List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Severe Acute Respiratory Syndrome Coronavirus 2 Vaccination Boosts Neutralizing Activity Against Seasonal Human Coronaviruses. Clinical Infectious Diseases, 2022, 75, e653-e661.	5.8	16
2	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
3	Divergent Rhabdovirus Discovered in a Patient with New-Onset Nodding Syndrome. Viruses, 2022, 14, 210.	3.3	6
4	Early-Life Colonization by Anelloviruses in Infants. Viruses, 2022, 14, 865.	3.3	9
5	Atypical Antibody Dynamics During Human Coronavirus HKU1 Infections. Frontiers in Microbiology, 2022, 13, 853410.	3.5	8
6	Diversity and Long-Term Dynamics of Human Blood Anelloviruses. Journal of Virology, 2022, 96, e0010922.	3.4	18
7	Women in the European Virus Bioinformatics Center. Viruses, 2022, 14, 1522.	3.3	1
8	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
9	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
10	A Versatile Processing Workflow to Enable Pathogen Detection in Clinical Samples from Organs Using VIDISCA. Diagnostics, 2021, 11, 791.	2.6	2
11	Comparable endemic coronavirus nucleoproteinâ€specific antibodies in mild and severe Covidâ€19 patients. Journal of Medical Virology, 2021, 93, 5614-5617.	5.0	5
12	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
13	Prevalence of Neutralising Antibodies to HCoV-NL63 in Healthy Adults in Australia. Viruses, 2021, 13, 1618.	3.3	3
14	Dynamics of the Enteric Virome in a Swine Herd Affected by Non-PCV2/PRRSV Postweaning Wasting Syndrome. Viruses, 2021, 13, 2538.	3.3	4
15	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
16	Entamoeba and Giardia parasites implicated as hosts of CRESS viruses. Nature Communications, 2020, 11, 4620.	12.8	34
17	Atypical Porcine Pestivirus Circulation and Molecular Evolution within an Affected Swine Herd. Viruses, 2020, 12, 1080.	3.3	12
18	Aerosol persistence in relation to possible transmission of SARS-CoV-2. Physics of Fluids, 2020, 32, 107108.	4.0	84

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19	Seasonal coronavirus protective immunity is short-lasting. Nature Medicine, 2020, 26, 1691-1693.	30.7	637
20	SARS-CoV-2 Re-infections: Lessons from Other Coronaviruses. Med, 2020, 1, 23-28.	4.4	3
21	Human anelloviruses: diverse, omnipresent and commensal members of the virome. FEMS Microbiology Reviews, 2020, 44, 305-313.	8.6	90
22	Atypical Porcine Pestivirus Molecular Evolution within a Persistently Infected Swine Farm. Proceedings (mdpi), 2020, 50, 29.	0.2	0
23	Revolution in Evolution of Human "Anellome― Proceedings (mdpi), 2020, 50, .	0.2	0
24	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
25	Emerging orthobunyaviruses associated with CNS disease. PLoS Neglected Tropical Diseases, 2020, 14, e0008856.	3.0	14
26	Novel Orthobunyavirus Identified in the Cerebrospinal Fluid of a Ugandan Child With Severe Encephalopathy. Clinical Infectious Diseases, 2019, 68, 139-142.	5.8	35
27	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
28	Viral Metagenomics on Cerebrospinal Fluid. Genes, 2019, 10, 332.	2.4	38
29	Enhanced bioinformatic profiling of VIDISCA libraries for virus detection and discovery. Virus Research, 2019, 263, 21-26.	2.2	18
30	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
31	Human Coronavirus NL63 Molecular Epidemiology and Evolutionary Patterns in Rural Coastal Kenya. Journal of Infectious Diseases, 2018, 217, 1728-1739.	4.0	116
32	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
33	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
34	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
35	Atypical Porcine Pestivirus: A Possible Cause of Congenital Tremor Type Aâ€ll in Newborn Piglets. Viruses, 2016, 8, 271	3.3	106
36	Genome Sequences of a Novel Vietnamese Bat Bunyavirus. Genome Announcements, 2016, 4, .	0.8	2

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37	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
38	The Vietnam Initiative on Zoonotic Infections (VIZIONS): A Strategic Approach to Studying Emerging Zoonotic Infectious Diseases. EcoHealth, 2015, 12, 726-735.	2.0	47
39	A novel genus in the order Picornavirales detected in human stool. Journal of General Virology, 2015, 96, 3440-3443.	2.9	23
40	A Novel Virus Causes Scale Drop Disease in Lates calcarifer. PLoS Pathogens, 2015, 11, e1005074.	4.7	80
41	Metagenomic analysis of a sample from a patient with respiratory tract infection reveals the presence of a ÅŽÅ ³ -papillomavirus. Frontiers in Microbiology, 2014, 5, 347.	3.5	15
42	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
43	Unexplained diarrhoea in HIV-1 infected individuals. BMC Infectious Diseases, 2014, 14, 22.	2.9	48
44	Virus discovery: are we scientists or genome collectors?. Trends in Microbiology, 2014, 22, 229-231.	7.7	17
45	Limited geographic distribution of the novel cyclovirus CyCV-VN. Scientific Reports, 2014, 4, 3967.	3.3	11
46	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
47	Identification of a new genotype of Torque Teno Mini virus. Virology Journal, 2013, 10, 323.	3.4	17
48	Performance of VIDISCA-454 in Feces-Suspensions and Serum. Viruses, 2012, 4, 1328-1334.	3.3	37
49	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
50	Sequence-independent VIDISCA-454 technique to discover new viruses in canine livers. Journal of Virological Methods, 2012, 185, 152-155.	2.1	11
51	The dominance of human coronavirus OC43 and NL63 infections in infants. Journal of Clinical Virology, 2012, 53, 135-139.	3.1	161
52	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
53	A Sensitive Assay for Virus Discovery in Respiratory Clinical Samples. PLoS ONE, 2011, 6, e16118.	2.5	80
54	Renin–angiotensin system in human coronavirus pathogenesis. Future Virology, 2010, 5, 145-161.	1.8	46

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55	Burden of disease due to human coronavirus NL63 infections and periodicity of infection. Journal of Clinical Virology, 2010, 48, 104-108.	3.1	33
56	Me Tri virus: a Semliki Forest virus strain from Vietnam?. Journal of General Virology, 2008, 89, 2132-2135.	2.9	15
57	Human Coronavirus NL63 and 229E Seroconversion in Children. Journal of Clinical Microbiology, 2008, 46, 2368-2373.	3.9	171
58	Mosaic Structure of Human Coronavirus NL63, One Thousand Years of Evolution. Journal of Molecular Biology, 2006, 364, 964-973.	4.2	149
59	Human coronavirus NL63, a new respiratory virus. FEMS Microbiology Reviews, 2006, 30, 760-773.	8.6	163
60	Croup Is Associated with the Novel Coronavirus NL63. PLoS Medicine, 2005, 2, e240.	8.4	239
61	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
62	Identification of a new human coronavirus. Nature Medicine, 2004, 10, 368-373.	30.7	1,573