

John Ziebuhr

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

76
papers

9,244
citations

39
h-index

83
g-index

83
ext. papers

10,941
ext. citations

8.2
avg, IF

5.85
L-index

#	Paper	IF	Citations
76	IFITM3 Interacts with the HBV/HDV Receptor NTCP and Modulates Virus Entry and Infection.. <i>Viruses</i> , 2022 , 14,	6.2	2
75	Rocaglates as Antivirals: Comparing the Effects on Viral Resistance, Anti-Coronaviral Activity, RNA-Clamping on eIF4A and Immune Cell Toxicity.. <i>Viruses</i> , 2022 , 14,	6.2	2
74	Inhibition of SARS-CoV-2 coronavirus proliferation by designer antisense-circRNAs. <i>Nucleic Acids Research</i> , 2021 , 49, 12502-12516	20.1	2
73	Targeting the DEAD-Box RNA Helicase eIF4A with Rocaglates-A Pan-Antiviral Strategy for Minimizing the Impact of Future RNA Virus Pandemics. <i>Microorganisms</i> , 2021 , 9,	4.9	6
72	Hallmarks of and non-structural protein 7+8 complexes. <i>Science Advances</i> , 2021 , 7,	14.3	9
71	Multilevel proteomics reveals host perturbations by SARS-CoV-2 and SARS-CoV. <i>Nature</i> , 2021 , 594, 246-252	35.4	150
70	SARS-CoV-2 Variants of Interest and Concern naming scheme conducive for global discourse. <i>Nature Microbiology</i> , 2021 , 6, 821-823	26.6	91
69	Conflicting and ambiguous names of overlapping ORFs in the SARS-CoV-2 genome: A homology-based resolution. <i>Virology</i> , 2021 , 558, 145-151	3.6	15
68	Immunoglobulin deficiency as an indicator of disease severity in patients with COVID-19. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2021 , 320, L590-L599	5.8	5
67	Structure-Activity Relationships of Benzamides and Isoindolines Designed as SARS-CoV Protease Inhibitors Effective against SARS-CoV-2. <i>ChemMedChem</i> , 2021 , 16, 340-354	3.7	17
66	Coronavirus replication-transcription complex: Vital and selective NMPylation of a conserved site in nsp9 by the NiRAN-RdRp subunit. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	39
65	The rocaglate CR-31-B (-) inhibits SARS-CoV-2 replication at non-cytotoxic, low nanomolar concentrations in vitro and ex vivo. <i>Antiviral Research</i> , 2021 , 186, 105012	10.8	12
64	Reprogramming of sRNA target specificity by the leader peptide peTrpL in response to antibiotic exposure. <i>Nucleic Acids Research</i> , 2021 , 49, 2894-2915	20.1	3
63	Multi-level inhibition of coronavirus replication by chemical ER stress. <i>Nature Communications</i> , 2021 , 12, 5536	17.4	19
62	Comparison of broad-spectrum antiviral activities of the synthetic rocaglate CR-31-B (-) and the eIF4A-inhibitor Silvestrol. <i>Antiviral Research</i> , 2020 , 175, 104706	10.8	22
61	Structural basis for catalysis and substrate specificity of a 3C-like cysteine protease from a mosquito mesonivirus. <i>Virology</i> , 2019 , 533, 21-33	3.6	5
60	Transcription attenuation-derived small RNA rnTrpL regulates tryptophan biosynthesis gene expression in trans. <i>Nucleic Acids Research</i> , 2019 , 47, 6396-6410	20.1	17

