

Martin Schwemmler

List of Publications by Year in descending order

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Version: 2024-02-01

137
papers

8,899
citations

38720

50
h-index

51562

86
g-index

151
all docs

151
docs citations

151
times ranked

10204
citing authors

#	ARTICLE	IF	CITATIONS
1	Different but Not Unique: Deciphering the Immunity of the Jamaican Fruit Bat by Studying Its Viriome. <i>Viruses</i> , 2022, 14, 238.	1.5	3
2	Biparatopic nanobodies protect mice from lethal challenge with SARS-CoV-2 variants of concern. <i>EMBO Reports</i> , 2022, 23, e53865.	2.0	18
3	Antibody escape and global spread of SARS-CoV-2 lineage A.27. <i>Nature Communications</i> , 2022, 13, 1152.	5.8	20
4	SARS-CoV-2-specific T-cell epitope repertoire in convalescent and mRNA-vaccinated individuals. <i>Nature Microbiology</i> , 2022, 7, 675-679.	5.9	29
5	Paradoxical immunodeficiencies—When failures of innate immunity cause immunopathology. <i>European Journal of Immunology</i> , 2022, 52, 1419-1430.	1.6	3
6	Are pigs overestimated as a source of zoonotic influenza viruses?. <i>Porcine Health Management</i> , 2022, 8, .	0.9	13
7	Influenza A Viruses: Understanding Human Host Determinants. <i>Trends in Molecular Medicine</i> , 2021, 27, 104-112.	3.5	24
8	Characterization of pre-existing and induced SARS-CoV-2-specific CD8+ T cells. <i>Nature Medicine</i> , 2021, 27, 78-85.	15.2	295
9	Bat-Borne Influenza A Viruses: An Awakening. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2021, 11, a038612.	2.9	6
10	Egyptian Fruit Bats (<i>Rousettus aegyptiacus</i>) Were Resistant to Experimental Inoculation with Avian-Origin Influenza A Virus of Subtype H9N2, But Are Susceptible to Experimental Infection with Bat-Borne H9N2 Virus. <i>Viruses</i> , 2021, 13, 672.	1.5	7
11	ITN—VIROINF: Understanding (Harmful) Virus-Host Interactions by Linking Virology and Bioinformatics. <i>Viruses</i> , 2021, 13, 766.	1.5	5
12	Selective Janus kinase inhibition preserves interferon-Î-mediated antiviral responses. <i>Science Immunology</i> , 2021, 6, .	5.6	16
13	Prevalence of SARS-CoV-2 Infection in Children and Their Parents in Southwest Germany. <i>JAMA Pediatrics</i> , 2021, 175, 586.	3.3	124
14	Rapid and stable mobilization of CD8+ T cells by SARS-CoV-2 mRNA vaccine. <i>Nature</i> , 2021, 597, 268-273.	13.7	279
15	BRD9 is a druggable component of interferon-stimulated gene expression and antiviral activity. <i>EMBO Reports</i> , 2021, 22, e52823.	2.0	11
16	Rare variant <i>MX1</i> alleles increase human susceptibility to zoonotic H7N9 influenza virus. <i>Science</i> , 2021, 373, 918-922.	6.0	41
17	Multisystem inflammation and susceptibility to viral infections in human ZNFX1 deficiency. <i>Journal of Allergy and Clinical Immunology</i> , 2021, 148, 381-393.	1.5	40
18	2021 Taxonomic update of phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2021, 166, 3513-3566.	0.9	62

