## Randal J Schoepp

List of Publications by Year in descending order

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PANDAL I SCHOEDD

#	Article	IF	CITATIONS
1	Pathology of Fatal West Nile Virus Infections in Native and Exotic Birds during the 1999 Outbreak in New York City, New York. Veterinary Pathology, 2000, 37, 208-224.	1.7	429
2	Molecular Evidence of Sexual Transmission of Ebola Virus. New England Journal of Medicine, 2015, 373, 2448-2454.	27.0	380
3	Three-dimensional structure of a membrane-containing virus Proceedings of the National Academy of Sciences of the United States of America, 1993, 90, 9095-9099.	7.1	205
4	Comprehensive Panel of Real-Time TaqManâ,,¢ Polymerase Chain Reaction Assays for Detection and Absolute Quantification of Filoviruses, Arenaviruses, and New World Hantaviruses. American Journal of Tropical Medicine and Hygiene, 2010, 82, 954-960.	1.4	195
5	Long-term sequelae after Ebola virus disease in Bundibugyo, Uganda: a retrospective cohort study. Lancet Infectious Diseases, The, 2015, 15, 905-912.	9.1	193
6	Possible sexual transmission of Ebola virus - Liberia, 2015. Morbidity and Mortality Weekly Report, 2015, 64, 479-81.	15.1	132
7	Undiagnosed Acute Viral Febrile Illnesses, Sierra Leone. Emerging Infectious Diseases, 2014, 20, 1176-1182.	4.3	122
8	FDA-ARGOS is a database with public quality-controlled reference genomes for diagnostic use and regulatory science. Nature Communications, 2019, 10, 3313.	12.8	101
9	Evolution and Spread of Ebola Virus in Liberia, 2014–2015. Cell Host and Microbe, 2015, 18, 659-669.	11.0	87
10	Seroprevalence and distribution of arboviral infections among rural Kenyan adults: A cross-sectional study. Virology Journal, 2011, 8, 371.	3.4	85
11	Nomenclature- and Database-Compatible Names for the Two Ebola Virus Variants that Emerged in Guinea and the Democratic Republic of the Congo in 2014. Viruses, 2014, 6, 4760-4799.	3.3	83
12	Monitoring of Ebola Virus Makona Evolution through Establishment of Advanced Genomic Capability in Liberia. Emerging Infectious Diseases, 2015, 21, 1135-1143.	4.3	79
13	Lassa virus-like particles displaying all major immunological determinants as a vaccine candidate for Lassa hemorrhagic fever. Virology Journal, 2010, 7, 279.	3.4	77
14	Early Events in the Pathogenesis of Eastern Equine Encephalitis Virus in Mice. American Journal of Pathology, 2005, 166, 159-171.	3.8	71
15	Crimean-Congo Hemorrhagic Fever, Afghanistan, 2009. Emerging Infectious Diseases, 2011, 17, 1940-1941.	4.3	66
16	Isolation and characterisation of Ebolavirus-specific recombinant antibody fragments from murine and shark immune libraries. Molecular Immunology, 2011, 48, 2027-2037.	2.2	63
17	Serosurveillance of viral pathogens circulating in West Africa. Virology Journal, 2016, 13, 163.	3.4	57
18	Detection of dengue virus serotypes 1, 2 and 3 in selected regions of Kenya: 2011–2014. Virology Journal, 2016, 13, 182.	3.4	55

