Dong Hae Shin

List of Publications by Year in descending order

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687363 361022 1,262 46 13 35 citations h-index g-index papers 46 46 46 2146 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Inhibition of SARS-CoV 3CL protease by flavonoids. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 145-151.	5.2	508
2	Characteristics of flavonoids as potent MERSâ€CoV 3Câ€like protease inhibitors. Chemical Biology and Drug Design, 2019, 94, 2023-2030.	3.2	191
3	Flavonoids with inhibitory activity against SARS-CoV-2 3CLpro. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 1539-1544.	5. 2	113
4	Structure-based functional inference in structural genomics. Journal of Structural and Functional Genomics, 2003, 4, 129-135.	1.2	60
5	Structural and Molecular Genetic Insight into a Widespread Sulfur Oxidation Pathway. Journal of Molecular Biology, 2008, 384, 1287-1300.	4.2	60
6	Crystal structure of conserved hypothetical protein Aq1575 from Aquifex aeolicus. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 7980-7985.	7.1	30
7	Structural Genomics of Minimal Organisms and Protein Fold Space. Journal of Structural and Functional Genomics, 2005, 6, 63-70.	1.2	29
8	Structure-based inference of molecular functions of proteins of unknown function from Berkeley Structural Genomics Center. Journal of Structural and Functional Genomics, 2007, 8, 99-105.	1.2	26
9	Inhibition of African swine fever virus protease by myricetin and myricitrin. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 1045-1049.	5 . 2	25
10	The Tnfaip8-PE complex is a novel upstream effector in the anti-autophagic action of insulin. Scientific Reports, 2017, 7, 6248.	3.3	21
11	Small molecule activator of Nm23/NDPK as an inhibitor of metastasis. Scientific Reports, 2018, 8, 10909.	3. 3	16
12	Structural and enzymatic characterization of DR1281: A calcineurinâ€like phosphoesterase from ⟨i⟩Deinococcus radiodurans⟨/i⟩. Proteins: Structure, Function and Bioinformatics, 2008, 70, 1000-1009.	2.6	15
13	Dimerized Translationally Controlled Tumor Protein-Binding Peptide Ameliorates Atopic Dermatitis in NC/Nga Mice. International Journal of Molecular Sciences, 2017, 18, 256.	4.1	14
14	Crystal structure of TM1457 from Thermotoga maritima. Journal of Structural Biology, 2005, 152, 113-117.	2.8	12
15	Crystal structure of a conserved hypothetical protein from Escherichia coli. Journal of Structural and Functional Genomics, 2002, 2, 53-66.	1.2	11
16	General assay for enzymes in the heptose biosynthesis pathways using electrospray ionization mass spectrometry. Applied Microbiology and Biotechnology, 2017, 101, 4521-4532.	3.6	11
17	Flexible loop and helix 2 domains of TCTP are the functional domains of dimerized TCTP. Scientific Reports, 2020, 10, 197.	3.3	11
18	Crystal structure of pharmaceutical-grade human serum albumin. International Journal of Biological Macromolecules, 2021, 166, 221-228.	7.5	10