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19	A Genome-Wide CRISPR/Cas9 Screen Reveals the Requirement of Host Sphingomyelin Synthase 1 for Infection with Pseudorabies Virus Mutant gDâ€“Pass. <i>Viruses</i> , 2021, 13, 1574.	1.5	9
20	An affinity-enhanced, broadly neutralizing heavy chainâ€“only antibody protects against SARS-CoV-2 infection in animal models. <i>Science Translational Medicine</i> , 2021, 13, eabi7826.	5.8	41
21	Within-host evolution of SARS-CoV-2 in an immunosuppressed COVID-19 patient as a source of immune escape variants. <i>Nature Communications</i> , 2021, 12, 6405.	5.8	128
22	Pre-existing immunity and vaccine history determine hemagglutinin-specific CD4 T cell and IgG response following seasonal influenza vaccination. <i>Nature Communications</i> , 2021, 12, 6720.	5.8	33
23	Influenza A Viruses and Zoonotic Eventsâ€“Are We Creating Our Own Reservoirs?. <i>Viruses</i> , 2021, 13, 2250.	1.5	26
24	Zoonotic spillover infections with Borna disease virus 1 leading to fatal human encephalitis, 1999â€“2019: an epidemiological investigation. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 467-477.	4.6	96
25	Are human Borna disease virus 1 infections zoonotic and fatal? â€“ Authors' reply. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 651.	4.6	10
26	Surveillance of European Domestic Pig Populations Identifies an Emerging Reservoir of Potentially Zoonotic Swine Influenza A Viruses. <i>Cell Host and Microbe</i> , 2020, 28, 614-627.e6.	5.1	76
27	Discrete spatio-temporal regulation of tyrosine phosphorylation directs influenza A virus M1 protein towards its function in virion assembly. <i>PLoS Pathogens</i> , 2020, 16, e1008775.	2.1	6
28	Influenza virus repurposes the antiviral protein IFIT2 to promote translation of viral mRNAs. <i>Nature Microbiology</i> , 2020, 5, 1490-1503.	5.9	45
29	2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. <i>Archives of Virology</i> , 2020, 165, 3023-3072.	0.9	184
30	A Genome-Wide CRISPR-Cas9 Screen Reveals the Requirement of Host Cell Sulfation for Schmallenberg Virus Infection. <i>Journal of Virology</i> , 2020, 94, .	1.5	18
31	Characterization of Experimental Oro-Nasal Inoculation of Sebaâ€™s Short-Tailed Bats (<i>Carollia</i>) Tj ETQq1 1 0.784314 rgBT /Overlock	1.5	5
32	Bats reveal the true power of influenza A virus adaptability. <i>PLoS Pathogens</i> , 2020, 16, e1008384.	2.1	21
33	A modified live bat influenza A virus-based vaccine prototype provides full protection against HPAIV H5N1. <i>Npj Vaccines</i> , 2020, 5, 40.	2.9	6
34	Multilineage murine stem cells generate complex organoids to model distal lung development and disease. <i>EMBO Journal</i> , 2020, 39, e103476.	3.5	44
35	Human bornavirus research: Back on track!. <i>PLoS Pathogens</i> , 2019, 15, e1007873.	2.1	41
36	Influenza restriction factor MxA functions as inflammasome sensor in the respiratory epithelium. <i>Science Immunology</i> , 2019, 4, .	5.6	39

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37	Zika Virus-Mediated Death of Hippocampal Neurons Is Independent From Maturation State. <i>Frontiers in Cellular Neuroscience</i> , 2019, 13, 389.	1.8	18
38	Bat influenza viruses transmit among bats but are poorly adapted to non-bat species. <i>Nature Microbiology</i> , 2019, 4, 2298-2309.	5.9	42
39	Poly-ADP Ribosyl Polymerase 1 (PARP1) Regulates Influenza A Virus Polymerase. <i>Advances in Virology</i> , 2019, 2019, 1-11.	0.5	13
40	Molecular mechanism for the control of virulent <i>Toxoplasma gondii</i> infections in wild-derived mice. <i>Nature Communications</i> , 2019, 10, 1233.	5.8	24
41	A protein-interaction network of interferon-stimulated genes extends the innate immune system landscape. <i>Nature Immunology</i> , 2019, 20, 493-502.	7.0	139
42	Human MxA is a potent interspecies barrier for the novel bat-derived influenza A-like virus H18N11. <i>Emerging Microbes and Infections</i> , 2019, 8, 556-563.	3.0	11
43	MHC class II proteins mediate cross-species entry of bat influenza viruses. <i>Nature</i> , 2019, 567, 109-112.	13.7	91
44	Packaging of the Influenza Virus Genome Is Governed by a Plastic Network of RNA- and Nucleoprotein-Mediated Interactions. <i>Journal of Virology</i> , 2019, 93, .	1.5	35
45	Eurasian Avian-Like Swine Influenza A Viruses Escape Human MxA Restriction through Distinct Mutations in Their Nucleoprotein. <i>Journal of Virology</i> , 2019, 93, .	1.5	26
46	Taxonomy of the order Mononegavirales: update 2018. <i>Archives of Virology</i> , 2018, 163, 2283-2294.	0.9	153
47	SMARCA2-regulated host cell factors are required for MxA restriction of influenza A viruses. <i>Scientific Reports</i> , 2018, 8, 2092.	1.6	12
48	Unexpected Functional Divergence of Bat Influenza Virus NS1 Proteins. <i>Journal of Virology</i> , 2018, 92, .	1.5	9
49	Specific Mutations in the PB2 Protein of Influenza A Virus Compensate for the Lack of Efficient Interferon Antagonism of the NS1 Protein of Bat Influenza A-Like Viruses. <i>Journal of Virology</i> , 2018, 92, .	1.5	11
50	Fatal Encephalitic Borna Disease Virus 1 in Solid-Organ Transplant Recipients. <i>New England Journal of Medicine</i> , 2018, 379, 1377-1379.	13.9	106
51	Partial Inactivation of the Chromatin Remodelers SMARCA2 and SMARCA4 in Virus-Infected Cells by Caspase-Mediated Cleavage. <i>Journal of Virology</i> , 2018, 92, .	1.5	9
52	Influenza A Virus Induces Autophagosomal Targeting of Ribosomal Proteins. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 1909-1921.	2.5	22
53	Taxonomy of the order Mononegavirales: update 2017. <i>Archives of Virology</i> , 2017, 162, 2493-2504.	0.9	173
54	In vivo evasion of MxA by avian influenza viruses requires human signature in the viral nucleoprotein. <i>Journal of Experimental Medicine</i> , 2017, 214, 1239-1248.	4.2	44

