

# John Ziebuhr

## List of Publications by Year in descending order

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77  
papers

12,288  
citations

66315

42  
h-index

71651

76  
g-index

83  
all docs

83  
docs citations

83  
times ranked

13782  
citing authors

#	ARTICLE	IF	CITATIONS
1	Coronavirus Main Proteinase (3CLpro) Structure: Basis for Design of Anti-SARS Drugs. <i>Science</i> , 2003, 300, 1763-1767.	6.0	1,514
2	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.	2.0	1,092
3	Commentary: Middle East Respiratory Syndrome Coronavirus (MERS-CoV): Announcement of the Coronavirus Study Group. <i>Journal of Virology</i> , 2013, 87, 7790-7792.	1.5	1,012
4	Virus-encoded proteinases and proteolytic processing in the Nidovirales. <i>Journal of General Virology</i> , 2000, 81, 853-879.	1.3	855
5	Mechanisms and enzymes involved in SARS coronavirus genome expression. <i>Journal of General Virology</i> , 2003, 84, 2305-2315.	1.3	767
6	Nidovirales: Evolving the largest RNA virus genome. <i>Virus Research</i> , 2006, 117, 17-37.	1.1	757
7	Design of Wide-Spectrum Inhibitors Targeting Coronavirus Main Proteases. <i>PLoS Biology</i> , 2005, 3, e324.	2.6	547
8	Structure of coronavirus main proteinase reveals combination of a chymotrypsin fold with an extra alpha-helical domain. <i>EMBO Journal</i> , 2002, 21, 3213-3224.	3.5	538
9	Discovery of an RNA virus 3'→5' exoribonuclease 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, 5108-5113.	3.3	524
10	Multilevel proteomics reveals host perturbations by SARS-CoV-2 and SARS-CoV. <i>Nature</i> , 2021, 594, 246-252.	13.7	475
11	Conservation of substrate specificities among coronavirus main proteases. <i>Journal of General Virology</i> , 2002, 83, 595-599.	1.3	256
12	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, 12694-12699.	3.3	254
13	SARS-CoV-2 Variants of Interest and Concern naming scheme conducive for global discourse. <i>Nature Microbiology</i> , 2021, 6, 821-823.	5.9	221
14	Molecular biology of severe acute respiratory syndrome coronavirus. <i>Current Opinion in Microbiology</i> , 2004, 7, 412-419.	2.3	192
15	Early endonuclease-mediated evasion of RNA sensing ensures efficient coronavirus replication. <i>PLoS Pathogens</i> , 2017, 13, e1006195.	2.1	184
16	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.	2.4	178
17	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, 11892-11897.	3.3	161
18	Broad-spectrum antiviral activity of the eIF4A inhibitor silvestrol against corona- and picornaviruses. <i>Antiviral Research</i> , 2018, 150, 123-129.	1.9	160

