

# Benjamin W Neuman

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7774993/publications.pdf>

Version: 2024-02-01

53  
papers

4,326  
citations

186209

28  
h-index

182361

51  
g-index

55  
all docs

55  
docs citations

55  
times ranked

6585  
citing authors

#	ARTICLE	IF	CITATIONS
1	Ribose 2â€²-O-methylation provides a molecular signature for the distinction of self and non-self mRNA dependent on the RNA sensor Mda5. <i>Nature Immunology</i> , 2011, 12, 137-143.	7.0	640
2	A structural analysis of M protein in coronavirus assembly and morphology. <i>Journal of Structural Biology</i> , 2011, 174, 11-22.	1.3	625
3	Severe Acute Respiratory Syndrome Coronavirus Nonstructural Proteins 3, 4, and 6 Induce Double-Membrane Vesicles. <i>MBio</i> , 2013, 4, .	1.8	399
4	Supramolecular Architecture of Severe Acute Respiratory Syndrome Coronavirus Revealed by Electron Cryomicroscopy. <i>Journal of Virology</i> , 2006, 80, 7918-7928.	1.5	350
5	Structural Basis of Severe Acute Respiratory Syndrome Coronavirus ADP-Ribose-1â€³-Phosphate Dephosphorylation by a Conserved Domain of nsP3. <i>Structure</i> , 2005, 13, 1665-1675.	1.6	175
6	Proteomics Analysis Unravels the Functional Repertoire of Coronavirus Nonstructural Protein 3. <i>Journal of Virology</i> , 2008, 82, 5279-5294.	1.5	167
7	Ribonucleocapsid Formation of Severe Acute Respiratory Syndrome Coronavirus through Molecular Action of the N-Terminal Domain of N Protein. <i>Journal of Virology</i> , 2007, 81, 3913-3921.	1.5	125
8	Crystal Structure of Nonstructural Protein 10 from the Severe Acute Respiratory Syndrome Coronavirus Reveals a Novel Fold with Two Zinc-Binding Motifs. <i>Journal of Virology</i> , 2006, 80, 7894-7901.	1.5	110
9	Inhibition of Cytosolic Phospholipase A <sub>2</sub> Impairs an Early Step of Coronavirus Replication in Cell Culture. <i>Journal of Virology</i> , 2018, 92, .	1.5	107
10	Supramolecular Architecture of the Coronavirus Particle. <i>Advances in Virus Research</i> , 2016, 96, 1-27.	0.9	104
11	Inhibition, Escape, and Attenuated Growth of Severe Acute Respiratory Syndrome Coronavirus Treated with Antisense Morpholino Oligomers. <i>Journal of Virology</i> , 2005, 79, 9665-9676.	1.5	102
12	Rapid Identification of Emerging Pathogens: Coronavirus. <i>Emerging Infectious Diseases</i> , 2005, 11, 373-379.	2.0	94
13	Antiviral Effects of Antisense Morpholino Oligomers in Murine Coronavirus Infection Models. <i>Journal of Virology</i> , 2007, 81, 5637-5648.	1.5	82
14	Crystal Structure of a Monomeric Form of Severe Acute Respiratory Syndrome Coronavirus Endonuclease nsp15 Suggests a Role for Hexamerization as an Allosteric Switch. <i>Journal of Virology</i> , 2007, 81, 6700-6708.	1.5	80
15	Nuclear Magnetic Resonance Structure of the N-Terminal Domain of Nonstructural Protein 3 from the Severe Acute Respiratory Syndrome Coronavirus. <i>Journal of Virology</i> , 2007, 81, 12049-12060.	1.5	75
16	Complementarity in the Supramolecular Design of Arenaviruses and Retroviruses Revealed by Electron Cryomicroscopy and Image Analysis. <i>Journal of Virology</i> , 2005, 79, 3822-3830.	1.5	72
17	Bioinformatics and functional analyses of coronavirus nonstructural proteins involved in the formation of replicative organelles. <i>Antiviral Research</i> , 2016, 135, 97-107.	1.9	72
18	Antisense Morpholino-Oligomers Directed against the 5â€² End of the Genome Inhibit Coronavirus Proliferation and Growthâ€™. <i>Journal of Virology</i> , 2004, 78, 5891-5899.	1.5	71