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55	Role of influenza A virus NP acetylation on viral growth and replication. <i>Nature Communications</i> , 2017, 8, 1259.	5.8	46
56	Novel insights into bat influenza A viruses. <i>Journal of General Virology</i> , 2017, 98, 2393-2400.	1.3	25
57	The Feat of Packaging Eight Unique Genome Segments. <i>Viruses</i> , 2016, 8, 165.	1.5	23
58	An RNA-dependent RNA polymerase gene in bat genomes derived from an ancient negative-strand RNA virus. <i>Scientific Reports</i> , 2016, 6, 25873.	1.6	35
59	Taxonomy of the order Mononegavirales: update 2016. <i>Archives of Virology</i> , 2016, 161, 2351-2360.	0.9	407
60	Synthetically derived bat influenza A-like viruses reveal a cell type- but not species-specific tropism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 12797-12802.	3.3	41
61	Possibility and Challenges of Conversion of Current Virus Species Names to Linnaean Binomials. <i>Systematic Biology</i> , 2016, 66, syw096.	2.7	17
62	A conserved influenza A virus nucleoprotein code controls specific viral genome packaging. <i>Nature Communications</i> , 2016, 7, 12861.	5.8	40
63	Influenza A viruses escape from MxA restriction at the expense of efficient nuclear vRNP import. <i>Scientific Reports</i> , 2016, 6, 23138.	1.6	146
64	Chiropteran influenza viruses: flu from bats or a relic from the past?. <i>Current Opinion in Virology</i> , 2016, 16, 114-119.	2.6	12
65	Mx GTPases: dynamin-like antiviral machines of innate immunity. <i>Trends in Microbiology</i> , 2015, 23, 154-163.	3.5	378
66	The Nucleoprotein of Newly Emerged H7N9 Influenza A Virus Harbors a Unique Motif Conferring Resistance to Antiviral Human MxA. <i>Journal of Virology</i> , 2015, 89, 2241-2252.	1.5	56
67	Influenza Virus Adaptation PB2-627K Modulates Nucleocapsid Inhibition by the Pathogen Sensor RIG-I. <i>Cell Host and Microbe</i> , 2015, 17, 309-319.	5.1	118
68	Taxonomic reorganization of the family Bornaviridae. <i>Archives of Virology</i> , 2015, 160, 621-632.	0.9	97
69	Generation of a variety of stable Influenza A reporter viruses by genetic engineering of the NS gene segment. <i>Scientific Reports</i> , 2015, 5, 11346.	1.6	57
70	Novel Bat Influenza Virus NS1 Proteins Bind Double-Stranded RNA and Antagonize Host Innate Immunity. <i>Journal of Virology</i> , 2015, 89, 10696-10701.	1.5	16
71	Expected and Unexpected Features of the Newly Discovered Bat Influenza A-like Viruses. <i>PLoS Pathogens</i> , 2015, 11, e1004819.	2.1	37
72	Adaptive Mutations in the Nuclear Export Protein of Human-Derived H5N1 Strains Facilitate a Polymerase Activity-Enhancing Conformation. <i>Journal of Virology</i> , 2014, 88, 263-271.	1.5	22

