Maria Górna

List of Publications by Year in descending order

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687363 713466 1,536 23 13 21 citations h-index g-index papers 28 28 28 2896 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The Repeating, Modular Architecture of the HtrA Proteases. Biomolecules, 2022, 12, 793.	4.0	1
2	Applications of Bacterial Degrons and Degraders â€" Toward Targeted Protein Degradation in Bacteria. Frontiers in Molecular Biosciences, 2021, 8, 669762.	3.5	22
3	A simple model for the total number of SARS-CoV-2 infections on a national level. Epidemiology and Infection, 2021, 149, e80.	2.1	6
4	The PI3K pathway preserves metabolic health through MARCO-dependent lipid uptake by adipose tissue macrophages. Nature Metabolism, 2020, 2, 1427-1442.	11.9	24
5	A Geometric Definition of Short to Medium Range Hydrogen-Mediated Interactions in Proteins. Molecules, 2020, 25, 5326.	3.8	10
6	Self-analysis of repeat proteins reveals evolutionarily conserved patterns. BMC Bioinformatics, 2020, 21, 179.	2.6	5
7	Structural, Biochemical, and Evolutionary Characterizations of Glyoxylate/Hydroxypyruvate Reductases Show Their Division into Two Distinct Subfamilies. Biochemistry, 2018, 57, 963-977.	2.5	12
8	Conservation and variability in hydrogen bonding in proteins. Acta Crystallographica Section A: Foundations and Advances, 2018, 74, e232-e232.	0.1	0
9	NDEL1-PDGFRB fusion gene in a myeloid malignancy with eosinophilia associated with resistance to tyrosine kinase inhibitors. Leukemia, 2017, 31, 237-240.	7.2	11
10	Mutations in <i>LONP1</i> , a mitochondrial matrix protease, cause CODAS syndrome. American Journal of Medical Genetics, Part A, 2015, 167, 1501-1509.	1.2	61
11	Viral RNA binding by the human IFIT1-IFIT3 protein complex in the innate immune response Acta Crystallographica Section A: Foundations and Advances, 2015, 71, s250-s250.	0.1	0
12	The SH2 Domain Regulates c-Abl Kinase Activation by a Cyclin-Like Mechanism and Remodulation of the Hinge Motion. PLoS Computational Biology, 2014, 10, e1003863.	3.2	26
13	IFITs: Emerging Roles as Key Anti-Viral Proteins. Frontiers in Immunology, 2014, 5, 94.	4.8	105
14	<i>MMP13</i> mutations are the cause of recessive metaphyseal dysplasia, Spahr type. American Journal of Medical Genetics, Part A, 2014, 164, 1175-1179.	1.2	14
15	Structural basis for viral 5′-PPP-RNA recognition by human IFIT proteins. Nature, 2013, 494, 60-64.	27.8	193
16	FAM111A Mutations Result in Hypoparathyroidism and Impaired Skeletal Development. American Journal of Human Genetics, 2013, 92, 990-995.	6.2	114
17	From conformational chaos to robust regulation: the structure and function of the multi-enzyme RNA degradosome. Quarterly Reviews of Biophysics, 2012, 45, 105-145.	5 . 7	71
18	The Seed Region of a Small RNA Drives the Controlled Destruction of the Target mRNA by the Endoribonuclease RNase E. Molecular Cell, 2012, 47, 943-953.	9.7	192

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#	Article	IF	CITATIONS
19	IFIT1 is an antiviral protein that recognizes 5′-triphosphate RNA. Nature Immunology, 2011, 12, 624-630.	14.5	422
20	Functional Dissection of the TBK1 Molecular Network. PLoS ONE, 2011, 6, e23971.	2.5	110
21	The regulatory protein RraA modulates RNA-binding and helicase activities of the <i>E. coli</i> RNA degradosome. Rna, 2010, 16, 553-562.	3.5	61
22	Reconstitution and Analysis of the Multienzyme Escherichia coli RNA Degradosome. Journal of Molecular Biology, 2008, 382, 870-883.	4.2	69
23	Design and chance in the self-assembly of macromolecules. Biochemical Society Transactions, 2007, 35, 502-507.	3.4	7