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19	ADP-Ribose-1"-Monophosphatase: a Conserved Coronavirus Enzyme That Is Dispensable for Viral Replication in Tissue Culture. <i>Journal of Virology</i> , 2005, 79, 12721-12731.	1.5	142
20	The human coronavirus 229E superfamily 1 helicase has RNA and DNA duplex-unwinding activities with 5â€²-to-3â€² polarity. <i>Rna</i> , 2000, 6, 1056-1068.	1.6	134
21	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, 33220-33232.	1.6	131
22	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
23	An Insect Nidovirus Emerging from a Primary Tropical Rainforest. <i>MBio</i> , 2011, 2, e00077-11.	1.8	100
24	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, 1623-1628.	0.9	98
25	Biochemical Characterization of Arterivirus Nonstructural Protein 11 Reveals the Nidovirus-Wide Conservation of a Replicative Endoribonuclease. <i>Journal of Virology</i> , 2009, 83, 5671-5682.	1.5	93
26	Coronavirus replicationâ€™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, .	3.3	90
27	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, 177-185.	1.5	90
28	The NF-Î±B-dependent and -independent transcriptome and chromatin landscapes of human coronavirus 229E-infected cells. <i>PLoS Pathogens</i> , 2017, 13, e1006286.	2.1	89
29	The ADP-ribose-1â€™3-monophosphatase 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.	1.3	88
30	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, 9586-9593.	1.5	78
31	Rapid identification of coronavirus replicase inhibitors using a selectable replicon RNA. <i>Journal of General Virology</i> , 2004, 85, 1717-1725.	1.3	76
32	Mutational analysis of the active centre of coronavirus 3C-like proteases. <i>Journal of General Virology</i> , 2002, 83, 581-593.	1.3	68
33	Nidovirus ribonucleases: Structures and functions in viral replication. <i>RNA Biology</i> , 2011, 8, 295-304.	1.5	68
34	Identification and Characterization of Genetically Divergent Members of the Newly Established Family Mesoniviridae. <i>Journal of Virology</i> , 2013, 87, 6346-6358.	1.5	67
35	The 3C-Like Proteinase of an Invertebrate Nidovirus Links Coronavirus and Potyvirus Homologs. <i>Journal of Virology</i> , 2003, 77, 1415-1426.	1.5	64
36	Characterization of White Bream Virus Reveals a Novel Genetic Cluster of Nidoviruses. <i>Journal of Virology</i> , 2006, 80, 11598-11609.	1.5	60

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37	Influenza Virus-Induced Caspase-Dependent Enlargement of Nuclear Pores Promotes Nuclear Export of Viral Ribonucleoprotein Complexes. <i>Journal of Virology</i> , 2015, 89, 6009-6021.	1.5	57
38	Identification and Characterization of a Human Coronavirus 229E Nonstructural Protein 8-Associated RNA 3'â€²-Terminal Adenylyltransferase Activity. <i>Journal of Virology</i> , 2019, 93, .	1.5	54
39	Multi-level inhibition of coronavirus replication by chemical ER stress. <i>Nature Communications</i> , 2021, 12, 5536.	5.8	54
40	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.	1.6	53
41	Human Coronavirus 229E Papain-Like Proteases Have Overlapping Specificities but Distinct Functions in Viral Replication. <i>Journal of Virology</i> , 2007, 81, 3922-3932.	1.5	51
42	Rational Design of Novel Highly Potent and Selective Phosphatidylinositol 4-Kinase III $\beta$ (PI4KB) Inhibitors as Broad-Spectrum Antiviral Agents and Tools for Chemical Biology. <i>Journal of Medicinal Chemistry</i> , 2017, 60, 100-118.	2.9	50
43	Structural and functional conservation of cis-acting RNA elements in coronavirus 5'-terminal genome regions. <i>Virology</i> , 2018, 517, 44-55.	1.1	46
44	RNA structure analysis of alphacoronavirus terminal genome regions. <i>Virus Research</i> , 2014, 194, 76-89.	1.1	45
45	Identification of protease and ADP-ribose 1 $\beta$ -monophosphatase activities associated with transmissible gastroenteritis virus non-structural protein 3. <i>Journal of General Virology</i> , 2006, 87, 651-656.	1.3	43
46	Conflicting and ambiguous names of overlapping ORFs in the SARS-CoV-2 genome: A homology-based resolution. <i>Virology</i> , 2021, 558, 145-151.	1.1	40
47	Comparison of broad-spectrum antiviral activities of the synthetic rocaglate CR-31-B (â€²) and the eIF4A-inhibitor Silvestrol. <i>Antiviral Research</i> , 2020, 175, 104706.	1.9	36
48	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.	1.6	36
49	Inhibition of SARS-CoV-2 coronavirus proliferation by designer antisense-circRNAs. <i>Nucleic Acids Research</i> , 2021, 49, 12502-12516.	6.5	27
50	The rocaglate CR-31-B (â€²) inhibits SARS-CoV-2 replication at non-cytotoxic, low nanomolar concentrations in vitro and ex vivo. <i>Antiviral Research</i> , 2021, 186, 105012.	1.9	26
51	Transcription attenuation-derived small RNA mTrpL regulates tryptophan biosynthesis gene expression in trans. <i>Nucleic Acids Research</i> , 2019, 47, 6396-6410.	6.5	24
52	D, L-lysine acetylsalicylate + glycine Impairs Coronavirus Replication. <i>Journal of Antivirals &amp; Antiretrovirals</i> , 2016, 08, .	0.1	21
53	Hallmarks of <i>Alpha-</i> and <i>Betacoronavirus-</i> non-structural protein 7+8 complexes. <i>Science Advances</i> , 2021, 7, .	4.7	20
54	Characterization of Bafinivirus Main Protease Autoprocessing Activities. <i>Journal of Virology</i> , 2011, 85, 1348-1359.	1.5	19