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73	An infectious bat-derived chimeric influenza virus harbouring the entry machinery of an influenza A virus. <i>Nature Communications</i> , 2014, 5, 4448.	5.8	80
74	Influenza, a One Health paradigm—Novel therapeutic strategies to fight a zoonotic pathogen with pandemic potential. <i>International Journal of Medical Microbiology</i> , 2014, 304, 894-901.	1.5	24
75	Phosphorylation of Highly Conserved Serine Residues in the Influenza A Virus Nuclear Export Protein NEP Plays a Minor Role in Viral Growth in Human Cells and Mice. <i>Journal of Virology</i> , 2014, 88, 7668-7673.	1.5	13
76	The Nuclear Export Protein of H5N1 Influenza A Viruses Recruits Matrix 1 (M1) Protein to the Viral Ribonucleoprotein to Mediate Nuclear Export. <i>Journal of Biological Chemistry</i> , 2014, 289, 20067-20077.	1.6	55
77	Absence of a robust innate immune response in rat neurons facilitates persistent infection of Borna disease virus in neuronal tissue. <i>Cellular and Molecular Life Sciences</i> , 2013, 70, 4399-4410.	2.4	12
78	Pandemic Influenza A Viruses Escape from Restriction by Human MxA through Adaptive Mutations in the Nucleoprotein. <i>PLoS Pathogens</i> , 2013, 9, e1003279.	2.1	156
79	Analysis of Borna Disease Virus Trafficking in Live Infected Cells by Using a Virus Encoding a Tetracycline-Tagged P Protein. <i>Journal of Virology</i> , 2013, 87, 12339-12348.	1.5	31
80	Borna disease virus-induced neuronal degeneration dependent on host genetic background and prevented by soluble factors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 1899-1904.	3.3	19
81	Adaptation of Avian Influenza A Virus Polymerase in Mammals To Overcome the Host Species Barrier. <i>Journal of Virology</i> , 2013, 87, 7200-7209.	1.5	188
82	Adaptive mutations in NEP compensate for defective H5N1 RNA replication in cultured human cells. <i>Nature Communications</i> , 2012, 3, 802.	5.8	113
83	Bornavirus Closely Associates and Segregates with Host Chromosomes to Ensure Persistent Intranuclear Infection. <i>Cell Host and Microbe</i> , 2012, 11, 492-503.	5.1	94
84	Affinity Purification of Influenza Virus Ribonucleoprotein Complexes from the Chromatin of Infected Cells. <i>Journal of Visualized Experiments</i> , 2012, , e4028.	0.2	2
85	Identification of influenza virus inhibitors which disrupt of viral polymerase protein—protein interactions. <i>Methods</i> , 2011, 55, 188-191.	1.9	27
86	Viral interference with neuronal integrity: what can we learn from the Borna disease virus?. <i>Cell and Tissue Research</i> , 2011, 344, 13-16.	1.5	2
87	The Influenza A Virus NS1 Protein Interacts with the Nucleoprotein of Viral Ribonucleoprotein Complexes. <i>Journal of Virology</i> , 2011, 85, 5228-5231.	1.5	51
88	The Viral Nucleoprotein Determines Mx Sensitivity of Influenza A Viruses. <i>Journal of Virology</i> , 2011, 85, 8133-8140.	1.5	159
89	Identification of High-Affinity PB1-Derived Peptides with Enhanced Affinity to the PA Protein of Influenza A Virus Polymerase. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 696-702.	1.4	52
90	Disruption of the Viral Polymerase Complex Assembly as a Novel Approach to Attenuate Influenza A Virus. <i>Journal of Biological Chemistry</i> , 2011, 286, 8414-8424.	1.6	31

