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Automated docking using a Lamarckian genetic algorithm and an empirical binding free energy function

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270	Quantum chemical, spectroscopic and molecular docking investigations of potential pulmonary fibrosis drug methyl 2-chloro 4-iodonicotinate.	0
269	A New Naphthopyran Derivative Combines c-Myb Inhibition, Microtubule-Targeting Effects, and Antiangiogenic Properties.	1
268	An Evaluation of the Novel Biological Properties of Diterpenes Isolated from Plectranthus ornatus Codd. In Vitro and In Silico. <b>2022</b> , 11, 3243	0
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265	Oxygen mediated oxidative couplings of flavones in alkaline water. <b>2022</b> , 13,	0
264	Structure-Based Virtual Screening and De Novo Design to Identify Submicromolar Inhibitors of G2019S Mutant of Leucine-Rich Repeat Kinase 2. <b>2022</b> , 23, 12825	1
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237	Ultra-Small NIR J-aggregates of BODIPY for Potent Phototheranostics.	0
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233	Insight into the Interaction Mechanism of HSA with Aztreonam: A Multispectroscopic and Computational Approach. <b>2022</b> , 27, 7858	1
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229	Guava Leaf Essential Oil as a Potent Antioxidant and Anticancer Agent: Validated through Experimental and Computational Study. <b>2022</b> , 11, 2204	1
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224	Binding Properties of Photosynthetic Herbicides: Photosynthetic Activity and Molecular Docking Approach towards 1,4-Dihydropyridines Derivatives.	0

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222	An immunoinformatic approach employing molecular docking and molecular dynamics simulation for evaluation of l-asparaginase produced by <i>Bacillus velezensis</i> . 1-15	0
221	Amine-Reactive BODIPY Dye: Spectral Properties and Application for Protein Labeling. <b>2022</b> , 27, 7911	0
220	Antiviral activity of <i>Turbinaria ornata</i> against white spot syndrome virus in freshwater crab ( <i>Paratelphusa hydrodromous</i> ).	0
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211	Icotinib derivatives as tyrosine kinase inhibitors with anti-esophageal squamous carcinoma activity. 13,	0
210	Computational drug discovery under RNA times. 1-21	0
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183	New Ni(II) and Pd(II) complexes bearing derived sulfa drug ligands: synthesis, characterization, DFT calculations, and in silico and in vitro biological activity studies.	0
182	Zinc sulfide nano aqua formulation as antifungal nanopriming agent against phytopathogens of paddy seeds: in silico, in vitro and seed treatment studies.	1
181	An in-silico investigation of fluoride ions impact on pancreatic lipase.	0
180	Isolation, purification and biochemical characterization of Conopomorpha cramerella farnesol dehydrogenase.	0
179	Design, synthesis, characterization and analysis of anti-inflammatory properties of novel N-(benzo[d]thiazol-2-yl)-2-[phenyl(2-(piperidin-1-yl) ethylamino) benzamides and N-(benzo[d]thiazol-2-yl)-2-[phenyl (2-morpholino) ethylamino] benzamides derivatives through in vitro and in silico approach.	0
178	Amino carbonyl derivatives: Synthesis, Molecular Docking, ADMET, Molecular Dynamic and Herbicidal studies.. <b>2022</b> , 7,	3
177	Alpha-Lipoic Acid and Its Enantiomers Prevent Methemoglobin Formation and DNA Damage Induced by Dapsone Hydroxylamine: Molecular Mechanism and Antioxidant Action. <b>2023</b> , 24, 57	0
176	(-)-Epigallocatechin-3-gallate Directly Binds Cyclophilin D: A Potential Mechanism for Mitochondrial Protection. <b>2022</b> , 27, 8661	0
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172	SARS-CoV-2 main protease inhibitors: Structure-based enhancement to anti-viral pre-clinical GC376 encourages further development.	0
171	Chemical Survey of Three Species of the Genus <i>Rauhia</i> Traub (Amaryllidaceae). <b>2022</b> , 11, 3549	2
170	Molecular spectroscopic and docking analysis of the interaction of fluorescent thiadicarbocyanine dye with biomolecule bovine serum albumin. 1-11	0

