

# Cadhla Firth

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3372836/publications.pdf>

Version: 2024-02-01

35  
papers

2,961  
citations

279798

23  
h-index

377865

34  
g-index

36  
all docs

36  
docs citations

36  
times ranked

5084  
citing authors

#	ARTICLE	IF	CITATIONS
1	Phyldynamic Inference of Bacterial Outbreak Parameters Using Nanopore Sequencing. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	9
2	First detection of a novel "unknown host"™ flavivirus in a Malaysian rodent. <i>Access Microbiology</i> , 2021, 3, 000223.	0.5	1
3	The Geographic Distribution, Venom Components, Pathology and Treatments of Stonefish ( <i>Synanceia</i> ) Tj ETQq1 1 0.784314 rgBT /Ov	4.6	12
4	Evolutionary history of Simbu serogroup orthobunyaviruses in the Australian episytem. <i>Virology</i> , 2019, 535, 32-44.	2.4	11
5	Association of rodent-borne <i>Leptospira</i> spp. with urban environments in Malaysian Borneo. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007141.	3.0	42
6	Discovery of two highly divergent negative-sense RNA viruses associated with the parasitic nematode, <i>Capillaria hepatica</i> , in wild <i>Mus musculus</i> from New York City. <i>Journal of General Virology</i> , 2019, 100, 1350-1362.	2.9	16
7	High Prevalence of Rodent-Borne <i>Bartonella</i> spp. in Urbanizing Environments in Sarawak, Malaysian Borneo. <i>American Journal of Tropical Medicine and Hygiene</i> , 2019, 100, 506-509.	1.4	12
8	Rodent-Borne <i>Bartonella</i> Infection Varies According to Host Species Within and Among Cities. <i>EcoHealth</i> , 2017, 14, 771-782.	2.0	31
9	Genomic analysis of bluetongue virus episytems in Australia and Indonesia. <i>Veterinary Research</i> , 2017, 48, 82.	3.0	15
10	Genetic Characterization of Archived Bunyaviruses and their Potential for Emergence in Australia. <i>Emerging Infectious Diseases</i> , 2016, 22, 833-840.	4.3	11
11	Evolution of Genome Size and Complexity in the Rhabdoviridae. <i>PLoS Pathogens</i> , 2015, 11, e1004664.	4.7	149
12	Genomic Characterization of Yogue, Kasokero, Issyk-Kul, Keterah, Gossas, and Thiafora Viruses: Nairoviruses Naturally Infecting Bats, Shrews, and Ticks. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 93, 1041-1051.	1.4	36
13	Ledantavirus: A Proposed New Genus in the Rhabdoviridae has a Strong Ecological Association with Bats. <i>American Journal of Tropical Medicine and Hygiene</i> , 2015, 92, 405-410.	1.4	27
14	Detection of Zoonotic Pathogens and Characterization of Novel Viruses Carried by Commensal <i>Rattus norvegicus</i> in New York City. <i>MBio</i> , 2014, 5, e01933-14.	4.1	310
15	Koolpinyah and Yata viruses: Two newly recognised ephemeroviruses from tropical regions of Australia and Africa. <i>Veterinary Microbiology</i> , 2014, 174, 547-553.	1.9	10
16	Mesoniviruses are mosquito-specific viruses with extensive geographic distribution and host range. <i>Virology Journal</i> , 2014, 11, 97.	3.4	65
17	Urbanisation brings animals and diseases closer to home. <i>Ecos</i> , 2014, , .	0.0	0
18	Comparative full length genome sequence analysis of usutu virus isolates from Africa. <i>Virology Journal</i> , 2013, 10, 217.	3.4	31

#	ARTICLE	IF	CITATIONS
19	The Genomics of Emerging Pathogens. Annual Review of Genomics and Human Genetics, 2013, 14, 281-300.	6.2	50
20	Viral surveillance and discovery. Current Opinion in Virology, 2013, 3, 199-204.	5.4	57
21	Bats are a major natural reservoir for hepaciviruses and pegiviruses. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 8194-8199.	7.1	251
22	Diversity and Distribution of Hantaviruses in South America. Journal of Virology, 2012, 86, 13756-13766.	3.4	67
23	Worldwide emergence of multiple clades of enterovirus 68. Journal of General Virology, 2012, 93, 1952-1958.	2.9	191
24	Characterization of a canine homolog of hepatitis C virus. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 11608-11613.	7.1	250
25	Lack of Evidence for an Association between Iridovirus and Colony Collapse Disorder. PLoS ONE, 2011, 6, e21844.	2.5	17
26	Phylogenetic Analysis Reveals Rapid Evolutionary Dynamics in the Plant RNA Virus Genus Tobamovirus. Journal of Molecular Evolution, 2010, 71, 298-307.	1.8	49
27	Astrovirus Encephalitis in Boy with X-linked Agammaglobulinemia. Emerging Infectious Diseases, 2010, 16, 918-925.	4.3	283
28	Yellow Fever Virus Exhibits Slower Evolutionary Dynamics than Dengue Virus. Journal of Virology, 2010, 84, 765-772.	3.4	69
29	Using Time-Structured Data to Estimate Evolutionary Rates of Double-Stranded DNA Viruses. Molecular Biology and Evolution, 2010, 27, 2038-2051.	8.9	279
30	Insights into the Evolutionary History of an Emerging Livestock Pathogen: Porcine Circovirus 2. Journal of Virology, 2009, 83, 12813-12821.	3.4	208
31	Hantavirus Evolution in Relation to Its Rodent and Insectivore Hosts: No Evidence for Codivergence. Molecular Biology and Evolution, 2008, 26, 143-153.	8.9	209
32	Population Genetics of <i>Ambystoma jeffersonianum</i> and Sympatric Unisexuals Reveal Signatures of Both Gynogenetic and Sexual Reproduction. Copeia, 2008, 2008, 586-594.	1.3	13
33	High Rates of Molecular Evolution in Hantaviruses. Molecular Biology and Evolution, 2008, 25, 1488-1492.	8.9	117
34	A nonlethal method of identification of <i>Ambystoma laterale</i> , <i>A. jeffersonianum</i> and sympatric unisexuals. Molecular Ecology Notes, 2006, 6, 261-264.	1.7	24
35	An unexpected recent ancestor of unisexual <i>Ambystoma</i> . Molecular Ecology, 2006, 15, 3339-3351.	3.9	37