## Scott Sherrill-Mix

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6681014/publications.pdf

Version: 2024-02-01

45 papers 4,248 citations

172457 29 h-index 243625 44 g-index

52 all docs 52 docs citations

times ranked

52

7738 citing authors

#	Article	IF	Citations
1	A Summary of the Sixth International Workshop on Microbiome in HIV Pathogenesis, Prevention, and Treatment. AIDS Research and Human Retroviruses, 2022, 38, 173-180.	1.1	О
2	SARS-CoV-2 Variants Associated with Vaccine Breakthrough in the Delaware Valley through Summer 2021. MBio, 2022, 13, e0378821.	4.1	11
3	SARS-CoV-2 Genomic Variation in Space and Time in Hospitalized Patients in Philadelphia. MBio, 2021, 12, .	4.1	27
4	CD4 receptor diversity represents an ancient protection mechanism against primate lentiviruses. Proceedings of the National Academy of Sciences of the United States of America, $2021,118,118$ , .	7.1	9
5	Detection of SARS-CoV-2 RNA using RT-LAMP and molecular beacons. Genome Biology, 2021, 22, 169.	8.8	61
6	Predictors of Nonseroconversion after SARS-CoV-2 Infection. Emerging Infectious Diseases, 2021, 27, 2454-2458.	4.3	48
7	Heightened resistance to host type $1$ interferons characterizes HIV- $1$ at transmission and after antiretroviral therapy interruption. Science Translational Medicine, 2021, $13$ , .	12.4	54
8	DNA binding to TLR9 expressed by red blood cells promotes innate immune activation and anemia. Science Translational Medicine, 2021, 13, eabj1008.	12.4	90
9	Molecular Beacons Allow Specific RT-LAMP Detection of B.1.1.7 Variant SARS-CoV-2. Journal of Biomolecular Techniques, 2021, 32, 98-101.	1.5	17
10	Loop-Mediated Isothermal Amplification Detection of SARS-CoV-2 and Myriad Other Applications. Journal of Biomolecular Techniques, 2021, 32, 228-275.	1.5	28
11	A Summary of the Fifth Annual Virology Education HIV Microbiome Workshop. AIDS Research and Human Retroviruses, 2020, 36, 886-895.	1.1	2
12	Single-cell transcriptional landscapes reveal HIV-1 $\hat{a}$ e "driven aberrant host gene transcription as a potential therapeutic target. Science Translational Medicine, 2020, 12, .	12.4	75
13	CpG Frequency in the $5\hat{a}\in^2$ Third of the <i>env</i> Gene Determines Sensitivity of Primary HIV-1 Strains to the Zinc-Finger Antiviral Protein. MBio, 2020, 11, .	4.1	46
14	A cone-beam computed tomographic evaluation of alveolar bone dimensional changes and the periodontal limits of mandibular incisor advancement in skeletal Class II patients. Angle Orthodontist, 2020, 90, 330-338.	2.4	25
15	The stepwise assembly of the neonatal virome is modulated by breastfeeding. Nature, 2020, 581, 470-474.	27.8	185
16	Virus structures constrain transmission modes. Nature Microbiology, 2019, 4, 1778-1780.	13.3	20
17	CD4 receptor diversity in chimpanzees protects against SIV infection. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 3229-3238.	7.1	21
18	CD19-targeting CAR T cell immunotherapy outcomes correlate with genomic modification by vector integration. Journal of Clinical Investigation, 2019, 130, 673-685.	8.2	78

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19	Allometry and Ecology of the Bilaterian Gut Microbiome. MBio, 2018, 9, .	4.1	29
20	HIV-1 latent reservoir size and diversity are stable following brief treatment interruption. Journal of Clinical Investigation, 2018, 128, 3102-3115.	8.2	88
21	Lack of detection of a human placenta microbiome in samples from preterm and term deliveries. Microbiome, 2018, 6, 196.	11.1	221
22	Resistance to type $1$ interferons is a major determinant of HIV- $1$ transmission fitness. Proceedings of the National Academy of Sciences of the United States of America, 2017, $114$ , E590-E599.	7.1	137
23	Optimizing methods and dodging pitfalls in microbiome research. Microbiome, 2017, 5, 52.	11.1	420
24	Wild bonobos host geographically restricted malaria parasites including a putative new Laverania species. Nature Communications, 2017, 8, 1635.	12.8	45
25	Guidelines for Genome-Scale Analysis of Biological Rhythms. Journal of Biological Rhythms, 2017, 32, 380-393.	2.6	237
26	Comparison of placenta samples with contamination controls does not provide evidence for a distinct placenta microbiota. Microbiome, 2016, 4, 29.	11.1	447
27	Structural Basis for Inhibitor-Induced Aggregation of HIV Integrase. PLoS Biology, 2016, 14, e1002584.	5.6	56
28	Gene activity in primary T cells infected with HIV89.6: intron retention and induction of genomic repeats. Retrovirology, 2015, 12, 79.	2.0	40
29	A Reverse Transcription Loop-Mediated Isothermal Amplification Assay Optimized to Detect Multiple HIV Subtypes. PLoS ONE, 2015, 10, e0117852.	2.5	23
30	Covalent Modification of Bacteriophage T4 DNA Inhibits CRISPR-Cas9. MBio, 2015, 6, e00648.	4.1	87
31	Power and sample-size estimation for microbiome studies using pairwise distances and PERMANOVA. Bioinformatics, 2015, 31, 2461-2468.	4.1	326
32	Phylogeography and Molecular Epidemiology of an Epidemic Strain of Dengue Virus Type $1$ in Sri Lanka. American Journal of Tropical Medicine and Hygiene, 2014, 91, 225-234.	1.4	16
33	HIV latency and integration site placement in five cell-based models. Retrovirology, 2013, 10, 90.	2.0	104
34	<i>Plasmodium falciparum</i> -like parasites infecting wild apes in southern Cameroon do not represent a recurrent source of human malaria. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 7020-7025.	7.1	53
35	Dynamic regulation of HIV-1 mRNA populations analyzed by single-molecule enrichment and long-read sequencing. Nucleic Acids Research, 2012, 40, 10345-10355.	14.5	114
36	A tool kit for quantifying eukaryotic rRNA gene sequences from human microbiome samples. Genome Biology, 2012, 13, R60.	9.6	121

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37	U1 snRNP Determines mRNA Length and Regulates Isoform Expression. Cell, 2012, 150, 53-64.	28.9	392
38	Engineering HIV-Resistant Human CD4+ T Cells with CXCR4-Specific Zinc-Finger Nucleases. PLoS Pathogens, 2011, 7, e1002020.	4.7	130
39	Hepatitis C Virus Transmission Bottlenecks Analyzed by Deep Sequencing. Journal of Virology, 2010, 84, 6218-6228.	3.4	135
40	HIV Sequence Variation Associated With env Antisense Adoptive T-cell Therapy in the hNSG Mouse Model. Molecular Therapy, 2010, 18, 803-811.	8.2	19
41	Migration cues and timing in leatherback sea turtles. Behavioral Ecology, 2008, 19, 231-236.	2.2	37
42	Evaluating potential tagging effects on leatherback sea turtles. Endangered Species Research, 2008, 4, 187-193.	2.4	17
43	Saving endangered whales at no cost. Current Biology, 2007, 17, R10-R11.	3.9	19
44	Population characteristics and seasonal migrations of leatherback sea turtles at high latitudes. Marine Ecology - Progress Series, 2007, 337, 245-254.	1.9	59
45	Canadian waters provide critical foraging habitat for leatherback sea turtles. Biological Conservation, 2006, 133, 347-357.	4.1	50