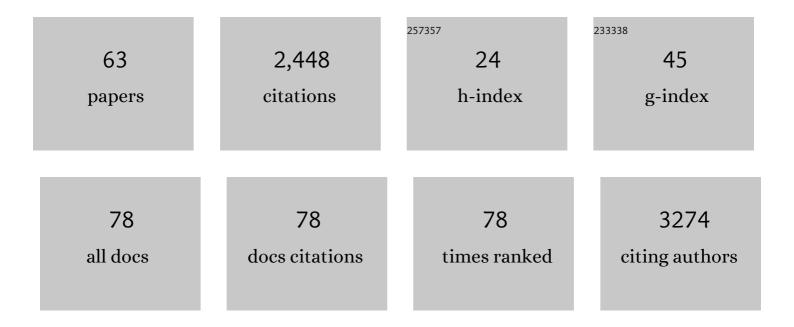
## Shelby L O'connor

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A rhesus macaque model of Asian-lineage Zika virus infection. Nature Communications, 2016, 7, 12204.	5.8	353
2	Highly efficient maternal-fetal Zika virus transmission in pregnant rhesus macaques. PLoS Pathogens, 2017, 13, e1006378.	2.1	201
3	Simian Immunodeficiency Virus SIVmac239 Infection of Major Histocompatibility Complex-Identical Cynomolgus Macaques from Mauritius. Journal of Virology, 2007, 81, 349-361.	1.5	157
4	Heterologous Protection against Asian Zika Virus Challenge in Rhesus Macaques. PLoS Neglected Tropical Diseases, 2016, 10, e0005168.	1.3	125
5	Comprehensive characterization of MHC class II haplotypes in Mauritian cynomolgus macaques. Immunogenetics, 2007, 59, 449-462.	1.2	122
6	Quantitation of Productively Infected Monocytes and Macrophages of Simian Immunodeficiency Virus-Infected Macaques. Journal of Virology, 2016, 90, 5643-5656.	1.5	93
7	Ocular and uteroplacental pathology in a macaque pregnancy with congenital Zika virus infection. PLoS ONE, 2018, 13, e0190617.	1.1	89
8	Infection via mosquito bite alters Zika virus tissue tropism and replication kinetics in rhesus macaques. Nature Communications, 2017, 8, 2096.	5.8	87
9	MHC Heterozygote Advantage in Simian Immunodeficiency Virus–Infected Mauritian Cynomolgus Macaques. Science Translational Medicine, 2010, 2, 22ra18.	5.8	80
10	Mauritian Cynomolgus Macaques Share Two Exceptionally Common Major Histocompatibility Complex Class I Alleles That Restrict Simian Immunodeficiency Virus-Specific CD8 <sup>+</sup> T Cells. Journal of Virology, 2009, 83, 6011-6019.	1.5	72
11	Zika viruses of African and Asian lineages cause fetal harm in a mouse model of vertical transmission. PLoS Neglected Tropical Diseases, 2019, 13, e0007343.	1.3	70
12	Characterization of a new SARS-CoV-2 variant that emerged in Brazil. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	63
13	Ultradeep Pyrosequencing Detects Complex Patterns of CD8 <sup>+</sup> T-Lymphocyte Escape in Simian Immunodeficiency Virus-Infected Macaques. Journal of Virology, 2009, 83, 8247-8253.	1.5	61
14	Specific CD8 <sup>+</sup> T Cell Responses Correlate with Control of Simian Immunodeficiency Virus Replication in Mauritian Cynomolgus Macaques. Journal of Virology, 2012, 86, 7596-7604.	1.5	56
15	MHC class I characterization of Indonesian cynomolgus macaques. Immunogenetics, 2008, 60, 339-51.	1.2	52
16	ALT-803 Transiently Reduces Simian Immunodeficiency Virus Replication in the Absence of Antiretroviral Treatment. Journal of Virology, 2018, 92, .	1.5	52
17	Transcriptionally Abundant Major Histocompatibility Complex Class I Alleles Are Fundamental to Nonhuman Primate Simian Immunodeficiency Virus-Specific CD8 <sup>+</sup> T Cell Responses. Journal of Virology, 2011, 85, 3250-3261.	1.5	47
18	Using barcoded Zika virus to assess virus population structure in vitro and in Aedes aegypti mosquitoes. Virology, 2018, 521, 138-148.	1.1	43

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#	Article	IF	CITATIONS
19	Characterization of 47 MHC class I sequences in Filipino cynomolgus macaques. Immunogenetics, 2009, 61, 177-187.	1.2	41
20	Revealing fine-scale spatiotemporal differences in SARS-CoV-2 introduction and spread. Nature Communications, 2020, 11, 5558.	5.8	39