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91	Reversion of PB2-627E to -627K during Replication of an H5N1 Clade 2.2 Virus in Mammalian Hosts Depends on the Origin of the Nucleoprotein. <i>Journal of Virology</i> , 2011, 85, 10691-10698.	1.5	52
92	Targeting of the Influenza A Virus Polymerase PB1-PB2 Interface Indicates Strain-Specific Assembly Differences. <i>Journal of Virology</i> , 2011, 85, 13298-13309.	1.5	25
93	Influenza Virus Ribonucleoprotein Complexes Gain Preferential Access to Cellular Export Machinery through Chromatin Targeting. <i>PLoS Pathogens</i> , 2011, 7, e1002187.	2.1	58
94	Protein kinase C-dependent phosphorylation of Borna disease virus P protein is required for efficient viral spread. <i>Archives of Virology</i> , 2010, 155, 789-793.	0.9	7
95	Fusion-active glycoprotein G mediates the cytotoxicity of vesicular stomatitis virus M mutants lacking host shut-off activity. <i>Journal of General Virology</i> , 2010, 91, 2782-2793.	1.3	79
96	Lambda Interferon Renders Epithelial Cells of the Respiratory and Gastrointestinal Tracts Resistant to Viral Infections. <i>Journal of Virology</i> , 2010, 84, 5670-5677.	1.5	369
97	A Polymorphism in the Hemagglutinin of the Human Isolate of a Highly Pathogenic H5N1 Influenza Virus Determines Organ Tropism in Mice. <i>Journal of Virology</i> , 2010, 84, 8316-8321.	1.5	21
98	Limited Compatibility of Polymerase Subunit Interactions in Influenza A and B Viruses. <i>Journal of Biological Chemistry</i> , 2010, 285, 16704-16712.	1.6	23
99	Identification of a PA-Binding Peptide with Inhibitory Activity against Influenza A and B Virus Replication. <i>PLoS ONE</i> , 2009, 4, e7517.	1.1	75
100	Attenuation of Rabies Virus Replication and Virulence by Picornavirus Internal Ribosome Entry Site Elements. <i>Journal of Virology</i> , 2009, 83, 1911-1919.	1.5	31
101	Mutation of the Protein Kinase C Site in Borna Disease Virus Phosphoprotein Abrogates Viral Interference with Neuronal Signaling and Restores Normal Synaptic Activity. <i>PLoS Pathogens</i> , 2009, 5, e1000425.	2.1	30
102	The Interferon Antagonist ML Protein of Thogoto Virus Targets General Transcription Factor IIB. <i>Journal of Virology</i> , 2008, 82, 11446-11453.	1.5	24
103	Functional Characterization of the Major and Minor Phosphorylation Sites of the P Protein of Borna Disease Virus. <i>Journal of Virology</i> , 2007, 81, 5497-5507.	1.5	26
104	Peptide-Mediated Interference with Influenza A Virus Polymerase. <i>Journal of Virology</i> , 2007, 81, 7801-7804.	1.5	119
105	Identification of Cellular Interaction Partners of the Influenza Virus Ribonucleoprotein Complex and Polymerase Complex Using Proteomic-Based Approaches. <i>Journal of Proteome Research</i> , 2007, 6, 672-682.	1.8	200
106	Borna Disease Virus Matrix Protein Is an Integral Component of the Viral Ribonucleoprotein Complex That Does Not Interfere with Polymerase Activity. <i>Journal of Virology</i> , 2007, 81, 743-749.	1.5	30
107	Isolation of viral ribonucleoprotein complexes from infected cells by tandem affinity purification. <i>Proteomics</i> , 2005, 5, 4483-4487.	1.3	19
108	Borna Disease Virus Replication in Organotypic Hippocampal Slice Cultures from Rats Results in Selective Damage of Dentate Granule Cells. <i>Journal of Virology</i> , 2005, 79, 11716-11723.	1.5	30

