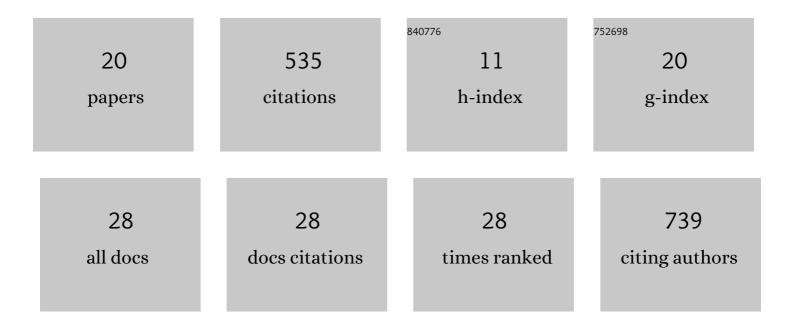
Anouk Willemsen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6468532/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	An ancient history of gene duplications, fusions and losses in the evolution of APOBEC3 mutators in mammals. BMC Evolutionary Biology, 2012, 12, 71.	3.2	130
2	EcPV2 DNA in equine genital squamous cell carcinomas and normal genital mucosa. Veterinary Microbiology, 2012, 158, 33-41.	1.9	44
3	On the stability of sequences inserted into viral genomes. Virus Evolution, 2019, 5, vez045.	4.9	41
4	Experimental Evolution of Pseudogenization and Gene Loss in a Plant RNA Virus. Molecular Biology and Evolution, 2014, 31, 121-134.	8.9	39
5	Origin and evolution of papillomavirus (onco)genes and genomes. Philosophical Transactions of the Royal Society B: Biological Sciences, 2019, 374, 20180303.	4.0	37
6	Genetic Variation and Possible Mechanisms Driving the Evolution of Worldwide <i>Fig mosaic virus</i> Isolates. Phytopathology, 2014, 104, 108-114.	2.2	33
7	Temporal Dynamics of Intrahost Molecular Evolution for a Plant RNA Virus. Molecular Biology and Evolution, 2015, 32, 1132-1147.	8.9	33
8	Emergence and Phylodynamics of Citrus tristeza virus in Sicily, Italy. PLoS ONE, 2013, 8, e66700.	2.5	32
9	Multiple Barriers to the Evolution of Alternative Gene Orders in a Positive-Strand RNA Virus. Genetics, 2016, 202, 1503-1521.	2.9	31
10	Predicting the Stability of Homologous Gene Duplications in a Plant RNA Virus. Genome Biology and Evolution, 2016, 8, 3065-3082.	2.5	20
11	Genome Plasticity in Papillomaviruses and De Novo Emergence of E5 Oncogenes. Genome Biology and Evolution, 2019, 11, 1602-1617.	2.5	14
12	Relocation of the NIb Gene in the Tobacco Etch Potyvirus Genome. Journal of Virology, 2014, 88, 4586-4590.	3.4	12
13	<i>2b</i> or not <i>2b</i> : Experimental evolution of functional exogenous sequences in a plant RNA virus. Genome Biology and Evolution, 2017, 9, evw300.	2.5	12
14	Genetic variation and evolutionary forces shaping <i>Cucumber vein yellowing virus</i> populations: risk of emergence of virulent isolates in Europe. Plant Pathology, 2016, 65, 847-856.	2.4	10
15	Going, going, gone: predicting the fate of genomic insertions in plant RNA viruses. Heredity, 2018, 121, 499-509.	2.6	10
16	<scp>B</scp> razilian <i><scp>P</scp>otato virus <scp>Y</scp></i> isolates identified as members of a new clade facilitate the reconstruction of evolutionary traits within this species. Plant Pathology, 2015, 64, 799-807.	2.4	9
17	Papillomaviruses infecting cetaceans exhibit signs of genome adaptation following a recombination event. Virus Evolution, 2020, 6, veaa038.	4.9	8
18	Genetic variability and evolutionary analysis of parietaria mottle virus: role of selection and genetic exchange. Archives of Virology, 2015, 160, 2611-2616.	2.1	5

#	Article	IF	CITATIONS
19	High virulence does not necessarily impede viral adaptation to a new host: a case study using a plant RNA virus. BMC Evolutionary Biology, 2017, 17, 25.	3.2	5
20	Genomic and phylogenetic characterization of ChPV2, a novel goat PV closely related to the Xi-PV1 species infecting bovines. Virology Journal, 2020, 17, 167.	3.4	4