59	Identification and Characterization of a Human Coronavirus 229E Nonstructural Protein 8-Associated RNA 3STerminal Adenylyltransferase Activity. <i>Journal of Virology</i> , 2019 , 93,	6.6	41
58	Direct RNA nanopore sequencing of full-length coronavirus genomes provides novel insights into structural variants and enables modification analysis. <i>Genome Research</i> , 2019 , 29, 1545-1554	9.7	112
57	Antiviral activity of K22 against members of the order Nidovirales. <i>Virus Research</i> , 2018 , 246, 28-34	6.4	14
56	Broad-spectrum antiviral activity of the eIF4A inhibitor silvestrol against corona- and picornaviruses. <i>Antiviral Research</i> , 2018 , 150, 123-129	10.8	103
55	Reverse Genetics for Type I Feline Coronavirus Field Isolate To Study the Molecular Pathogenesis of Feline Infectious Peritonitis. <i>MBio</i> , 2018 , 9,	7.8	7
54	A Single-Center Study of Viral Respiratory Tract Infections in Hospitalized Children From the Kurdistan Region of Iraq. <i>Global Pediatric Health</i> , 2018 , 5, 2333794X18784996	1.2	3
53	Characterization of a bafinivirus exoribonuclease activity. <i>Journal of General Virology</i> , 2018 , 99, 1253-1260	4.9	8
52	Inhibition of Cytosolic Phospholipase A α Impairs an Early Step of Coronavirus Replication in Cell Culture. <i>Journal of Virology</i> , 2018 , 92,	6.6	70
51	Structural and functional conservation of cis-acting RNA elements in coronavirus 5STerminal genome regions. <i>Virology</i> , 2018 , 517, 44-55	3.6	27
50	Rational Design of Novel Highly Potent and Selective Phosphatidylinositol 4-Kinase III β (PI4KB) Inhibitors as Broad-Spectrum Antiviral Agents and Tools for Chemical Biology. <i>Journal of Medicinal Chemistry</i> , 2017 , 60, 100-118	8.3	34
49	Early endonuclease-mediated evasion of RNA sensing ensures efficient coronavirus replication. <i>PLoS Pathogens</i> , 2017 , 13, e1006195	7.6	131
48	The NF- κ B-dependent and -independent transcriptome and chromatin landscapes of human coronavirus 229E-infected cells. <i>PLoS Pathogens</i> , 2017 , 13, e1006286	7.6	61
47	Identification and characterization of a Golgi retention signal in feline coronavirus accessory protein 7b. <i>Journal of General Virology</i> , 2017 , 98, 2017-2029	4.9	3
46	Characterization of the 3rd International Standard for hepatitis B virus surface antigen (HBsAg). <i>Journal of Clinical Virology</i> , 2016 , 82, 166-172	14.5	4
45	Characterization of monoclonal antibodies against feline coronavirus accessory protein 7b. <i>Veterinary Microbiology</i> , 2016 , 184, 11-9	3.3	1
44	Phylogenetic analysis of human influenza A/H3N2 viruses isolated in 2015 in Germany indicates significant genetic divergence from vaccine strains. <i>Archives of Virology</i> , 2016 , 161, 1505-15	2.6	12
43	D, L-lysine acetylsalicylate + glycine Impairs Coronavirus Replication. <i>Journal of Antivirals & Antiretrovirals</i> , 2016 , 08,	2	16
42	The PB1 segment of an influenza A virus H1N1 2009pdm isolate enhances the replication efficiency of specific influenza vaccine strains in cell culture and embryonated eggs. <i>Journal of General Virology</i> , 2016 , 97, 620-631	4.9	11

41	Proteolytic processing of mesonivirus replicase polyproteins by the viral 3C-like protease. <i>Journal of General Virology</i> , 2016 , 97, 1439-1445	4.9	5
40	Identification of specific residues in avian influenza A virus NS1 that enhance viral replication and pathogenicity in mammalian systems. <i>Journal of General Virology</i> , 2016 , 97, 2135-2148	4.9	10
39	Influenza virus-induced caspase-dependent enlargement of nuclear pores promotes nuclear export of viral ribonucleoprotein complexes. <i>Journal of Virology</i> , 2015 , 89, 6009-21	6.6	46
38	In Silico Prediction and Experimental Confirmation of HA Residues Conferring Enhanced Human Receptor Specificity of H5N1 Influenza A Viruses. <i>Scientific Reports</i> , 2015 , 5, 11434	4.9	32
37	Studies of nosocomial outbreaks of hepatitis B in nursing homes in Germany suggest a major role of hepatitis B e antigen expression in disease severity and progression. <i>International Journal of Medical Microbiology</i> , 2015 , 305, 663-72	3.7	8
36	Development and evaluation of reverse transcription loop-mediated isothermal amplification assay for the detection of the fathead minnow nidovirus. <i>Journal of Virological Methods</i> , 2014 , 202, 39-45	2.6	7
35	Coronavirus Replicative Proteins 2014 , 65-81		7
34	RNA structure analysis of alphacoronavirus terminal genome regions. <i>Virus Research</i> , 2014 , 194, 76-89	6.4	27
33	Characterization of an alphamesonivirus 3C-like protease defines a special group of nidovirus main proteases. <i>Journal of Virology</i> , 2014 , 88, 13747-58	6.6	12
32	Middle East respiratory syndrome coronavirus (MERS-CoV): announcement of the Coronavirus Study Group. <i>Journal of Virology</i> , 2013 , 87, 7790-2	6.6	796
31	Identification and characterization of genetically divergent members of the newly established family Mesoniviridae. <i>Journal of Virology</i> , 2013 , 87, 6346-58	6.6	59
30	Mesoniviridae: a proposed new family in the order Nidovirales formed by a single species of mosquito-borne viruses. <i>Archives of Virology</i> , 2012 , 157, 1623-8	2.6	89
29	The ADP-ribose-1Smonophosphatase domains of severe acute respiratory syndrome coronavirus and human coronavirus 229E mediate resistance to antiviral interferon responses. <i>Journal of General Virology</i> , 2011 , 92, 1899-1905	4.9	67
28	An insect nidovirus emerging from a primary tropical rainforest. <i>MBio</i> , 2011 , 2, e00077-11	7.8	89
27	Nidovirus ribonucleases: Structures and functions in viral replication. <i>RNA Biology</i> , 2011 , 8, 295-304	4.8	49
26	Characterization of Bafinivirus main protease autoprocessing activities. <i>Journal of Virology</i> , 2011 , 85, 1348-59	6.6	17
25	Biochemical characterization of arterivirus nonstructural protein 11 reveals the nidovirus-wide conservation of a replicative endoribonuclease. <i>Journal of Virology</i> , 2009 , 83, 5671-82	6.6	82
24	Human coronavirus 229E papain-like proteases have overlapping specificities but distinct functions in viral replication. <i>Journal of Virology</i> , 2007 , 81, 3922-32	6.6	44

