bunyavirus particles can cooperatively support virus infection and spread. PLoS Biology, 2022, 20, e3001870.	2.6	8

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#	Article	I	IF	CITATIONS	
23	Impact of CHIKV Replication on the Global Proteome of Aedes albopictus Cells. Proteomes, 2022, 10,	38. :	1.7	3	
24	Plant Virus Adaptation to New Hosts: A Multi-scale Approach. Current Topics in Microbiology and Immunology, 2023, , 167-196.		0.7	1	
25	Low Transmission of Chikungunya Virus by AedesÂaegypti from Vientiane Capital, Lao PDR. Pathogen 2023, 12, 31.	S, _	1.2	2	
27	Does arbovirus emergence in humans require adaptation to domestic mosquitoes?. Current Opinion in Virology, 2023, 60, 101315.	ר ז	2.6	4	
28	Dynamical demographic phases explain how population growth and mutation control the evolutionary impact of bottlenecks. Physical Review Research, 2023, 5, .	:	1.3	1	
29	Determinants of Chikungunya and O'nyong-Nyong Virus Specificity for Infection of Aedes and Anopheles Mosquito Vectors. Viruses, 2023, 15, 589.		1.5	3	
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31	Chikungunya fever. Nature Reviews Disease Primers, 2023, 9, .		18.1	26	