

Luis P B Scott

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

262
citations

933447

10
h-index

996975

15
g-index

22
all docs

22
docs citations

22
times ranked

942
citing authors

#	ARTICLE	IF	CITATIONS
1	Virtual screening and inÂvitro assays of novel hits as promising DPP-4 inhibitors. <i>Biochimie</i> , 2022, 194, 43-50.	2.6	2
2	Sampling of Protein Conformational Space Using Hybrid Simulations: A Critical Assessment of Recent Methods. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 832847.	3.5	14
3	Structural modelling and thermostability of a serine protease inhibitor belonging to the Kunitz-BPTI family from the <i>Rhipicephalus microplus</i> tick. <i>Biochimie</i> , 2021, 181, 226-233.	2.6	4
4	Nifuroxazide as JAK2 inhibitor: A binding mode proposal and Hel cell proliferation assay. <i>European Journal of Pharmaceutical Sciences</i> , 2021, 162, 105822.	4.0	3
5	Towards gaining sight of multiscale events: utilizing network models and normal modes in hybrid methods. <i>Current Opinion in Structural Biology</i> , 2020, 64, 34-41.	5.7	32
6	Unveiling functional motions based on point mutations in biased signaling systems: A normal mode study on nerve growth factor bound to TrkA. <i>PLoS ONE</i> , 2020, 15, e0231542.	2.5	5
7	Integrated Protocol to Design Potential Inhibitors of Dipeptidyl Peptidase- 4 (DPP-4). <i>Current Topics in Medicinal Chemistry</i> , 2020, 20, 209-226.	2.1	2
8	Methyl divanillate: redox properties and binding affinity with albumin of an antioxidant and potential NADPH oxidase inhibitor. <i>RSC Advances</i> , 2019, 9, 19983-19992.	3.6	6
9	Predicting regions prone to protein aggregation based on SVM algorithm. <i>Applied Mathematics and Computation</i> , 2019, 359, 502-511.	2.2	9
10	Interaction between 1-pyrenesulfonic acid and albumin: Moving inside the protein. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2019, 208, 243-254.	3.9	14
11	PINK1 Interacts with VCP/p97 and Activates PKA to Promote NSFL1C/p47 Phosphorylation and Dendritic Arborization in Neurons. <i>ENeuro</i> , 2018, 5, ENEURO.0466-18.2018.	1.9	34
12	Structural Dynamics of DPP-4 and Its Influence on the Projection of Bioactive Ligands. <i>Molecules</i> , 2018, 23, 490.	3.8	25
13	UDP-glucuronosyltransferases: Structure, Function and Drug Design Studies. <i>Current Medicinal Chemistry</i> , 2018, 25, 3247-3255.	2.4	33
14	The Role of QSAR and Virtual Screening Studies in Type 2 Diabetes Drug Discovery. <i>Medicinal Chemistry</i> , 2017, 13, 706-720.	1.5	7
15	Structure and functional dynamics characterization of the ion channel of the human respiratory syncytial virus (hRSV) small hydrophobic protein (SH) transmembrane domain by combining molecular dynamics with excited normal modes. <i>Journal of Molecular Modeling</i> , 2016, 22, 286.	1.8	13
16	Recovery of the wild type atomic flexibility in the HIV-1 protease double mutants. <i>Journal of Molecular Graphics and Modelling</i> , 2015, 59, 107-116.	2.4	3
17	A cell-based reporter assay for screening for EcR agonist/antagonist activity of natural ecdysteroids in Lepidoptera (Bm5) and Diptera (S2) cell cultures, followed by modeling of ecdysteroid-EcR interactions and normal mode analysis. <i>Pesticide Biochemistry and Physiology</i> , 2013, 107, 309-320.	3.6	16
18	Self-Assembly of Argâ€Phe Nanostructures via the Solidâ€Vapor Phase Method. <i>Journal of Physical Chemistry B</i> , 2013, 117, 733-740.	2.6	27

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19	Relation between flexibility and positively selected HIV-1 protease mutants against inhibitors. Proteins: Structure, Function and Bioinformatics, 2012, 80, 2680-2691.	2.6	13