

# Catherine A A Beauchemin

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7467647/publications.pdf>

Version: 2024-02-01

44  
papers

2,592  
citations

218592

26  
h-index

289141

40  
g-index

47  
all docs

47  
docs citations

47  
times ranked

2457  
citing authors

#	ARTICLE	IF	CITATIONS
1	Kinetics of Influenza A Virus Infection in Humans. <i>Journal of Virology</i> , 2006, 80, 7590-7599.	1.5	630
2	A review of mathematical models of influenza A infections within a host or cell culture: lessons learned and challenges ahead. <i>BMC Public Health</i> , 2011, 11, S7.	1.2	191
3	Agent-based modeling of host-pathogen systems: The successes and challenges. <i>Information Sciences</i> , 2009, 179, 1379-1389.	4.0	174
4	A simple cellular automaton model for influenza A viral infections. <i>Journal of Theoretical Biology</i> , 2005, 232, 223-234.	0.8	154
5	Modeling amantadine treatment of influenza A virus in vitro. <i>Journal of Theoretical Biology</i> , 2008, 254, 439-451.	0.8	114
6	The H275Y Neuraminidase Mutation of the Pandemic A/H1N1 Influenza Virus Lengthens the Eclipse Phase and Reduces Viral Output of Infected Cells, Potentially Compromising Fitness in Ferrets. <i>Journal of Virology</i> , 2012, 86, 10651-10660.	1.5	99
7	Computer immunology. <i>Immunological Reviews</i> , 2007, 216, 176-197.	2.8	96
8	Assessing Mathematical Models of Influenza Infections Using Features of the Immune Response. <i>PLoS ONE</i> , 2013, 8, e57088.	1.1	96
9	Viral factors in influenza pandemic risk assessment. <i>ELife</i> , 2016, 5, .	2.8	82
10	Exploring the effect of biological delays in kinetic models of influenza within a host or cell culture. <i>BMC Public Health</i> , 2011, 11, S10.	1.2	76
11	Characterizing T Cell Movement within Lymph Nodes in the Absence of Antigen. <i>Journal of Immunology</i> , 2007, 178, 5505-5512.	0.4	74
12	Higher Level of Replication Efficiency of 2009 (H1N1) Pandemic Influenza Virus than Those of Seasonal and Avian Strains: Kinetics from Epithelial Cell Culture and Computational Modeling. <i>Journal of Virology</i> , 2011, 85, 1125-1135.	1.5	64
13	Assessing the In Vitro Fitness of an Oseltamivir-Resistant Seasonal A/H1N1 Influenza Strain Using a Mathematical Model. <i>PLoS ONE</i> , 2011, 6, e14767.	1.1	54
14	A method to determine the duration of the eclipse phase for in vitro infection with a highly pathogenic SHIV strain. <i>Scientific Reports</i> , 2015, 5, 10371.	1.6	51
15	Probing the effects of the well-mixed assumption on viral infection dynamics. <i>Journal of Theoretical Biology</i> , 2006, 242, 464-477.	0.8	50
16	Impact of the H275Y and I223V Mutations in the Neuraminidase of the 2009 Pandemic Influenza Virus In Vitro and Evaluating Experimental Reproducibility. <i>PLoS ONE</i> , 2015, 10, e0126115.	1.1	46
17	Neuraminidase inhibitors for treatment of human and avian strain influenza: A comparative modeling study. <i>Journal of Theoretical Biology</i> , 2011, 269, 234-244.	0.8	42
18	A mathematical model describing the localization and spread of influenza A virus infection within the human respiratory tract. <i>PLoS Computational Biology</i> , 2020, 16, e1007705.	1.5	39