23	Identification of protease and ADP-ribose 1Smonophosphatase activities associated with transmissible gastroenteritis virus non-structural protein 3. <i>Journal of General Virology</i> , 2006 , 87, 651-658	4.9	40
22	Characterization of White bream virus reveals a novel genetic cluster of nidoviruses. <i>Journal of Virology</i> , 2006 , 80, 11598-609	6.6	54
21	Discovery of an RNA virus 3S>5Sexoribonuclease that is critically involved in coronavirus RNA synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 5108-13	11.5	396
20	Crystal structure and mechanistic determinants of SARS coronavirus nonstructural protein 15 define an endoribonuclease family. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 11892-7	11.5	138
19	Nidovirales: evolving the largest RNA virus genome. <i>Virus Research</i> , 2006 , 117, 17-37	6.4	615
18	Design of wide-spectrum inhibitors targeting coronavirus main proteases. <i>PLoS Biology</i> , 2005 , 3, e324	9.7	392
17	ADP-ribose-1"-monophosphatase: a conserved coronavirus enzyme that is dispensable for viral replication in tissue culture. <i>Journal of Virology</i> , 2005 , 79, 12721-31	6.6	122
16	Major genetic marker of nidoviruses encodes a replicative endoribonuclease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 12694-9	11.5	210
15	Rapid identification of coronavirus replicase inhibitors using a selectable replicon RNA. <i>Journal of General Virology</i> , 2004 , 85, 1717-1725	4.9	64
14	Molecular biology of severe acute respiratory syndrome coronavirus. <i>Current Opinion in Microbiology</i> , 2004 , 7, 412-9	7.9	150
13	The 3C-like proteinase of an invertebrate nidovirus links coronavirus and potyvirus homologs. <i>Journal of Virology</i> , 2003 , 77, 1415-26	6.6	55
12	Unique and conserved features of genome and proteome of SARS-coronavirus, an early split-off from the coronavirus group 2 lineage. <i>Journal of Molecular Biology</i> , 2003 , 331, 991-1004	6.5	947
11	Coronavirus main proteinase (3CLpro) structure: basis for design of anti-SARS drugs. <i>Science</i> , 2003 , 300, 1763-7	33.3	1140
10	Mechanisms and enzymes involved in SARS coronavirus genome expression. <i>Journal of General Virology</i> , 2003 , 84, 2305-2315	4.9	641
9	Structure of coronavirus main proteinase reveals combination of a chymotrypsin fold with an extra alpha-helical domain. <i>EMBO Journal</i> , 2002 , 21, 3213-24	13	426
8	Mutational analysis of the active centre of coronavirus 3C-like proteases. <i>Journal of General Virology</i> , 2002 , 83, 581-593	4.9	57
7	Conservation of substrate specificities among coronavirus main proteases. <i>Journal of General Virology</i> , 2002 , 83, 595-599	4.9	190
6	The autocatalytic release of a putative RNA virus transcription factor from its polyprotein precursor involves two paralogous papain-like proteases that cleave the same peptide bond. <i>Journal of Biological Chemistry</i> , 2001 , 276, 33220-32	5.4	114

5	The human coronavirus 229E superfamily 1 helicase has RNA and DNA duplex-unwinding activities with 5Sto-3Spolarity. <i>Rna</i> , 2000 , 6, 1056-68	5.8	112
4	Biochemical characterization of the equine arteritis virus helicase suggests a close functional relationship between arterivirus and coronavirus helicases. <i>Journal of Virology</i> , 2000 , 74, 9586-93	6.6	70
3	Virus-encoded proteinases and proteolytic processing in the Nidovirales. <i>Journal of General Virology</i> , 2000 , 81, 853-79	4.9	697
2	Processing of the human coronavirus 229E replicase polyproteins by the virus-encoded 3C-like proteinase: identification of proteolytic products and cleavage sites common to pp1a and pp1ab. <i>Journal of Virology</i> , 1999 , 73, 177-85	6.6	78
1	Coronaviruses, Toroviruses, and Arteriviruses		4