Napat Songtawee

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/448578/publications.pdf

Version: 2024-02-01

22 338 11 papers citations h-index

23 23 23 613 all docs docs citations times ranked citing authors

18

g-index

#	Article	IF	CITATIONS
1	Extracellular Vesicle-Mediated IL-1 Signaling in Response to Doxorubicin Activates PD-L1 Expression in Osteosarcoma Models. Cells, 2022, 11, 1042.	4.1	11
2	Identification of tripeptides against tyrosine kinase domain of EGFR for lung cancer cell inhibition by in silico and in vitro studies. Chemical Biology and Drug Design, 2022, 99, 456-469.	3.2	5
3	Modulatory Effects of Alpha-Mangostin Mediated by SIRT1/3-FOXO3a Pathway in Oxidative Stress-Induced Neuronal Cells. Frontiers in Nutrition, 2021, 8, 714463.	3.7	9
4	Butein, isoliquiritigenin, and scopoletin attenuate neurodegeneration <i>via</i> antioxidant enzymes and SIRT1/ADAM10 signaling pathway. RSC Advances, 2020, 10, 16593-16606.	3.6	20
5	Potential tripeptides against the tyrosine kinase domain of human epidermal growth factor receptor (HER) 2 through computational and kinase assay approaches. Journal of Molecular Graphics and Modelling, 2020, 97, 107564.	2.4	4
6	Repurposing of Nitroxoline Drug for the Prevention of Neurodegeneration. Chemical Research in Toxicology, 2019, 32, 2182-2191.	3.3	22
7	Yield improvement and enzymatic dissection of Plasmodium falciparum plasmepsin V. Molecular and Biochemical Parasitology, 2019, 231, 111188.	1.1	5
8	Production and Characterization of Recombinant Wild Type Uricase from Indonesian Coelacanth (L.) Tj ETQq0 0 Bridges Engineering. International Journal of Molecular Sciences, 2019, 20, 1269.	0 rgBT /Ov 4.1	verlock 10 Tf 5
9	Rational Design of Colchicine Derivatives as anti-HIV Agents via QSAR and Molecular Docking. Medicinal Chemistry, 2019, 15, 328-340.	1.5	17
10	Neuroprotective Effects of Phenolic and Carboxylic Acids on Oxidative Stress-Induced Toxicity in Human Neuroblastoma SH-SY5Y Cells. Neurochemical Research, 2018, 43, 619-636.	3.3	63
11	Molecular dynamics simulations of asymmetric heterodimers of HER1/HER2 complexes. Journal of Molecular Modeling, 2018, 24, 30.	1.8	1
12	InÂvitro and in silico studies of naphthoquinones and peptidomimetics toward Plasmodium falciparum plasmepsin V. Biochimie, 2018, 152, 159-173.	2.6	9
13	Improving enzymatic activities and thermostability of a tri-functional enzyme with SOD, catalase and cell-permeable activities. Journal of Biotechnology, 2017, 247, 50-59.	3.8	7
14	Engineering of a novel tri-functional enzyme with MnSOD, catalase and cell-permeable activities. International Journal of Biological Macromolecules, 2016, 85, 451-459.	7.5	6
15	Structural and biochemical characterization of two heme binding sites on α 1 -microglobulin using site directed mutagenesis and molecular simulation. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2016, 1864, 29-41.	2.3	20
16	Navigating the chemical space of dipeptidyl peptidase-4 inhibitors. Drug Design, Development and Therapy, 2015, 9, 4515.	4.3	20
17	Quinoline-based clioquinol and nitroxoline exhibit anticancer activity inducing FoxM1 inhibition in cholangiocarcinoma cells. Drug Design, Development and Therapy, 2015, 9, 2033.	4.3	34
18	Understanding the molecular basis of EGFR kinase domain/MIG-6 peptide recognition complex using computational analyses. BMC Bioinformatics, 2015, 16, 103.	2.6	18

#	Article	IF	CITATION
19	Molecular dynamics of the asymmetric dimers of EGFR: Simulations on the active and inactive conformations of the kinase domain. Journal of Molecular Graphics and Modelling, 2015, 58, 16-29.	2.4	11
20	Probing the origins of $17\hat{l}^2$ -hydroxysteroid dehydrogenase type 1 inhibitory activity via QSAR and molecular docking. European Journal of Medicinal Chemistry, 2015, 96, 231-237.	5. 5	4
21	Molecular characterization of Galectin-8 from Nile tilapia (Oreochromis niloticus Linn.) and its response to bacterial infection. Molecular Immunology, 2015, 68, 585-596.	2.2	16
22	Computational study of EGFR inhibition: molecular dynamics studies on the active and inactive protein conformations. Journal of Molecular Modeling, 2013, 19, 497-509.	1.8	22