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55	Antiviral activity of K22 against members of the order Nidovirales. <i>Virus Research</i> , 2018, 246, 28-34.	1.1	17
56	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.	1.3	17
57	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.	1.3	17
58	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, 1505-1515.	0.9	16
59	Call for Papers: The Pathophysiology of COVID-19 and SARS-CoV-2 Infection. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2020, 318, L1016-L1019.	1.3	16
60	Targeting the DEAD-Box RNA Helicase eIF4A with Rocaglatesâ€™A Pan-Antiviral Strategy for Minimizing the Impact of Future RNA Virus Pandemics. <i>Microorganisms</i> , 2021, 9, 540.	1.6	16
61	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.	1.3	16
62	Characterization of an Alphamesonivirus 3C-Like Protease Defines a Special Group of Nidovirus Main Proteases. <i>Journal of Virology</i> , 2014, 88, 13747-13758.	1.5	13
63	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, 663-672.	1.5	12
64	Reverse Genetics for Type I Feline Coronavirus Field Isolate To Study the Molecular Pathogenesis of Feline Infectious Peritonitis. <i>MBio</i> , 2018, 9, .	1.8	12
65	IFITM3 Interacts with the HBV/HDV Receptor NTCP and Modulates Virus Entry and Infection. <i>Viruses</i> , 2022, 14, 727.	1.5	11
66	Coronavirus Replicative Proteins. , 2014, , 65-81.		10
67	Structural basis for catalysis and substrate specificity of a 3C-like cysteine protease from a mosquito mesonivirus. <i>Virology</i> , 2019, 533, 21-33.	1.1	10
68	Reprogramming of sRNA target specificity by the leader peptide peTrpL in response to antibiotic exposure. <i>Nucleic Acids Research</i> , 2021, 49, 2894-2915.	6.5	9
69	Identification and characterization of a Golgi retention signal in feline coronavirus accessory protein 7b. <i>Journal of General Virology</i> , 2017, 98, 2017-2029.	1.3	8
70	Characterization of a bafinivirus exoribonuclease activity. <i>Journal of General Virology</i> , 2018, 99, 1253-1260.	1.3	8
71	Thapsigargin: key to new host-directed coronavirus antivirals?. <i>Trends in Pharmacological Sciences</i> , 2022, 43, 557-568.	4.0	8
72	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.	1.0	7

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73	Characterization of the 3rd International Standard for hepatitis B virus surface antigen (HBsAg). Journal of Clinical Virology, 2016, 82, 166-172.	1.6	7
74	Proteolytic processing of mesonivirus replicase polyproteins by the viral 3C-like protease. Journal of General Virology, 2016, 97, 1439-1445.	1.3	6
75	Rocaglates as Antivirals: Comparing the Effects on Viral Resistance, Anti-Coronaviral Activity, RNA-Clamping on eIF4A and Immune Cell Toxicity. Viruses, 2022, 14, 519.	1.5	4
76	A Single-Center Study of Viral Respiratory Tract Infections in Hospitalized Children From the Kurdistan Region of Iraq. Global Pediatric Health, 2018, 5, 2333794X1878499.	0.3	3
77	Characterization of monoclonal antibodies against feline coronavirus accessory protein 7b. Veterinary Microbiology, 2016, 184, 11-19.	0.8	2