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19	Circulating microRNA profiles of Ebola virus infection. Scientific Reports, 2016, 6, 24496.	3.3	50
20	Lassa fever diagnostics: past, present, and future. Current Opinion in Virology, 2019, 37, 132-138.	5.4	47
21	Recombinant Chimeric Western and Eastern Equine Encephalitis Viruses as Potential Vaccine Candidates. Virology, 2002, 302, 299-309.	2.4	46
22	Evolution of Ebola Virus Disease from Exotic Infection to Global Health Priority, Liberia, Mid-2014. Emerging Infectious Diseases, 2015, 21, 578-584.	4.3	43
23	Evaluation of ViroCyt® Virus Counter for Rapid Filovirus Quantitation. Viruses, 2015, 7, 857-872.	3.3	42
24	Capacity building permitting comprehensive monitoring of a severe case of Lassa hemorrhagic fever in Sierra Leone with a positive outcome: Case Report. Virology Journal, 2011, 8, 314.	3.4	41
25	Detection of viral RNA from paraffin-embedded tissues after prolonged formalin fixation. Journal of Clinical Virology, 2009, 44, 39-42.	3.1	37
26	Lateral Flow Immunoassays for Ebola Virus Disease Detection in Liberia. Journal of Infectious Diseases, 2016, 214, S222-S228.	4.0	36
27	Virus-encoded miRNAs in Ebola virus disease. Scientific Reports, 2018, 8, 6480.	3.3	34
28	Comparison of MagPix Assays and Enzyme-Linked Immunosorbent Assay for Detection of Hemorrhagic Fever Viruses. Journal of Clinical Microbiology, 2017, 55, 68-78.	3.9	33
29	Directed Mutagenesis of a Sindbis Virus Pathogenesis Site. Virology, 1993, 193, 149-159.	2.4	32
30	Comparison of Transcriptomic Platforms for Analysis of Whole Blood from Ebola-Infected Cynomolgus Macaques. Scientific Reports, 2017, 7, 14756.	3.3	32
31	Persistent Crimean-Congo hemorrhagic fever virus infection in the testes and within granulomas of non-human primates with latent tuberculosis. PLoS Pathogens, 2019, 15, e1008050.	4.7	32
32	Chikungunya and O'nyong-nyong Viruses in Uganda: Implications for Diagnostics. Open Forum Infectious Diseases, 2019, 6, ofz001.	0.9	29
33	Development and Evaluation of a Panel of Filovirus Sequence Capture Probes for Pathogen Detection by Next-Generation Sequencing. PLoS ONE, 2014, 9, e107007.	2.5	28
34	Bacterial-based systems for expression and purification of recombinant Lassa virus proteins of immunological relevance. Virology Journal, 2008, 5, 74.	3.4	24
35	Shedding of soluble glycoprotein 1 detected during acute Lassa virus infection in human subjects. Virology Journal, 2010, 7, 306.	3.4	23
36	Department of Defense influenza and other respiratory disease surveillance during the 2009 pandemic. BMC Public Health, 2011, 11, S6.	2.9	20

