

Anouk Willemsen

List of Publications by Year in descending order

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papers

535
citations

840119

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docs citations

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739
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomic and phylogenetic characterization of ChPV2, a novel goat PV closely related to the Xi-PV1 species infecting bovines. <i>Virology Journal</i> , 2020, 17, 167.	1.4	4
2	Papillomaviruses infecting cetaceans exhibit signs of genome adaptation following a recombination event. <i>Virus Evolution</i> , 2020, 6, veaa038.	2.2	8
3	Genome Plasticity in Papillomaviruses and De Novo Emergence of E5 Oncogenes. <i>Genome Biology and Evolution</i> , 2019, 11, 1602-1617.	1.1	14
4	Origin and evolution of papillomavirus (onco)genes and genomes. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2019, 374, 20180303.	1.8	37
5	On the stability of sequences inserted into viral genomes. <i>Virus Evolution</i> , 2019, 5, vez045.	2.2	41
6	Going, going, gone: predicting the fate of genomic insertions in plant RNA viruses. <i>Heredity</i> , 2018, 121, 499-509.	1.2	10
7	High virulence does not necessarily impede viral adaptation to a new host: a case study using a plant RNA virus. <i>BMC Evolutionary Biology</i> , 2017, 17, 25.	3.2	5
8	<i>2b</i> or not <i>2b</i> : Experimental evolution of functional exogenous sequences in a plant RNA virus. <i>Genome Biology and Evolution</i> , 2017, 9, eww300.	1.1	12
9	Genetic variation and evolutionary forces shaping <i>Cucumber vein yellowing virus</i> populations: risk of emergence of virulent isolates in Europe. <i>Plant Pathology</i> , 2016, 65, 847-856.	1.2	10
10	Predicting the Stability of Homologous Gene Duplications in a Plant RNA Virus. <i>Genome Biology and Evolution</i> , 2016, 8, 3065-3082.	1.1	20
11	Multiple Barriers to the Evolution of Alternative Gene Orders in a Positive-Strand RNA Virus. <i>Genetics</i> , 2016, 202, 1503-1521.	1.2	31
12	Temporal Dynamics of Intrahost Molecular Evolution for a Plant RNA Virus. <i>Molecular Biology and Evolution</i> , 2015, 32, 1132-1147.	3.5	33
13	<sc>B</sc>razilian <i><sc>P</sc>otato virus <sc>Y</sc></i> isolates identified as members of a new clade facilitate the reconstruction of evolutionary traits within this species. <i>Plant Pathology</i> , 2015, 64, 799-807.	1.2	9
14	Genetic variability and evolutionary analysis of parietaria mottle virus: role of selection and genetic exchange. <i>Archives of Virology</i> , 2015, 160, 2611-2616.	0.9	5
15	Experimental Evolution of Pseudogenization and Gene Loss in a Plant RNA Virus. <i>Molecular Biology and Evolution</i> , 2014, 31, 121-134.	3.5	39
16	Genetic Variation and Possible Mechanisms Driving the Evolution of Worldwide <i>Fig mosaic virus</i> Isolates. <i>Phytopathology</i> , 2014, 104, 108-114.	1.1	33
17	Relocation of the Nlb Gene in the Tobacco Etch Potyvirus Genome. <i>Journal of Virology</i> , 2014, 88, 4586-4590.	1.5	12
18	Emergence and Phylodynamics of Citrus tristeza virus in Sicily, Italy. <i>PLoS ONE</i> , 2013, 8, e66700.	1.1	32

#	ARTICLE	IF	CITATIONS
19	An ancient history of gene duplications, fusions and losses in the evolution of APOBEC3 mutators in mammals. <i>BMC Evolutionary Biology</i> , 2012, 12, 71.	3.2	130
20	EcPV2 DNA in equine genital squamous cell carcinomas and normal genital mucosa. <i>Veterinary Microbiology</i> , 2012, 158, 33-41.	0.8	44