Lia van der Hoek

List of Publications by Year in descending order

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186265 128289 4,784 62 28 60 citations h-index g-index papers 63 63 63 7753 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Identification of a new human coronavirus. Nature Medicine, 2004, 10, 368-373.	30.7	1,573
2	Seasonal coronavirus protective immunity is short-lasting. Nature Medicine, 2020, 26, 1691-1693.	30.7	637
3	Croup Is Associated with the Novel Coronavirus NL63. PLoS Medicine, 2005, 2, e240.	8.4	239
4	Human Coronavirus NL63 and 229E Seroconversion in Children. Journal of Clinical Microbiology, 2008, 46, 2368-2373.	3.9	171
5	Human coronavirus NL63, a new respiratory virus. FEMS Microbiology Reviews, 2006, 30, 760-773.	8.6	163
6	The dominance of human coronavirus OC43 and NL63 infections in infants. Journal of Clinical Virology, 2012, 53, 135-139.	3.1	161
7	Mosaic Structure of Human Coronavirus NL63, One Thousand Years of Evolution. Journal of Molecular Biology, 2006, 364, 964-973.	4.2	149
8	Replication-dependent downregulation of cellular angiotensin-converting enzyme 2 protein expression by human coronavirus NL63. Journal of General Virology, 2012, 93, 1924-1929.	2.9	128
9	Human Coronavirus NL63 Molecular Epidemiology and Evolutionary Patterns in Rural Coastal Kenya. Journal of Infectious Diseases, 2018, 217, 1728-1739.	4.0	116
10	Atypical Porcine Pestivirus: A Possible Cause of Congenital Tremor Type Aâ€II in Newborn Piglets. Viruses, 2016, 8, 271.	3.3	106
11	Human anelloviruses: diverse, omnipresent and commensal members of the virome. FEMS Microbiology Reviews, 2020, 44, 305-313.	8.6	90
12	Aerosol persistence in relation to possible transmission of SARS-CoV-2. Physics of Fluids, 2020, 32, 107108.	4.0	84
13	A Sensitive Assay for Virus Discovery in Respiratory Clinical Samples. PLoS ONE, 2011, 6, e16118.	2.5	80
14	A Novel Virus Causes Scale Drop Disease in Lates calcarifer. PLoS Pathogens, 2015, 11, e1005074.	4.7	80
15	Full Genome Virus Detection in Fecal Samples Using Sensitive Nucleic Acid Preparation, Deep Sequencing, and a Novel Iterative Sequence Classification Algorithm. PLoS ONE, 2014, 9, e93269.	2.5	71
16	Unbiased whole-genome deep sequencing of human and porcine stool samples reveals circulation of multiple groups of rotaviruses and a putative zoonotic infection. Virus Evolution, 2016, 2, vew027.	4.9	52
17	Unexplained diarrhoea in HIV-1 infected individuals. BMC Infectious Diseases, 2014, 14, 22.	2.9	48
18	The Vietnam Initiative on Zoonotic Infections (VIZIONS): A Strategic Approach to Studying Emerging Zoonotic Infectious Diseases. EcoHealth, 2015, 12, 726-735.	2.0	47

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19	Renin–angiotensin system in human coronavirus pathogenesis. Future Virology, 2010, 5, 145-161.	1.8	46
20	Sexual Behavior and Its Determinants During COVID-19 Restrictions Among Men Who Have Sex With Men in Amsterdam. Journal of Acquired Immune Deficiency Syndromes (1999), 2021, 86, 288-296.	2.1	46
21	Recommendations for the introduction of metagenomic high-throughput sequencing in clinical virology, part I: Wet lab procedure. Journal of Clinical Virology, 2021, 134, 104691.	3.1	42
22	Viral Metagenomics on Cerebrospinal Fluid. Genes, 2019, 10, 332.	2.4	38
23	Performance of VIDISCA-454 in Feces-Suspensions and Serum. Viruses, 2012, 4, 1328-1334.	3.3	37
24	Differentiation between Human Coronaviruses NL63 and 229E Using a Novel Double-Antibody Sandwich Enzyme-Linked Immunosorbent Assay Based on Specific Monoclonal Antibodies. Vaccine Journal, 2011, 18, 113-118.	3.1	36
25	Novel Orthobunyavirus Identified in the Cerebrospinal Fluid of a Ugandan Child With Severe Encephalopathy. Clinical Infectious Diseases, 2019, 68, 139-142.	5.8	35
26	Entamoeba and Giardia parasites implicated as hosts of CRESS viruses. Nature Communications, 2020, 11, 4620.	12.8	34
27	Burden of disease due to human coronavirus NL63 infections and periodicity of infection. Journal of Clinical Virology, 2010, 48, 104-108.	3.1	33
28	Diverging trends in incidence of HIV versus other sexually transmitted infections in HIV-negative MSM in Amsterdam. Aids, 2020, 34, 301-309.	2.2	33
29	Characterization of Posa and Posa-like virus genomes in fecal samples from humans, pigs, rats, and bats collected from a single location in Vietnam. Virus Evolution, 2017, 3, vex022.	4.9	25
30	A novel genus in the order Picornavirales detected in human stool. Journal of General Virology, 2015, 96, 3440-3443.	2.9	23
31	Two serological approaches for detection of antibodies to SARS-CoV-2 in different scenarios: a screening tool and a point-of-care test. Diagnostic Microbiology and Infectious Disease, 2020, 98, 115167.	1.8	21
32	Increased Multinucleoside Drug Resistance and Decreased Replicative Capacity of a Human Immunodeficiency Virus Type 1 Variant with an 8-Amino-Acid Insert in the Reverse Transcriptase. Journal of Virology, 2005, 79, 3536-3543.	3.4	20
33	Enhanced bioinformatic profiling of VIDISCA libraries for virus detection and discovery. Virus Research, 2019, 263, 21-26.	2.2	18
34	Diversity and Long-Term Dynamics of Human Blood Anelloviruses. Journal of Virology, 2022, 96, e0010922.	3.4	18
35	Identification of a new genotype of Torque Teno Mini virus. Virology Journal, 2013, 10, 323.	3.4	17
36	Virus discovery: are we scientists or genome collectors?. Trends in Microbiology, 2014, 22, 229-231.	7.7	17