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37	Capacity-building efforts by the AFHSC-GEIS program. BMC Public Health, 2011, 11, S4.	2.9	19
38	Infection of Aedes Albopictus and Aedes Aegypti Mosquitoes with Dengue Parent and Progeny Candidate Vaccine Viruses: a Possible Marker of Human Attenuation. American Journal of Tropical Medicine and Hygiene, 1991, 45, 202-210.	1.4	19
39	Uncoupling GP1 and GP2 expression in the Lassa virus glycoprotein complex: implications for GP1 ectodomain shedding. Virology Journal, 2008, 5, 161.	3.4	18
40	Growth and Stability of a Cholesterol-Independent Semliki Forest Virus Mutant in Mosquitoes. Virology, 1999, 262, 452-456.	2.4	16
41	Seroprevalance of Crimean–Congo haemorrhagic fever in Bulgarian livestock. Biotechnology and Biotechnological Equipment, 2014, 28, 540-542.	1.3	16
42	Detection and identification of Variola virus in fixed human tissue after prolonged archival storage. Laboratory Investigation, 2004, 84, 41-48.	3.7	15
43	Crimean-Congo Hemorrhagic Fever Virus, Mongolia, 2013–2014. Emerging Infectious Diseases, 2018, 24, 2202-2209.	4.3	14
44	Effects of La Crosse Virus Infection on Pregnant Domestic Rabbits and Mongolian Gerbils. American Journal of Tropical Medicine and Hygiene, 1996, 55, 384-390.	1.4	13
45	Dengue 3 Virus Infection of Aedes Albopictus and Aedes Aegypti: Comparison of Parent and Progeny Candidate Vaccine Viruses. American Journal of Tropical Medicine and Hygiene, 1990, 42, 89-96.	1.4	12
46	Detection of Bluetongue Virus RNA by in Situ Hybridization: Comparison with Virus Isolation and Antigen Detection. Journal of Veterinary Diagnostic Investigation, 1991, 3, 22-28.	1.1	11
47	Sindbis virus pathogenesis: phenotypic reversion of an attenuated strain to virulence by second-site intragenic suppressor mutations. Journal of General Virology, 1993, 74, 1691-1695.	2.9	11
48	Rodent-borne infections in rural Ghanaian farming communities. PLoS ONE, 2019, 14, e0215224.	2.5	11
49	Development of a bead-based immunoassay using virus-like particles for detection of alphaviral humoral response. Journal of Virological Methods, 2019, 270, 12-17.	2.1	11
50	Rapid discovery and optimization of therapeutic antibodies against emerging infectious diseases. Protein Engineering, Design and Selection, 2008, 21, 495-505.	2.1	10
51	Associations Between Antibody Fc-Mediated Effector Functions and Long-Term Sequelae in Ebola Virus Survivors. Frontiers in Immunology, 2021, 12, 682120.	4.8	9
52	Molecular Characteristics of Rickettsia in Ticks Collected along the Southern Border of Mongolia. Pathogens, 2020, 9, 943.	2.8	7
53	Enhancing laboratory capacity during Ebola virus disease (EVD) heightened surveillance in Liberia: lessons learned and recommendations. Pan African Medical Journal, 2019, 33, 8.	0.8	7
54	Comparative sequence analysis of the eastern equine encephalitis virus pathogenic strains FL91-4679 and GA97 to other north american strains. DNA Sequence, 2005, 16, 308-320.	0.7	5

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55	High heart rate at admission as a predictive factor of mortality in hospitalized patients with Lassa fever: An observational cohort study in Sierra Leone. Journal of Infection, 2020, 80, 671-693.	3.3	5
56	Molecular analysis of the 2012 Bundibugyo virus disease outbreak. Cell Reports Medicine, 2021, 2, 100351.	6.5	4
57	Evidence of presence of antibodies against selected arboviruses in Ijara and Marigat Districts, Kenya. International Journal of Infectious Diseases, 2016, 45, 188-189.	3.3	3
58	Draft Genome Sequences of Eight Crimean-Congo Hemorrhagic Fever Virus Strains. Genome Announcements, 2017, 5, .	0.8	3
59	Corning HYPERFlask® for viral amplification and production of diagnostic reagents. Journal of Virological Methods, 2017, 242, 9-13.	2.1	3
60	Conversion of a mouse Fab into a whole humanized IgG antibody for detecting botulinum toxin. Human Antibodies, 2007, 15, 125-132.	1.5	2
61	Development of a sustainable diagnostic toolbox for serosurveillance of West African infectious diseases. International Journal of Infectious Diseases, 2019, 79, 24-25.	3.3	1
62	Detection and identification of Variola virus in fixed human tissue after prolonged archival storage. Laboratory Investigation, 2004, 84, 41-48.	3.7	1
63	Specificity of molecular hybridization techniques for the detection of bluetongue virus serotypes in Culicoides variipennis. Molecular and Cellular Probes, 1992, 6, 431-438.	2.1	0
64	Development of a multiplexed antigen detection immunoassay for detection of viral agents. International Journal of Infectious Diseases, 2019, 79, 106.	3.3	0
65	The pathogenesis of genetically diverse strains of Crimean-Congo hemorrhagic fever virus in the cynomolgus macaque model. International Journal of Infectious Diseases, 2019, 79, 16.	3.3	0
66	Chikungunya and Zika Viruses Not Detected Among Patients With Dengue-Like Illness, Sarawak, Malaysia. Asia-Pacific Journal of Public Health, 2021, 33, 101053952110076.	1.0	0
67	Filoviruses. , 2014, , 65-80.		0