

# Margaret A Johnson

## List of Publications by Year in descending order

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15  
papers

547  
citations

1040056  
9  
h-index

1058476  
14  
g-index

16  
all docs

16  
docs citations

16  
times ranked

1054  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteomics Analysis Unravels the Functional Repertoire of Coronavirus Nonstructural Protein 3. Journal of Virology, 2008, 82, 5279-5294.	3.4	167
2	Protein Splicing: How Inteins Escape from Precursor Proteins. Journal of Biological Chemistry, 2014, 289, 14498-14505.	3.4	115
3	SARS Coronavirus Unique Domain: Three-Domain Molecular Architecture in Solution and RNA Binding. Journal of Molecular Biology, 2010, 400, 724-742.	4.2	54
4	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
5	NMR structure of a KlbA intein precursor from <i>Methanococcus jannaschii</i> . Protein Science, 2007, 16, 1316-1328.	7.6	44
6	Poly(ADP-Ribose) Polymerases in Host-Pathogen Interactions, Inflammation, and Immunity. Microbiology and Molecular Biology Reviews, 2019, 83, .	6.6	31
7	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
8	Solution structure of Asl1650, an acyl carrier protein from <i>Anabaenasp. PCC 7120</i> with a variant phosphopantetheinylation-site sequence. Protein Science, 2006, 15, 1030-1041.	7.6	19
9	Structural and Functional Studies of Peptide-“Carbohydrate Mimicry. Topics in Current Chemistry, 2008, 273, 55-116.	4.0	18
10	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
11	SARS-unique fold in the <i>Rousettus</i> bat coronavirus HKU9. Protein Science, 2017, 26, 1726-1737.	7.6	6
12	NMR assignment of a KlbA intein precursor from <i>Methanococcus jannaschii</i> . Biomolecular NMR Assignments, 2007, 1, 19-21.	0.8	2
13	Structure of the SARS-Unique Domain C From the Bat Coronavirus HKU4. Natural Product Communications, 2019, 14, 1934578X1984920.	0.5	2
14	SARS Coronavirus Unique Domain: Three-Domain Molecular Architecture in Solution and RNA Binding. , 2021, , 155-173.	0	0
15	Computational and Experimental Studies of ADP-Ribosylation. Methods in Molecular Biology, 2017, 1608, 475-513.	0.9	0