## Andrew D Davidson

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Neuropilin-1 is a host factor for SARS-CoV-2 infection. Science, 2020, 370, 861-865.	6.0	1,015
2	The furin cleavage site in the SARS-CoV-2 spike protein is required for transmission in ferrets. Nature Microbiology, 2021, 6, 899-909.	5.9	556
3	Characterisation of the transcriptome and proteome of SARS-CoV-2 reveals a cell passage induced in-frame deletion of the furin-like cleavage site from the spike glycoprotein. Genome Medicine, 2020, 12, 68.	3.6	386
4	Free fatty acid binding pocket in the locked structure of SARS-CoV-2 spike protein. Science, 2020, 370, 725-730.	6.0	348
5	The SARS-CoV-2 Spike protein disrupts human cardiac pericytes function through CD147 receptor-mediated signalling: a potential non-infective mechanism of COVID-19 microvascular disease. Clinical Science, 2021, 135, 2667-2689.	1.8	97
6	The ADP-ribose-1″-monophosphatase domains of severe acute respiratory syndrome coronavirus and human coronavirus 229E mediate resistance to antiviral interferon responses. Journal of General Virology, 2011, 92, 1899-1905.	1.3	88
7	Development of a chimeric Zika vaccine using a licensed live-attenuated flavivirus vaccine as backbone. Nature Communications, 2018, 9, 673.	5.8	84
8	The dynamics of SARS-CoV-2 infectivity with changes in aerosol microenvironment. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	84
9	KIR2DS2 recognizes conserved peptides derived from viral helicases in the context of HLA-C. Science Immunology, 2017, 2, .	5.6	78
10	Molecular Simulations suggest Vitamins, Retinoids and Steroids as Ligands of the Free Fatty Acid Pocket of the SARSâ€CoVâ€2 Spike Protein**. Angewandte Chemie - International Edition, 2021, 60, 7098-7110.	7.2	77
11	Chapter 2 New Insights into Flavivirus Nonstructural Protein 5. Advances in Virus Research, 2009, 74, 41-101.	0.9	76
12	Sensing of Immature Particles Produced by Dengue Virus Infected Cells Induces an Antiviral Response by Plasmacytoid Dendritic Cells. PLoS Pathogens, 2014, 10, e1004434.	2.1	65
13	High-Throughput Quantitative Proteomic Analysis of Dengue Virus Type 2 Infected A549 Cells. PLoS ONE, 2014, 9, e93305.	1.1	62
14	Amplicon-Based Detection and Sequencing of SARS-CoV-2 in Nasopharyngeal Swabs from Patients With COVID-19 and Identification of Deletions in the Viral Genome That Encode Proteins Involved in Interferon Antagonism. Viruses, 2020, 12, 1164.	1.5	51
15	Proteomics informed by transcriptomics for characterising active transposable elements and genome annotation in Aedes aegypti. BMC Genomics, 2017, 18, 101.	1.2	49
16	Limitations of using feline coronavirus spike protein gene mutations to diagnose feline infectious peritonitis. Veterinary Research, 2017, 48, 60.	1.1	47
17	SARS-CoV-2 vaccine ChAdOx1 nCoV-19 infection of human cell lines reveals low levels of viral backbone gene transcription alongside very high levels of SARS-CoV-2 S glycoprotein gene transcription. Genome Medicine, 2021, 13, 43.	3.6	44
18	Aedes aegypti (Aag2)-derived clonal mosquito cell lines reveal the effects of pre-existing persistent infection with the insect-specific bunyavirus Phasi Charoen-like virus on arbovirus replication. PLoS Neglected Tropical Diseases, 2019, 13, e0007346.	1.3	38