169	Ligand-guided investigation of a series of formamidine-based thiuram disulfides as potential dual-inhibitors of COX-1 and COX-2.	0
168	Discovery of superinsecticide-resistant dengue mosquitoes in Asia: Threats of concomitant knockdown resistance mutations. <b>2022</b> , 8,	1
167	Unconventional interactions of the TRPV4 ion channel with beta-adrenergic receptor ligands. <b>2023</b> , 6, e202201704	0
166	Bioactive compounds from <i>Pandanous fascicularis</i> as potential therapeutic candidate to tackle hepatitis a inhibition: Docking and molecular dynamics simulation study. 1-17	2
165	Comprehensive Survey of Consensus Docking for High-Throughput Virtual Screening. <b>2023</b> , 28, 175	0
164	Decoding the binding interaction of steroidal pyridines with bovine serum albumin using spectroscopic and molecular docking techniques. <b>2022</b> , 109156	0
163	3D-QSAR, drug-likeness, ADMET prediction, and molecular docking studies in silico of novel 5-oxo-1-thioxo-4,5-dihydro-1H-thiazolo[3,4-a]quinazoline derivatives as MALT1 protease inhibitors for the treatment of B cell lymphoma.	1
162	Absolute binding free energies of the antiviral peptide ATN-161 with protein targets of SARS-CoV-2. 1-12	0
161	A new serotonin 2A receptor antagonist with potential benefits in non-alcoholic fatty liver disease. <b>2022</b> , 121315	0
160	Evaluating <i>Khaya senegalensis</i> for dipeptidyl peptidaseIV inhibition using in vitro analysis and molecular dynamic simulation of identified bioactive compounds.	0
159	A Biologically Active Chromone from <i>Bomarea setacea</i> (alstroemeriaceae): Leishmanicidal, Antioxidant and Multilevel Computational Studies. <b>2022</b> , 7,	0
158	In Vitro Study of Cytotoxic Mechanisms of Alkylphospholipids and Alkyltriazoles in Acute Lymphoblastic Leukemia Models. <b>2022</b> , 27, 8633	0
157	Destabilizers of the thymidylate synthase homodimer accelerate its proteasomal degradation and inhibit cancer growth. 11,	0
156	Synthesis, characterization and biological evaluation of novel 2-aminopridinium nicotinate invitro antifungal, insilico ADME and molecular docking studies. <b>2022</b> , 134794	0
155	Investigation of the Interaction between Aloe vera Anthraquinone Metabolites and c-Myc and C-Kit G-Quadruplex DNA Structures. <b>2022</b> , 23, 16018	0
154	Design, synthesis, biological evaluation and molecular docking of cyclic biguanidine compounds as cholinesterase inhibitors. 1-15	0
153	The Impact of Software Used and the Type of Target Protein on Molecular Docking Accuracy. <b>2022</b> , 27, 9041	0
152	Scanning aldoxime dehydratase sequence space and characterization of a new aldoxime dehydratase from <i>Fusarium vanettenii</i> . <b>2022</b> , 110187	1



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- 145 Synthesis, crystal structure, DFT, Hirshfeld surface analysis, energy framework, docking and molecular dynamic simulations of (E)-4-(4-methylbenzyl)-6-styrylpyridazin-3(2H)-one as anticancer agent. 1-20 ○
- 144 Pyridine derivatives complexes of Co (II) and Ni (II) 3-Bromobenzoates: Crystal Structure, in silico Anti-SARS-CoV-2 potential, Serum Albumin Binding Properties and Cytotoxicity. ○
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- 134 Neural Networks in the Design of Molecules with Affinity to Selected Protein Domains. **2023**, 24, 1762 ○

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- 114 Identification of thrombin inhibiting antithrombin-III like protein from Punica granatum using in silico approach and in vitro validation of thrombin inhibition activity in crude protein. 1-13 ○
- 113 In silico Identification of Triclosan Derivatives as Potential Inhibitors of Mutant Mycobacterium tuberculosis InhA. ○
- 112 Novel Benzo[4,5]imidazo[1,2-a]pyrimidine derivatives as selective Cyclooxygenase-2 Inhibitors: Design, synthesis, docking studies, and biological evaluation. ○
- 111 Interaction of copper potential metallodrugs with TMPRSS2: A comparative study of docking tools and its implications on COVID-19. 11, ○
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