21	Comparable Genital Tract Infection, Pathology, and Immunity in Rhesus Macaques Inoculated with Wild-Type or Plasmid-Deficient Chlamydia trachomatis Serovar D. Infection and Immunity, 2015, 83, 4056-4067.	1.0	38
22	Molecularly barcoded Zika virus libraries to probe in vivo evolutionary dynamics. PLoS Pathogens, 2018, 14, e1006964.	2.1	38
23	Latent Mycobacterium tuberculosis Infection Is Associated With a Higher Frequency of Mucosal-Associated Invariant T and Invariant Natural Killer T Cells. Frontiers in Immunology, 2018, 9, 1394.	2.2	33
24	Conditional CD8 <sup>+</sup> T Cell Escape during Acute Simian Immunodeficiency Virus Infection. Journal of Virology, 2012, 86, 605-609.	1.5	29
25	MAIT cells are functionally impaired in a Mauritian cynomolgus macaque model of SIV and Mtb co-infection. PLoS Pathogens, 2020, 16, e1008585.	2.1	28
26	A cautionary perspective regarding the isolation and serial propagation of SARS-CoV-2 in Vero cells. Npj Vaccines, 2021, 6, 83.	2.9	25
27	Acute-Phase CD8 T Cell Responses That Select for Escape Variants Are Needed to Control Live Attenuated Simian Immunodeficiency Virus. Journal of Virology, 2013, 87, 9353-9364.	1.5	24
28	Characterization of full-length MHC class II sequences in Indonesian and Vietnamese cynomolgus macaques. Immunogenetics, 2011, 63, 611-618.	1.2	23
29	Preexisting Simian Immunodeficiency Virus Infection Increases Susceptibility to Tuberculosis in Mauritian Cynomolgus Macaques. Infection and Immunity, 2018, 86, .	1.0	23
30	Characterization of the SARS-CoV-2 B.1.621 (Mu) variant. Science Translational Medicine, 2022, 14, eabm4908.	5.8	21
31	CD8 T Cell Response Maturation Defined by Anentropic Specificity and Repertoire Depth Correlates with SIVΔnef-induced Protection. PLoS Pathogens, 2015, 11, e1004633.	2.1	19
32	Propagation of SARS-CoV-2 in Calu-3 Cells to Eliminate Mutations in the Furin Cleavage Site of Spike. Viruses, 2021, 13, 2434.	1.5	19
33	Therapeutic Potential of IL-15 and N-803 in HIV/SIV Infection. Viruses, 2021, 13, 1750.	1.5	18
34	Vaccination with Live Attenuated Simian Immunodeficiency Virus (SIV) Protects from Mucosal, but Not Necessarily Intravenous, Challenge with a Minimally Heterologous SIV. Journal of Virology, 2016, 90, 5541-5548.	1.5	15
35	Prior infection with SARS-CoV-2 WA1/2020 partially protects rhesus macaques against reinfection with B.1.1.7 and B.1.351 variants. Science Translational Medicine, 2021, 13, eabj2641.	5.8	15
36	Acute-Phase CD4 <sup>+</sup> T Cell Responses Targeting Invariant Viral Regions Are Associated with Control of Live Attenuated Simian Immunodeficiency Virus. Journal of Virology, 2018, 92, .	1.5	13

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#	Article	IF	CITATIONS
37	Characterization of T Cells Specific for CFP-10 and ESAT-6 in Mycobacterium tuberculosis-Infected Mauritian Cynomolgus Macaques. Infection and Immunity, 2017, 85, .	1.0	12
38	Spondweni virus causes fetal harm in Ifnar1 mice and is transmitted by Aedes aegypti mosquitoes. Virology, 2020, 547, 35-46.	1.1	12
39	SIV progenitor evolution toward HIV: A humanized mouse surrogate model for SIVsm adaptation toward HIVâ€2. Journal of Medical Primatology, 2018, 47, 298-301.	0.3	11