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19	Crystal structure of the DUF16 domain of MPN010 from Mycoplasma pneumoniae. Protein Science, 2006, 15, 921-928.	7.6	7
20	Cloning, expression, purification, crystallization and preliminary X-ray diffraction analysis of DsrEFH fromAllochromatium vinosum. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 890-892.	0.7	7
21	Expression and localization of translationally controlled tumor protein in rat urinary organs. Microscopy Research and Technique, 2012, 75, 1576-1581.	2.2	7
22	Stepwise oxidations play key roles in the structural and functional regulations of DJ-1. Biochemical Journal, 2021, 478, 3505-3525.	3.7	7
23	Crystal structure of Dâ€ <i>glycero</i> â€ <i>Î'</i> â€Dâ€ <i>manno</i> â€heptoseâ€1â€phosphate adenylyltransfer from <i>Burkholderia pseudomallei</i> . Proteins: Structure, Function and Bioinformatics, 2018, 86, 124-131.	ase 2.6	6
24	Tyrosine 51 residue of the syndecan-2 extracellular domain is involved in the interaction with and activation of pro-matrix metalloproteinase-7. Scientific Reports, 2019, 9, 10625.	3.3	6
25	A Study of a Potent Inhibitor Against a GDP-6-Deoxy-α- <scp>d</scp> - <i>Manno</i> -Heptose Biosynthesis Pathway as Antibiotic Candidates. Microbial Drug Resistance, 2020, 26, 385-390.	2.0	6
26	GTP Preference of d-Glycero-α-d-manno-Heptose-1-Phosphate Guanylyltransferase from Yersinia pseudotuberculosis. International Journal of Molecular Sciences, 2020, 21, 280.	4.1	6
27	A Study of 3CLpros as Promising Targets against SARS-CoV and SARS-CoV-2. Microorganisms, 2021, 9, 756.	3.6	6
28	A Preliminary X-ray Study of Murine Tnfaip8/Oxi- \hat{l} ±. International Journal of Molecular Sciences, 2014, 15, 4523-4530.	4.1	4
29	Crystal structure of d-glycero-α-d-manno-heptose-1-phosphate guanylyltransferase from Yersinia pseudotuberculosis. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2018, 1866, 482-487.	2.3	4
30	Inhibition of $\langle scp \rangle d \langle scp \rangle - \langle i \rangle g$ lycero $\langle i \rangle - \hat{l}^2 - \langle scp \rangle d \langle scp \rangle - \langle i \rangle manno \langle i \rangle - heptose 1-phosphate adenylyltransferase from \langle i \rangle Burkholderia pseudomallei \langle i \rangle by epigallocatechin gallate and myricetin. Biochemical Journal, 2021, 478, 235-245.$	3.7	4
31	Dimerization Tendency of 3CLpros of Human Coronaviruses Based on the X-ray Crystal Structure of the Catalytic Domain of SARS-CoV-2 3CLpro. International Journal of Molecular Sciences, 2022, 23, 5268.	4.1	4
32	Preliminary structural studies on the MtxX protein from <i>Methanococcus jannaschii</i> Crystallographica Section F: Structural Biology Communications, 2008, 64, 300-303.	0.7	3
33	Expression and crystallographic studies of scp>- <i>manno</i> -heptose-1-phosphate adenylyltransferase from <i>Burkholderia pseudomallei</i> - Acta Crystallographica Section F, Structural Biology Communications, 2017, 73, 90-94.	0.8	3
34	A study of Rose Bengal against a 2-keto-3-deoxy-d-manno-octulosonate cytidylyltransferase as an antibiotic candidate. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 1414-1421.	5.2	3
35	A study of inhibitors of d-glycero- \hat{l}^2 -d-manno-heptose-1-phosphate adenylyltransferase from Burkholderia pseudomallei as a potential antibiotic target. Journal of Enzyme Inhibition and Medicinal Chemistry, 2021, 36, 776-784.	5.2	3
36	A Preliminary X-Ray Study of a Refolded PTS EIIBfruc Protein from Escherichia coli. Protein and Peptide Letters, 2008, 15, 630-632.	0.9	2

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37	A preliminary X-ray study of D, D-heptose-1, 7-bisphosphate phosphatase from Burkholderia thailandensis E264. Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 160-162.	0.7	2
38	New molecular interaction of IIA ^{Ntr} and HPr from <i>Burkholderia pseudomallei</i> identified by Xâ€ray crystallography and docking studies. Proteins: Structure, Function and Bioinformatics, 2013, 81, 1499-1508.	2.6	2
39	A preliminary X-ray study of 3-deoxy-D-manno-oct-2-ulosonic acid 8-phosphate phosphatase (YrbI) fromBurkholderia pseudomallei. Acta Crystallographica Section F, Structural Biology Communications, 2015, 71, 790-793.	0.8	2
40	Crystal Structure of Hypothetical Fructose-Specific EIIB from Escherichia coli. Molecules and Cells, 2016, 39, 495-500.	2.6	2
41	Preliminary Structural Studies on MPN423 Expressed from an Orthologous ORFan of Mycoplasma pneumoniae. Protein and Peptide Letters, 2008, 15, 753-755.	0.9	1
42	Structure and <i>in silico</i> substrate-binding mode of ADP- <scp>-<i>glycero</i>-<scp>D</scp>-<i>manno</i>-heptose 6-epimerase from<i>Burkholderia thailandensis</i>-Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 658-668.</scp>	2.5	1
43	A Study of Drug Repurposing to Identify SARS-CoV-2 Main Protease (3CLpro) Inhibitors. International Journal of Molecular Sciences, 2022, 23, 6468.	4.1	1
44	An artificially constructed dimer through deformation of a short zinc-binding loop. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2018, 1866, 205-213.	2.3	0
45	Cover Image, Volume 86, Issue 1. Proteins: Structure, Function and Bioinformatics, 2018, 86, C1.	2.6	0
46	A tripleâ€targeting inhibitory activity of Rose Bengal on polysaccharide biosynthesis of Burkholderia pseudomallei. Archiv Der Pharmazie, 2021, 354, 2000360.	4.1	O