#	ARTICLE	IF	CITATIONS
19	Nuclear Magnetic Resonance Structure of the Nucleic Acid-Binding Domain of Severe Acute Respiratory Syndrome Coronavirus Nonstructural Protein 3. <i>Journal of Virology</i> , 2009, 83, 12998-13008.	1.5	63
20	Description and initial characterization of metatranscriptomic nidovirus-like genomes from the proposed new family Abysoviridae, and from a sister group to the Coronavirinae, the proposed genus Alphaletovirus. <i>Virology</i> , 2018, 524, 160-171.	1.1	63
21	Structural Genomics of the Severe Acute Respiratory Syndrome Coronavirus: Nuclear Magnetic Resonance Structure of the Protein nsP7. <i>Journal of Virology</i> , 2005, 79, 12905-12913.	1.5	58
22	SARS Coronavirus Unique Domain: Three-Domain Molecular Architecture in Solution and RNA Binding. <i>Journal of Molecular Biology</i> , 2010, 400, 724-742.	2.0	54
23	Mapping the Landscape of the Lymphocytic Choriomeningitis Virus Stable Signal Peptide Reveals Novel Functional Domains. <i>Journal of Virology</i> , 2007, 81, 5649-5657.	1.5	53
24	Atlas of coronavirus replicase structure. <i>Virus Research</i> , 2014, 194, 49-66.	1.1	52
25	Nuclear Magnetic Resonance Structure Shows that the Severe Acute Respiratory Syndrome Coronavirus-Unique Domain Contains a Macrodomain Fold. <i>Journal of Virology</i> , 2009, 83, 1823-1836.	1.5	50
26	Does form meet function in the coronavirus replicative organelle?. <i>Trends in Microbiology</i> , 2014, 22, 642-647.	3.5	39
27	A Fusion Peptide in the Spike Protein of MERS Coronavirus. <i>Viruses</i> , 2019, 11, 825.	1.5	38
28	Enhancing sensitivity of lateral flow assay with application to SARS-CoV-2. <i>Applied Physics Letters</i> , 2020, 117, 120601.	1.5	34
29	Bacteriophages as promising agents for the biological control of Moko disease ( <i>Ralstonia</i> ) Tj ETQq1 1 0.784314 rgBT/Overlock 10 Tf 50 1,4 31		
30	Evaluation of SARS-CoV-2 Main Protease Inhibitors Using a Novel Cell-Based Assay. <i>ACS Central Science</i> , 2022, 8, 192-204.	5.3	30
31	Competitive Fitness in Coronaviruses Is Not Correlated with Size or Number of Double-Membrane Vesicles under Reduced-Temperature Growth Conditions. <i>MBio</i> , 2014, 5, e01107-13.	1.8	28
32	Extensive coronavirus-induced membrane rearrangements are not a determinant of pathogenicity. <i>Scientific Reports</i> , 2016, 6, 27126.	1.6	28
33	Synthesis and Antiviral Properties of Spirocyclic [1,2,3]- $\Delta^4$ -Triazolooxazine Nucleosides. <i>Chemistry - A European Journal</i> , 2014, 20, 11685-11689.	1.7	25
34	Untangling Membrane Rearrangement in the <i>Nidovirales</i> . <i>DNA and Cell Biology</i> , 2014, 33, 122-127.	0.9	24
35	Development of Peptide-Conjugated Morpholino Oligomers as Pan-Arenavirus Inhibitors. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 4631-4638.	1.4	23
36	Identification and Characterization of a Ribose 2'-O-Methyltransferase Encoded by the Ronivirus Branch of Nidovirales. <i>Journal of Virology</i> , 2016, 90, 6675-6685.	1.5	22

#	ARTICLE	IF	CITATIONS
37	Infectious Bronchitis Virus Nonstructural Protein 4 Alone Induces Membrane Pairing. <i>Viruses</i> , 2018, 10, 477.	1.5	20
38	New insights on the role of paired membrane structures in coronavirus replication. <i>Virus Research</i> , 2015, 202, 33-40.	1.1	19
39	Purification of Coronavirus Virions for Cryo-EM and Proteomic Analysis. <i>Methods in Molecular Biology</i> , 2015, 1282, 99-108.	0.4	17
40	The proteome of the infectious bronchitis virus Beau-R virion. <i>Journal of General Virology</i> , 2015, 96, 3499-3506.	1.3	15
41	Inhibition and Escape of SARS-CoV Treated with Antisense Morpholino Oligomers. <i>Advances in Experimental Medicine and Biology</i> , 2006, 581, 567-571.	0.8	11
42	Purification and Electron Cryomicroscopy of Coronavirus Particles. <i>Methods in Molecular Biology</i> , 2008, 454, 129-136.	0.4	11
43	Arenavirus budding resulting from viral-protein-associated cell membrane curvature. <i>Journal of the Royal Society Interface</i> , 2013, 10, 20130403.	1.5	10
44	Synthesis and antiviral activity of novel spirocyclic nucleosides. <i>New Journal of Chemistry</i> , 2018, 42, 18363-18380.	1.4	10
45	Isolation, Characterisation and Experimental Evolution of Phage that Infect the Horse Chestnut Tree Pathogen, <i>Pseudomonas syringae</i> pv. <i>aesculi</i> . <i>Current Microbiology</i> , 2020, 77, 1438-1447.	1.0	10
46	Ultrastructure of SARS-CoV, FIPV, and MHV Revealed by Electron Cryomicroscopy. <i>Advances in Experimental Medicine and Biology</i> , 2006, 581, 181-185.	0.8	9
47	Identification of a Membrane Binding Peptide in the Envelope Protein of MHV Coronavirus. <i>Viruses</i> , 2020, 12, 1054.	1.5	9
48	Quantum optical immunoassay: upconversion nanoparticle-based neutralizing assay for COVID-19. <i>Scientific Reports</i> , 2022, 12, 1263.	1.6	8
49	Direct Observation of Membrane Insertion by Enveloped Virus Matrix Proteins by Phosphate Displacement. <i>PLoS ONE</i> , 2013, 8, e57916.	1.1	5
50	How the Double Spherules of Infectious Bronchitis Virus Impact Our Understanding of RNA Virus Replicative Organelles. <i>MBio</i> , 2013, 4, e00987-13.	1.8	4
51	Genomic Characterisation of Mushroom Pathogenic Pseudomonads and Their Interaction with Bacteriophages. <i>Viruses</i> , 2020, 12, 1286.	1.5	3
52	Supramolecular Architecture of the Coronavirus Particle. , 2014, , 201-210.		0
53	New CLEM Method to Reveal Ultrastructural Reorganization in the Host Cell during Coronavirus Infection. <i>Microscopy and Microanalysis</i> , 2015, 21, 537-538.	0.2	0