40	Pre-existing Simian Immunodeficiency Virus Infection Increases Expression of T Cell Markers Associated with Activation during Early <i>Mycobacterium tuberculosis</i> Coinfection and Impairs TNF Responses in Granulomas. Journal of Immunology, 2021, 207, 175-188.	0.4	11
41	Initial Evaluation of a Mobile SARS-CoV-2 RT-LAMP Testing Strategy. Journal of Biomolecular Techniques, 2021, 32, 137-147.	0.8	11
42	SIV Genome-Wide Pyrosequencing Provides a Comprehensive and Unbiased View of Variation within and outside CD8 T Lymphocyte Epitopes. PLoS ONE, 2012, 7, e47818.	1.1	9
43	CD8Î <sup>2</sup> Depletion Does Not Prevent Control of Viral Replication or Protection from Challenge in Macaques Chronically Infected with a Live Attenuated Simian Immunodeficiency Virus. Journal of Virology, 2019, 93, .	1.5	9
44	SIVcpz crossâ€species transmission and viral evolution toward HIVâ€1 in a humanized mouse model. Journal of Medical Primatology, 2020, 49, 40-43.	0.3	9
45	Loss of tetherin antagonism by Nef impairs SIV replication during acute infection of rhesus macaques. PLoS Pathogens, 2020, 16, e1008487.	2.1	8
46	Spontaneous Control of SIV Replication Does Not Prevent T Cell Dysregulation and Bacterial Dissemination in Animals Co-Infected with M. tuberculosis. Microbiology Spectrum, 2022, 10, e0172421.	1.2	8
47	T cell response specificity and magnitude against SIVmac239 are not concordant in major histocompatibility complex-matched animals. Retrovirology, 2013, 10, 116.	0.9	7
48	Conditional Immune Escape during Chronic Simian Immunodeficiency Virus Infection. Journal of Virology, 2016, 90, 545-552.	1.5	6
49	Zika Virus Infection of Pregnant <i>Ifnar1</i> <sup>â^'/â^'</sup> Mice Triggers Strain-Specific Differences in Fetal Outcomes. Journal of Virology, 2021, 95, e0081821.	1.5	6
50	Acute Viral Escape Selectively Impairs Nef-Mediated Major Histocompatibility Complex Class I Downmodulation and Increases Susceptibility to Antiviral T Cells. Journal of Virology, 2016, 90, 2119-2126.	1.5	5
51	Characterization of major histocompatibility complex-related molecule 1 sequence variants in non-human primates. Immunogenetics, 2019, 71, 109-121.	1.2	5
52	Mimicking SIV chimpanzee viral evolution toward HIVâ€1 during crossâ€species transmission. Journal of Medical Primatology, 2020, 49, 284-287.	0.3	5
53	Evolution of SIVsm in humanized mice towards HIVâ€⊋. Journal of Medical Primatology, 2020, 49, 280-283.	0.3	5
54	Mathematical modeling of N-803 treatment in SIV-infected non-human primates. PLoS Computational Biology, 2021, 17, e1009204.	1.5	3

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55	Validation of multiplex PCR sequencing assay of SIV. Virology Journal, 2021, 18, 21.	1.4	2
56	Translating viral vaccines into immunity. Science, 2021, 371, 460-461.	6.0	2
57	The mucosal barrier and anti-viral immune responses can eliminate portions of the viral population during transmission and early viral growth. PLoS ONE, 2021, 16, e0260010.	1.1	1
58	Monkeying around with MAIT Cells: Studying the Role of MAIT Cells in SIV and Mtb Co-Infection. Viruses, 2021, 13, 863.	1.5	0
59	Polycystic kidney disease in rhesus macaques (Macaca mulatta). FASEB Journal, 2007, 21, A1133.	0.2	Ο
60	Title is missing!. , 2020, 16, e1008585.		0
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