Phillip Aoto

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/726318/publications.pdf

Version: 2024-02-01

1125271 758635 19 370 12 13 citations h-index g-index papers 22 22 22 645 docs citations citing authors all docs times ranked

#	Article	IF	CITATIONS
1	Accurate scoring of non-uniform sampling schemes for quantitative NMR. Journal of Magnetic Resonance, 2014, 246, 31-35.	1.2	57
2	Globally correlated conformational entropy underlies positive and negative cooperativity in a kinase's enzymatic cycle. Nature Communications, 2019, 10, 799.	5.8	40
3	Conformation and dynamics of the kinase domain drive subcellular location and activation of LRRK2. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	35
4	Dynamic allostery-based molecular workings of kinase:peptide complexes. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 15052-15061.	3.3	33
5	Germline and Mosaic Variants in PRKACA and PRKACB Cause a Multiple Congenital Malformation Syndrome. American Journal of Human Genetics, 2020, 107, 977-988.	2.6	33
6	Structure and mechanism of monoclonal antibody binding to theÂjunctional epitope of Plasmodium falciparumÂcircumsporozoite protein. PLoS Pathogens, 2020, 16, e1008373.	2.1	30
7	Two PKA Rl \hat{l} ± holoenzyme states define ATP as an isoform-specific orthosteric inhibitor that competes with the allosteric activator, cAMP. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 16347-16356.	3.3	28
8	Defining the Structural Basis for Allosteric Product Release from <i>E. coli</i> Dihydrofolate Reductase Using NMR Relaxation Dispersion. Journal of the American Chemical Society, 2017, 139, 11233-11240.	6.6	27
9	Structural analyses of the PKA $RIll^2$ holoenzyme containing the oncogenic DnaJB1-PKAc fusion protein reveal protomer asymmetry and fusion-induced allosteric perturbations in fibrolamellar hepatocellular carcinoma. PLoS Biology, 2020, 18, e3001018.	2.6	22
10	NMR Characterization of Information Flow and Allosteric Communities in the MAP Kinase p38 \hat{l}^3 . Scientific Reports, 2016, 6, 28655.	1.6	19
11	Kinase Domain Is a Dynamic Hub for Driving LRRK2 Allostery. Frontiers in Molecular Neuroscience, 2020, 13, 538219.	1.4	18
12	LRRK2 dynamics analysis identifies allosteric control of the crosstalk between its catalytic domains. PLoS Biology, 2022, 20, e3001427.	2.6	18
13	A Dynamic Switch in Inactive $p38\hat{l}^3$ Leads to an Excited State on the Pathway to an Active Kinase. Biochemistry, 2019, 58, 5160-5172.	1.2	7
14	Title is missing!. , 2020, 18, e3001018.		0
15	Title is missing!. , 2020, 18, e3001018.		О
16	Title is missing!. , 2020, 18, e3001018.		0
17	Title is missing!. , 2020, 18, e3001018.		O
18	Title is missing!. , 2020, 18, e3001018.		0

ARTICLE IF CITATIONS

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