CITATION REPORT List of articles citing

Homologous crossovers among molecules of brome mosaic bromovirus RNA1 or RNA2 segments in vivo

DOI: 10.1128/jvi.79.9.5732-5742.2005 Journal of Virology, 2005, 79, 5732-42.

Source: https://exaly.com/paper-pdf/39010611/citation-report.pdf

Version: 2024-04-28

This report has been generated based on the citations recorded by exaly.com for the above article. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

#	Paper	IF	Citations
42	A universal BMV-based RNA recombination systemhow to search for general rules in RNA recombination. <i>Nucleic Acids Research</i> , 2005 , 33, e105	20.1	14
41	Dynamics of mutation and recombination in a replicating population of complementing, defective viral genomes. <i>Journal of Molecular Biology</i> , 2006 , 360, 558-72	6.5	39
40	Characterization of a Brome mosaic virus strain and its use as a vector for gene silencing in monocotyledonous hosts. <i>Molecular Plant-Microbe Interactions</i> , 2006 , 19, 1229-39	3.6	162
39	Characterization of a novel 5Tsubgenomic RNA3a derived from RNA3 of Brome mosaic bromovirus. Journal of Virology, 2006 , 80, 12357-66	6.6	16
38	Recombination in Plant RNA Viruses. 2008 , 133-156		36
37	Plant Virus Evolution. 2008,		12
36	The promiscuous evolutionary history of the family Bromoviridae. <i>Journal of General Virology</i> , 2008 , 89, 1739-1747	4.9	39
35	cis- and trans-acting functions of brome mosaic virus protein 1a in genomic RNA1 replication. <i>Journal of Virology</i> , 2008 , 82, 3045-53	6.6	31
34	Repair of lost 5Tterminal sequences in tombusviruses: Rapid recovery of promoter- and enhancer-like sequences in recombinant RNAs. <i>Virology</i> , 2010 , 404, 96-105	3.6	7
33	How RNA viruses maintain their genome integrity. Journal of General Virology, 2010, 91, 1373-87	4.9	59
32	Advances in plant virus evolution: translating evolutionary insights into better disease management. <i>Phytopathology</i> , 2011 , 101, 1136-48	3.8	64
31	RNA-RNA recombination in plant virus replication and evolution. <i>Annual Review of Phytopathology</i> , 2011 , 49, 415-43	10.8	114
30	The evolutionary genetics of emerging plant RNA viruses. <i>Molecular Plant-Microbe Interactions</i> , 2011 , 24, 287-93	3.6	88
29	Why do RNA viruses recombine?. <i>Nature Reviews Microbiology</i> , 2011 , 9, 617-26	22.2	381
28	Recombination of 5Tsubgenomic RNA3a with genomic RNA3 of Brome mosaic bromovirus in vitro and in vivo. <i>Virology</i> , 2011 , 410, 129-41	3.6	11
27	Distribution of the phenotypic effects of random homologous recombination between two virus species. <i>PLoS Pathogens</i> , 2011 , 7, e1002028	7.6	15
26	Mutations in the antiviral RNAi defense pathway modify Brome mosaic virus RNA recombinant profiles. <i>Molecular Plant-Microbe Interactions</i> , 2012 , 25, 97-106	3.6	32

25	Two types of defective RNAs arising from the tomato black ring virus genome. <i>Archives of Virology</i> , 2012 , 157, 569-72	2.6	10
24	The genome of murine cytomegalovirus is shaped by purifying selection and extensive recombination. <i>Virology</i> , 2013 , 435, 258-68	3.6	23
23	Plant feeding by insect vectors can affect life cycle, population genetics and evolution of plant viruses. <i>Functional Ecology</i> , 2013 , 27, 610-622	5.6	67
22	Genetic recombination in plant-infecting messenger-sense RNA viruses: overview and research perspectives. <i>Frontiers in Plant Science</i> , 2013 , 4, 68	6.2	30
21	Tissue-specific attenuation of oncolytic sindbis virus without compromised genetic stability. <i>Human Gene Therapy Methods</i> , 2014 , 25, 154-65	4.9	8
20	Replication of Plant Viruses. 2014 , 341-421		4
19	Estimation of the in vivo recombination rate for a plant RNA virus. <i>Journal of General Virology</i> , 2014 , 95, 724-732	4.9	29
18	Phylogeny and molecular evolution of the hepatitis C virus. <i>Infection, Genetics and Evolution</i> , 2014 , 21, 67-82	4.5	38
17	Co-infection with two strains of Brome mosaic bromovirus reveals common RNA recombination sites in different hosts. <i>Virus Evolution</i> , 2015 , 1, vev021	3.7	3
16	Clonality and intracellular polyploidy in virus evolution and pathogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 8887-92	11.5	14
15	The Strange Lifestyle of Multipartite Viruses. <i>PLoS Pathogens</i> , 2016 , 12, e1005819	7.6	54
14	Molecular Basis of Genetic Variation of Viruses. 2016 , 35-71		4
13	Evolutionary ecology of virus emergence. Annals of the New York Academy of Sciences, 2017, 1389, 124-	1 <u>4.</u> 6	23
12	Biophysical analysis of BMV virions purified using a novel method. <i>Journal of Chromatography B:</i> Analytical Technologies in the Biomedical and Life Sciences, 2017 , 1068-1069, 157-163	3.2	1
11	Diversity of Plant Virus Populations: A Valuable Tool for Epidemiological Studies. 2017,		2
10	Quasispecies and virus. European Biophysics Journal, 2018 , 47, 443-457	1.9	38
9	An Improved Silencing Vector: Greater Insert Stability and More Extensive VIGS. <i>Plant Physiology</i> , 2018 , 176, 496-510	6.6	26
8	Recombination of host cell mRNA with the Asia 1 foot-and-mouth disease virus genome in cell suspension culture. <i>Archives of Virology</i> , 2019 , 164, 41-50	2.6	2

7	Molecular basis of genetic variation of viruses: error-prone replication. 2020 , 35-71		10	
6	Phylogenetic network analysis revealed the recombinant origin of the SARS-CoV-2 VOC202012/01 (B.1.1.7) variant first discovered in U.K		1	
5	From RNA World to SARS-CoV-2: The Edited Story of RNA Viral Evolution. <i>Cells</i> , 2021 , 10,	7.9	4	
4	Appendix III. 2010 , 365-468			
3	BMV Propagation, Extraction and Purification Using Chromatographic Methods. <i>Bio-protocol</i> , 2018 , 8, e2935	0.9		
2	VirDB: Crowdsourced Database for Evaluation of Dynamical Viral Infection Models. <i>Current Bioinformatics</i> , 2019 , 14, 740-748	4.7		
1	Cryo-EM reconstructions of BMV-derived virus-like particles reveal assembly defects in the icosahedral lattice structure <i>Nanoscale</i> , 2022 ,	7.7	1	