Chi H Trinh

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4831308/publications.pdf

Version: 2024-02-01

777949 685536 24 653 13 24 citations h-index g-index papers 26 26 26 1455 docs citations citing authors all docs times ranked

#	Article	IF	CITATIONS
1	From Fragment to Lead: De Novo Design and Development toward a Selective FGFR2 Inhibitor. Journal of Medicinal Chemistry, 2022, 65, 1481-1504.	2.9	16
2	Understanding the interaction of 14â€3â€3 proteins with <i>h</i> DMX and <i>h</i> DM2: a structural and biophysical study. FEBS Journal, 2022, 289, 5341-5358.	2.2	3
3	Nanoscale Pattern Extraction from Relative Positions of Sparse 3D Localizations. Nano Letters, 2021, 21, 1213-1220.	4.5	19
4	Selective Affimers Recognise the BCLâ€2 Family Proteins BCLâ€x _L and MCLâ€1 through Noncanonical Structural Motifs**. ChemBioChem, 2021, 22, 232-240.	1.3	9
5	Identification of β-strand mediated protein–protein interaction inhibitors using ligand-directed fragment ligation. Chemical Science, 2021, 12, 2286-2293.	3.7	3
6	Substitution of histidine 30 by asparagine in manganese superoxide dismutase alters biophysical properties and supports proliferation in a K562 leukemia cell line. European Biophysics Journal, 2021, 50, 571-585.	1.2	5
7	RAS-inhibiting biologics identify and probe druggable pockets including an SII- $\hat{l}\pm 3$ allosteric site. Nature Communications, 2021, 12, 4045.	5.8	19
8	Characterization and applications of a Crimean-Congo hemorrhagic fever virus nucleoprotein-specific Affimer: Inhibitory effects in viral replication and development of colorimetric diagnostic tests. PLoS Neglected Tropical Diseases, 2020, 14, e0008364.	1.3	4
9	Directed Assembly of Homopentameric Cholera Toxin B-Subunit Proteins into Higher-Order Structures Using Coiled-Coil Appendages. Journal of the American Chemical Society, 2019, 141, 5211-5219.	6.6	18
10	Crystal structure of the putative cyclase IdmH from the indanomycin nonribosomal peptide synthase/polyketide synthase. IUCrJ, 2019, 6, 1120-1133.	1.0	8
11	A Single Mutation is Sufficient to Modify the Metal Selectivity and Specificity of a Eukaryotic Manganese Superoxide Dismutase to Encompass Iron. Chemistry - A European Journal, 2018, 24, 5303-5308.	1.7	10
12	The Structure of the Human Respiratory Syncytial Virus M2-1 Protein Bound to the Interaction Domain of the Phosphoprotein P Defines the Orientation of the Complex. MBio, $2018, 9,$	1.8	28
13	Oxygen Activation Switch in the Copper Amine Oxidase of <i>Escherichia coli</i> . Biochemistry, 2018, 57, 5301-5314.	1.2	8
14	Identification of the site of oxidase substrate binding in <i>Scytalidium thermophilum</i> catalase. Acta Crystallographica Section D: Structural Biology, 2018, 74, 979-985.	1.1	5
15	Extending enzyme molecular recognition with an expanded amino acid alphabet. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 2610-2615.	3.3	30
16	Generation of specific inhibitors of SUMO-1– and SUMO-2/3–mediated protein-protein interactions using Affimer (Adhiron) technology. Science Signaling, 2017, 10, .	1.6	44
17	Hypertrophic cardiomyopathy mutations in the calponin-homology domain of ACTN2 affect actin binding and cardiomyocyte Z-disc incorporation. Biochemical Journal, 2016, 473, 2485-2493.	1.7	42
18	Genetic analysis of Physcomitrella patens identifies ABSCISIC ACID NON-RESPONSIVE (ANR), a regulator of ABA responses unique to basal land plants and required for desiccation tolerance. Plant Cell, 2016, 28, tpc.00091.2016.	3.1	98

#	Article	IF	CITATION
19	The crystal structure of the Hazara virus nucleocapsid protein. BMC Structural Biology, 2015, 15, 24.	2.3	26
20	<scp>T</scp> he structure of the <i><scp>C</scp>aenorhabditis elegans</i> manganese superoxide dismutase <scp>M</scp> n <scp>SOD</scp> â€3â€azide complex. Protein Science, 2015, 24, 1777-1788.	3.1	18
21	Structure, recombinant expression and mutagenesis studies of the catalase with oxidase activity from <i>Scytalidium thermophilum </i> . Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 398-408.	2.5	10
22	Structures of alternatively spliced isoforms of human ketohexokinase. Acta Crystallographica Section D: Biological Crystallography, 2009, 65, 201-211.	2.5	32
23	Purification, crystallization and X-ray structures of the two manganese superoxide dismutases from <i>Caenorhabditis elegans </i> Acta Crystallographica Section F: Structural Biology Communications, 2008, 64, 1110-1114.	0.7	17
24	Crystal structure of monomeric human Â-2-microglobulin reveals clues to its amyloidogenic properties. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 9771-9776.	3.3	178