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37	Severe Acute Respiratory Syndrome Coronavirus 2 Vaccination Boosts Neutralizing Activity Against Seasonal Human Coronaviruses. Clinical Infectious Diseases, 2022, 75, e653-e661.	5.8	16
38	Me Tri virus: a Semliki Forest virus strain from Vietnam?. Journal of General Virology, 2008, 89, 2132-2135.	2.9	15
39	Metagenomic analysis of a sample from a patient with respiratory tract infection reveals the presence of a $\hat{A}\hat{Z}\hat{A}^3$ -papillomavirus. Frontiers in Microbiology, 2014, 5, 347.	3.5	15
40	Persistent viremia by a novel parvovirus in a slow loris (Nycticebus coucang) with diffuse histiocytic sarcoma. Frontiers in Microbiology, 2014, 5, 655.	3.5	14
41	Emerging orthobunyaviruses associated with CNS disease. PLoS Neglected Tropical Diseases, 2020, 14, e0008856.	3.0	14
42	Reduced maternal levels of common viruses during pregnancy predict offspring psychosis: Potential role of enhanced maternal immune activity?. Schizophrenia Research, 2015, 166, 248-254.	2.0	13
43	Atypical Porcine Pestivirus Circulation and Molecular Evolution within an Affected Swine Herd. Viruses, 2020, 12, 1080.	3.3	12
44	Sequence-independent VIDISCA-454 technique to discover new viruses in canine livers. Journal of Virological Methods, 2012, 185, 152-155.	2.1	11
45	Limited geographic distribution of the novel cyclovirus CyCV-VN. Scientific Reports, 2014, 4, 3967.	3.3	11
46	Virus discovery reveals frequent infection by diverse novel members of the Flaviviridae in wild lemurs. Archives of Virology, 2019, 164, 509-522.	2.1	11
47	Early-Life Colonization by Anelloviruses in Infants. Viruses, 2022, 14, 865.	3.3	9
48	Absence of accessory genes in a divergent simian T-lymphotropic virus type 1 isolated from a bonnet macaque (Macaca radiata). PLoS Neglected Tropical Diseases, 2019, 13, e0007521.	3.0	8
49	Atypical Antibody Dynamics During Human Coronavirus HKU1 Infections. Frontiers in Microbiology, 2022, 13, 853410.	3.5	8
50	No Evidence Known Viruses Play a Role in the Pathogenesis of Onchocerciasis-Associated Epilepsy. An Explorative Metagenomic Case-Control Study. Pathogens, 2021, 10, 787.	2.8	7
51	Divergent Rhabdovirus Discovered in a Patient with New-Onset Nodding Syndrome. Viruses, 2022, 14, 210.	3.3	6
52	Comparable endemic coronavirus nucleoproteinâ€specific antibodies in mild and severe Covidâ€19 patients. Journal of Medical Virology, 2021, 93, 5614-5617.	5.0	5
53	Dynamics of the Enteric Virome in a Swine Herd Affected by Non-PCV2/PRRSV Postweaning Wasting Syndrome. Viruses, 2021, 13, 2538.	3.3	4
54	SARS-CoV-2 Re-infections: Lessons from Other Coronaviruses. Med, 2020, 1, 23-28.	4.4	3

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55	Prevalence of Neutralising Antibodies to HCoV-NL63 in Healthy Adults in Australia. Viruses, 2021, 13, 1618.	3.3	3
56	Diagnostic performance of two serological assays for the detection of SARS-CoV-2 specific antibodies: surveillance after vaccination. Diagnostic Microbiology and Infectious Disease, 2022, 102, 115650.	1.8	3
57	Genome Sequences of a Novel Vietnamese Bat Bunyavirus. Genome Announcements, 2016, 4, .	0.8	2
58	Host switching pathogens, infectious outbreaks and zoonosis: A Marie SkÅ,odowska-Curie innovative training network (HONOURs). Virus Research, 2018, 257, 120-124.	2.2	2
59	A Versatile Processing Workflow to Enable Pathogen Detection in Clinical Samples from Organs Using VIDISCA. Diagnostics, 2021, 11, 791.	2.6	2
60	Women in the European Virus Bioinformatics Center. Viruses, 2022, 14, 1522.	3.3	1
61	Atypical Porcine Pestivirus Molecular Evolution within a Persistently Infected Swine Farm. Proceedings (mdpi), 2020, 50, 29.	0.2	0
62	Revolution in Evolution of Human "Anellome― Proceedings (mdpi), 2020, 50, .	0.2	0