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109	The negative regulator of Borna disease virus polymerase is a non-structural protein. <i>Journal of General Virology</i> , 2005, 86, 3163-3169.	1.3	20
110	Genome trimming: A unique strategy for replication control employed by Borna disease virus. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 3441-3446.	3.3	83
111	Borna disease virus interference with neuronal plasticity. <i>Virus Research</i> , 2005, 111, 224-234.	1.1	37
112	Overlap of Interaction Domains Indicates a Central Role of the P Protein in Assembly and Regulation of the Borna Disease Virus Polymerase Complex. <i>Journal of Biological Chemistry</i> , 2004, 279, 55290-55296.	1.6	25
113	The use of peptide arrays for the characterization of monospecific antibody repertoires from polyclonal sera of psychiatric patients suspected of infection by Borna Disease Virus. <i>Molecular Diversity</i> , 2004, 8, 247-250.	2.1	9
114	Active Borna Disease Virus Polymerase Complex Requires a Distinct Nucleoprotein-to-Phosphoprotein Ratio but No Viral X Protein. <i>Journal of Virology</i> , 2003, 77, 11781-11789.	1.5	70
115	Selective Virus Resistance Conferred by Expression of Borna Disease Virus Nucleocapsid Components. <i>Journal of Virology</i> , 2003, 77, 4283-4290.	1.5	66
116	Guanylate-Binding Protein-1 Expression Is Selectively Induced by Inflammatory Cytokines and Is an Activation Marker of Endothelial Cells during Inflammatory Diseases. <i>American Journal of Pathology</i> , 2002, 161, 1749-1759.	1.9	129
117	High-avidity human serum antibodies recognizing linear epitopes of borna disease virus proteins. <i>Biological Psychiatry</i> , 2002, 51, 979-987.	0.7	68
118	Borna disease virus infection in psychiatric patients: are we on the right track?. <i>Lancet Infectious Diseases</i> , The, 2001, 1, 46-52.	4.6	50
119	Conservation of coding potential and terminal sequences in four different isolates of Borna disease virus. <i>Journal of General Virology</i> , 2001, 82, 2681-2690.	1.3	49
120	Isolation and Characterization of a New Subtype of Borna Disease Virus. <i>Journal of Virology</i> , 2000, 74, 5655-5658.	1.5	89
121	Sequence Variability of Borna Disease Virus: Resistance to Superinfection May Contribute to High Genome Stability in Persistently Infected Cells. <i>Journal of Virology</i> , 2000, 74, 7878-7883.	1.5	38
122	Authentic Borna disease virus transcripts are spliced less efficiently than cDNA-derived viral RNAs. <i>Journal of General Virology</i> , 2000, 81, 1947-1954.	1.3	12
123	Epidemiology of Borna disease virus. <i>Journal of General Virology</i> , 2000, 81, 2123-2135.	1.3	200
124	Sequence similarities between human bornavirus isolates and laboratory strains question human origin. <i>Lancet</i> , The, 1999, 354, 1973-1974.	6.3	74
125	Nucleotide-binding characteristics of human guanylate-binding protein 1 (hGBP1) and identification of the third GTP-binding motif 1 Edited by P. E. Wright. <i>Journal of Molecular Biology</i> , 1999, 292, 321-332.	2.0	114
126	Interactions of the Borna Disease Virus P, N, and X Proteins and Their Functional Implications. <i>Journal of Biological Chemistry</i> , 1998, 273, 9007-9012.	1.6	77

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127	Borna Disease Virus-Induced Neurological Disorder in Mice: Infection of Neonates Results in Immunopathology. <i>Journal of Virology</i> , 1998, 72, 4379-4386.	1.5	92
128	Borna Disease Virus P-protein Is Phosphorylated by Protein Kinase C μ and Casein Kinase II. <i>Journal of Biological Chemistry</i> , 1997, 272, 21818-21823.	1.6	63
129	Borna disease virus in brains of North American and European people with schizophrenia and bipolar disorder. <i>Lancet, The</i> , 1997, 349, 1813-1814.	6.3	105
130	GTPase properties of the interferon-induced human guanylate-binding protein 2. <i>FEBS Letters</i> , 1996, 390, 69-72.	1.3	45
131	Prenylation of an interferon- β -induced GTP-binding protein: the human guanylate binding protein, huGBP1. <i>Journal of Leukocyte Biology</i> , 1996, 60, 423-431.	1.5	60
132	Chicken Guanylate-binding Protein. <i>Journal of Biological Chemistry</i> , 1996, 271, 10304-10308.	1.6	34
133	Vesicular stomatitis virus transcription inhibited by purified MxA protein. <i>Virology</i> , 1995, 206, 545-554.	1.1	80
134	Nuclear Localization of the Interferon-Inducible Protein Kinase PKR in Human Cells and Transfected Mouse Cells. <i>Experimental Cell Research</i> , 1995, 218, 17-27.	1.2	114
135	Isolation of rat cDNA clones coding for the autoantigen SS-B/La: detection of species-specific variations. <i>Gene</i> , 1993, 126, 265-268.	1.0	28
136	Comparative analysis of the regulation of the interferoninducible protein kinase PKR by Epstein - Barr virus RNAs EBER-1 and EBER-2 and adenovirus VA, RNA. <i>Nucleic Acids Research</i> , 1993, 21, 4483-4490.	6.5	189
137	Binding of Epstein-Barr virus small RNA EBER-1 to the double-stranded RNA-activated protein kinase DAI. <i>Nucleic Acids Research</i> , 1991, 19, 243-248.	6.5	180