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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Nsp3 of coronaviruses: Structures and functions of a large multi-domain protein. Antiviral Research, 2018, 149, 58-74.	4.1	542
2	SARS-CoV-2 M ^{pro} inhibitors with antiviral activity in a transgenic mouse model. Science, 2021, 371, 1374-1378.	12.6	324
3	Crystal structure of Zika virus NS2B-NS3 protease in complex with a boronate inhibitor. Science, 2016, 353, 503-505.	12.6	285
4	p53 down-regulates SARS coronavirus replication and is targeted by the SARS-unique domain and PL ^{pro} via E3 ubiquitin ligase RCHY1. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E5192-201.	7.1	172
5	Structural characterization of the C-terminal domain of SARS-CoV-2 nucleocapsid protein. Molecular Biomedicine, 2020, 1, 2.	4.4	76
6	Crystal structure of the papain-like protease of MERS coronavirus reveals unusual, potentially druggable active-site features. Antiviral Research, 2014, 109, 72-82.	4.1	74
7	<scp>RNA</scp> â€virus proteases counteracting host innate immunity. FEBS Letters, 2017, 591, 3190-3210.	2.8	64
8	An orally available Mpro inhibitor is effective against wild-type SARS-CoV-2 and variants including Omicron. Nature Microbiology, 2022, 7, 716-725.	13.3	62
9	Virus–host interactomes — antiviral drug discovery. Current Opinion in Virology, 2012, 2, 614-621.	5.4	40
10	Structural and mutational analysis of the interaction between the Middle-East respiratory syndrome coronavirus (MERS-CoV) papain-like protease and human ubiquitin. Virologica Sinica, 2016, 31, 288-299.	3.0	30
11	The Structure of the Zika Virus Protease, NS2B/NS3pro. Advances in Experimental Medicine and Biology, 2018, 1062, 131-145.	1.6	28
12	The SARSâ€unique domain (SUD) of SARSâ€CoV and SARSâ€CoVâ€2 interacts with human Paip1 to enhance vira RNA translation. EMBO Journal, 2021, 40, e102277.	7.8	26
13	SARS-CoV-2 impairs the disassembly of stress granules and promotes ALS-associated amyloid aggregation. Protein and Cell, 2022, 13, 602-614.	11.0	15
14	Lybatides from Lycium barbarum Contain An Unusual Cystine-stapled Helical Peptide Scaffold. Scientific Reports, 2017, 7, 5194.	3.3	13
15	Crystal structure of the middle domain of human poly(A)-binding protein-interacting protein 1. Biochemical and Biophysical Research Communications, 2011, 408, 680-685.	2.1	5
16	Characterization of an Allosteric Pocket in Zika Virus NS2B-NS3 Protease. Journal of Chemical Information and Modeling, 2022, 62, 945-957.	5.4	4