ANDREW D DAVIDSON

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19	Evaluation of the antiviral activity of orlistat (tetrahydrolipstatin) against dengue virus, Japanese encephalitis virus, Zika virus and chikungunya virus. Scientific Reports, 2020, 10, 1499.	1.6	38
20	Histidine 39 in the dengue virus type 2 M protein has an important role in virus assembly. Journal of General Virology, 2004, 85, 3627-3636.	1.3	37
21	A Modular Vaccine Platform Combining Selfâ€Assembled Peptide Cages and Immunogenic Peptides. Advanced Functional Materials, 2019, 29, 1807357.	7.8	36
22	Young infants exhibit robust functional antibody responses and restrained IFN-Î <sup>3</sup> production to SARS-CoV-2. Cell Reports Medicine, 2021, 2, 100327.	3.3	29
23	Imd pathway-specific immune assays reveal NF-κB stimulation by viral RNA PAMPs in Aedes aegypti Aag2 cells. PLoS Neglected Tropical Diseases, 2021, 15, e0008524.	1.3	28
24	BASP1 interacts with oestrogen receptor $\hat{I}\pm$ and modifies the tamoxifen response. Cell Death and Disease, 2017, 8, e2771-e2771.	2.7	26
25	Relationship between adenovirus DNA replication proteins and nucleolar proteins B23.1 and B23.2. Journal of General Virology, 2007, 88, 3244-3248.	1.3	26
26	Post-acute COVID-19 associated with evidence of bystander T-cell activation and a recurring antibiotic-resistant bacterial pneumonia. ELife, 2020, 9, .	2.8	26
27	Structural insights in cell-type specific evolution of intra-host diversity by SARS-CoV-2. Nature Communications, 2022, 13, 222.	5.8	23
28	The fatty acid site is coupled to functional motifs in the SARS-CoV-2 spike protein and modulates spike allosteric behaviour. Computational and Structural Biotechnology Journal, 2022, 20, 139-147.	1.9	19
29	ESCPE-1 mediates retrograde endosomal sorting of the SARS-CoV-2 host factor Neuropilin-1. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	17
30	Measuring stability of virus in aerosols under varying environmental conditions. Aerosol Science and Technology, 2021, 55, 1315-1320.	1.5	15
31	TMPRSS2 promotes SARS-CoV-2 evasion from NCOA7-mediated restriction. PLoS Pathogens, 2021, 17, e1009820.	2.1	13
32	Amplicon and Metagenomic Analysis of Middle East Respiratory Syndrome (MERS) Coronavirus and the Microbiome in Patients with Severe MERS. MSphere, 2021, 6, e0021921.	1.3	12
33	Identification of amino acids in the dengue virus type 2 envelope glycoprotein critical to virus infectivity. Journal of General Virology, 2009, 90, 2457-2461.	1.3	12
34	Nanopore ReCappable sequencing maps SARS-CoV-2 5′ capping sites and provides new insights into the structure of sgRNAs. Nucleic Acids Research, 2022, 50, 3475-3489.	6.5	12
35	Frontispiz: Molecular Simulations suggest Vitamins, Retinoids and Steroids as Ligands of the Free Fatty Acid Pocket of the SARS oVâ€2 Spike Protein. Angewandte Chemie, 2021, 133,	1.6	7
36	Molecular Simulations suggest Vitamins, Retinoids and Steroids as Ligands of the Free Fatty Acid Pocket of the SARS oVâ€2 Spike Protein**. Angewandte Chemie, 2021, 133, 7174-7186.	1.6	6

#	Article	IF	CITATIONS
37	Proteomics technique opens new frontiers in mobilome research. Mobile Genetic Elements, 2017, 7, 1-9.	1.8	4
38	Assessment of the red blood cell proteome in a dog with unexplained hemolytic anemia. Veterinary Clinical Pathology, 2018, 47, 377-385.	0.3	3
39	Development and Application of Dengue Virus Reverse Genetic Systems. Methods in Molecular Biology, 2014, 1138, 113-130.	0.4	2
40	Frontispiece: Molecular Simulations suggest Vitamins, Retinoids and Steroids as Ligands of the Free Fatty Acid Pocket of the SARSâ€CoVâ€2 Spike Protein. Angewandte Chemie - International Edition, 2021, 60, .	7.2	0