#	ARTICLE	IF	CITATIONS
19	The I222V Neuraminidase Mutation Has a Compensatory Role in Replication of an Oseltamivir-Resistant Influenza Virus A/H3N2 E119V Mutant. <i>Journal of Clinical Microbiology</i> , 2011, 49, 715-717.	1.8	38
20	Quantification system for the viral dynamics of a highly pathogenic simian/human immunodeficiency virus based on an in vitro experiment and a mathematical model. <i>Retrovirology</i> , 2012, 9, 18.	0.9	38
21	Progress and trends in mathematical modelling of influenza A virus infections. <i>Current Opinion in Systems Biology</i> , 2018, 12, 30-36.	1.3	37
22	Avian influenza viruses that cause highly virulent infections in humans exhibit distinct replicative properties in contrast to human H1N1 viruses. <i>Scientific Reports</i> , 2016, 6, 24154.	1.6	35
23	Exploring Cell Tropism as a Possible Contributor to Influenza Infection Severity. <i>PLoS ONE</i> , 2010, 5, e13811.	1.1	30
24	Design considerations in building <i>in silico</i> equivalents of common experimental influenza virus assays. <i>Autoimmunity</i> , 2011, 44, 282-293.	1.2	30
25	Duration of SHIV production by infected cells is not exponentially distributed: Implications for estimates of infection parameters and antiviral efficacy. <i>Scientific Reports</i> , 2017, 7, 42765.	1.6	30
26	Information processing mechanisms in microtubules at physiological temperature: Model predictions for experimental tests. <i>BioSystems</i> , 2009, 97, 28-34.	0.9	29
27	Modeling Influenza Viral Dynamics in Tissue. <i>Lecture Notes in Computer Science</i> , 2006, , 23-36.	1.0	26
28	A Drug-Disease Model Describing the Effect of Oseltamivir Neuraminidase Inhibition on Influenza Virus Progression. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 5388-5395.	1.4	23
29	Modelling the emergence of influenza drug resistance: The roles of surface proteins, the immune response and antiviral mechanisms. <i>PLoS ONE</i> , 2017, 12, e0180582.	1.1	23
30	Uncovering critical properties of the human respiratory syncytial virus by combining in vitro assays and in silico analyses. <i>PLoS ONE</i> , 2019, 14, e0214708.	1.1	18
31	Time to revisit the endpoint dilution assay and to replace the TCID50 as a measure of a virus sample's infection concentration. <i>PLoS Computational Biology</i> , 2021, 17, e1009480.	1.5	17
32	(In)validating experimentally derived knowledge about influenza A defective interfering particles. <i>Journal of the Royal Society Interface</i> , 2016, 13, 20160412.	1.5	14
33	The in vivo efficacy of neuraminidase inhibitors cannot be determined from the decay rates of influenza viral titers observed in treated patients. <i>Scientific Reports</i> , 2017, 7, 40210.	1.6	14
34	Exploring virus release as a bottleneck for the spread of influenza A virus infection in vitro and the implications for antiviral therapy with neuraminidase inhibitors. <i>PLoS ONE</i> , 2017, 12, e0183621.	1.1	11
35	A conservation law for virus infection kinetics in vitro. <i>Journal of Theoretical Biology</i> , 2015, 376, 39-47.	0.8	10
36	Quantifying mechanistic traits of influenza viral dynamics using in vitro data. <i>Epidemics</i> , 2020, 33, 100406.	1.5	10

#	ARTICLE	IF	CITATIONS
37	Quantification of Ebola virus replication kinetics in vitro. PLoS Computational Biology, 2020, 16, e1008375.	1.5	10
38	Modelling endurance and resumption times for repetitive one-hand pushing. Ergonomics, 2018, 61, 891-901.	1.1	8
39	The Maximum Energy of Shock-accelerated Electrons in a Microturbulent Magnetic Field. Astrophysical Journal, 2021, 906, 33.	1.6	8
40	Agent-Based Models in Infectious Disease and Immunology. , 2015, , 38-43.		0
41	Title is missing!. , 2020, 16, e1007705.		0
42	Title is missing!. , 2020, 16, e1007705.		0
43	Title is missing!. , 2020, 16, e1007705.		0
44	Title is missing!. , 2020, 16, e1007705.		0