

Margaret A Johnson

List of Publications by Year in descending order

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15
papers

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1040056

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16
docs citations

16
times ranked

1054
citing authors

#	ARTICLE	IF	CITATIONS
1	ADP-ribose and analogues bound to the deMARylating macrodomain from the bat coronavirus HKU4. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	7
2	SARS Coronavirus Unique Domain: Three-Domain Molecular Architecture in Solution and RNA Binding. , 2021, , 155-173.		0
3	Structure of the SARS-Unique Domain C From the Bat Coronavirus HKU4. Natural Product Communications, 2019, 14, 1934578X1984920.	0.5	2
4	Poly(ADP-Ribose) Polymerases in Host-Pathogen Interactions, Inflammation, and Immunity. Microbiology and Molecular Biology Reviews, 2019, 83, .	6.6	31
5	SARSâ€unique fold in the <i>Rousettus</i> bat coronavirus HKU9. Protein Science, 2017, 26, 1726-1737.	7.6	6
6	Computational and Experimental Studies of ADP-Ribosylation. Methods in Molecular Biology, 2017, 1608, 475-513.	0.9	0
7	Protein Splicing: How Inteins Escape from Precursor Proteins. Journal of Biological Chemistry, 2014, 289, 14498-14505.	3.4	115
8	SARS Coronavirus Unique Domain: Three-Domain Molecular Architecture in Solution and RNA Binding. Journal of Molecular Biology, 2010, 400, 724-742.	4.2	54
9	NMR Structure of the SARS-CoV Nonstructural Protein 7 in Solution at pH 6.5. Journal of Molecular Biology, 2010, 402, 619-628.	4.2	29
10	Nuclear Magnetic Resonance Structure Shows that the Severe Acute Respiratory Syndrome Coronavirus-Unique Domain Contains a Macrodomain Fold. Journal of Virology, 2009, 83, 1823-1836.	3.4	50
11	Structural and Functional Studies of Peptideâ€Carbohydrate Mimicry. Topics in Current Chemistry, 2008, 273, 55-116.	4.0	18
12	Proteomics Analysis Unravels the Functional Repertoire of Coronavirus Nonstructural Protein 3. Journal of Virology, 2008, 82, 5279-5294.	3.4	167
13	NMR structure of a KlbA intein precursor from Methanococcus jannaschii. Protein Science, 2007, 16, 1316-1328.	7.6	44
14	NMR assignment of a KlbA intein precursor from Methanococcus jannaschii. Biomolecular NMR Assignments, 2007, 1, 19-21.	0.8	2
15	Solution structure of Asl1650, an acyl carrier protein from Anabaena sp. PCC 7120 with a variant phosphopantetheinylation-site sequence. Protein Science, 2006, 15, 1030-1041.	7.6	19