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AutoDock4 and AutoDockTools4: Automated docking with selective receptor flexibility

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- 2182 A Catalytic Binding Site Together with a Distal Tyr in Myoglobin Affords Catalytic Efficiencies Similar to Natural Peroxidases.
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- 2156 Fast Dynamic Docking Guided by Adaptive Electrostatic Bias: The MD-Binding Approach.
- 2155 .
- 2154 .

2153 In Silico Fragment-Based Design Identifies Subfamily B1 Metallo--lactamase Inhibitors.

2152 Identification of Eukaryotic Translation Elongation Factor 1 1 Gamendazole-Binding Site for Binding of 3Hydroxy-4(1H)quinolinones as Novel Ligands with Anticancer Activity.

2151 Targeting DNA Repair in Tumor Cells via Inhibition of ERCC1XPF.

2150 Bcl-XL inhibits membrane permeabilization by competing with Bax. **2008**, 6, e147 232

2149 AutoDock Vina: improving the speed and accuracy of docking with a new scoring function, efficient optimization, and multithreading. *Journal of Computational Chemistry*, **2010**, 31, 455-61 3.5 11670

2148 Computational design tools for synthetic biology. **2009**, 20, 479-85 84

2147 N-acylpolyamine inhibitors of HDM2 and HDMX binding to p53. **2009**, 17, 7884-93 21

2146 Discovery of a novel protein kinase B inhibitor by structure-based virtual screening. **2009**, 19, 4634-8 16

2145 Activating mutations in TOR are in similar structures as oncogenic mutations in PI3KCalpha. **2009**, 4, 999-1015 30

2144 Chemoinformatics-applications in food chemistry. **2009**, 58, 33-56 15

2143 Virtual screening against alpha-cobratoxin. **2009**, 14, 1109-18 17

2142 Erybraedin C, a natural compound from the plant *Bituminaria bituminosa*, inhibits both the cleavage and religation activities of human topoisomerase I. **2010**, 425, 531-9 35

2141 Protein Flexibility in In Silico Screening. **2010**, 867-887

2140 Cooccupancy of the outer vestibule of voltage-gated sodium channels by micro-conotoxin KIIIA and saxitoxin or tetrodotoxin. **2010**, 104, 88-97 33

2139 Conformational flexibility and allosteric regulation of cathepsin K. **2010**, 429, 379-89 57

2138 Reactive-site mutants of N-TIMP-3 that selectively inhibit ADAMTS-4 and ADAMTS-5: biological and structural implications. **2010**, 431, 113-22 50

2137 Molecular dynamics simulations reveal fundamental role of water as factor determining affinity of binding of beta-blocker nebivolol to beta(2)-adrenergic receptor. **2010**, 114, 8374-86 30

2136 Multilayer coating of gold nanoparticles with drug-polymer coadsorbates. **2010**, 26, 16901-8 57

| | | |
|------|--|----|
| 2135 | Development of selective and reversible pyrazoline based MAO-A inhibitors: Synthesis, biological evaluation and docking studies. 2010 , 18, 1875-81 | 42 |
| 2134 | Novel estrone mimetics with high 17beta-HSD1 inhibitory activity. 2010 , 18, 3494-505 | 29 |
| 2133 | Binding of the general odorant binding protein of Bombyx mori BmorGOBP2 to the moth sex pheromone components. 2010 , 36, 1293-305 | 72 |
| 2132 | Structural insights into the function of P2X4: an ATP-gated cation channel of neuroendocrine cells. 2010 , 30, 1251-8 | 17 |
| 2131 | An efficient synthesis of a rationally designed 1,5 disubstituted imidazole AT(1) angiotensin II receptor antagonist: reorientation of imidazole pharmacophore groups in losartan reserves high receptor affinity and confirms docking studies. 2010 , 24, 749-58 | 15 |
| 2130 | NMR-based exploration of the acceptor binding site of human blood group B galactosyltransferase with molecular fragments. 2010 , 27, 349-58 | 17 |
| 2129 | A possible structural model of members of the CPF family of cuticular proteins implicating binding to components other than chitin. 2010 , 56, 1420-6 | 16 |
| 2128 | Structural underpinnings of nitrogen regulation by the prototypical nitrogen-responsive transcriptional factor NrpR. 2010 , 18, 1512-21 | 7 |
| 2127 | Community-driven computational biology with Debian Linux. 2010 , 11 Suppl 12, S5 | 37 |
| 2126 | Triplet-triplet energy transfer in the major intrinsic light-harvesting complex of <i>Amphidinium carterae</i> as revealed by ODMR and EPR spectroscopies. 2010 , 1797, 1759-67 | 27 |
| 2125 | Conformational flexibility of transmembrane helix VII of the human serotonin transporter impacts ion dependence and transport. 2010 , 80, 1418-26 | 5 |
| 2124 | Synthesis of N-hydroxycinnamides capped with a naturally occurring moiety as inhibitors of histone deacetylase. 2010 , 5, 598-607 | 33 |
| 2123 | In Pursuit of Fully Flexible Protein-Ligand Docking: Modeling the Bilateral Mechanism of Binding. 2010 , 29, 164-73 | 17 |
| 2122 | Role of Arg301 in substrate orientation and catalysis in subsite 2 of D-alanine:D-alanine (D-lactate) ligase from <i>Leuconostoc mesenteroides</i> : a molecular docking study. 2010 , 28, 728-34 | 5 |
| 2121 | Tetrapyrrole binding affinity of the murine and human p22HBP heme-binding proteins. 2010 , 29, 396-405 | 9 |
| 2120 | Computational approaches for protein function prediction: a combined strategy from multiple sequence alignment to molecular docking-based virtual screening. 2010 , 1804, 1695-712 | 68 |
| 2119 | Structure-based virtual screening of novel tubulin inhibitors and their characterization as anti-mitotic agents. 2010 , 18, 7092-100 | 33 |
| 2118 | Molecular docking studies of phlorotannins from <i>Eisenia bicyclis</i> with BACE1 inhibitory activity. 2010 , 20, 3211-5 | 86 |

| | | |
|------|--|-----|
| 2117 | Modification at the acidic domain of RXR agonists has little effect on permissive RXR-heterodimer activation. 2010 , 20, 5139-42 | 21 |
| 2116 | Triazolyl tryptoline derivatives as α -secretase inhibitors. 2010 , 20, 6572-6 | 20 |
| 2115 | A comparative study of flavonoid analogues on streptozotocin-nicotinamide induced diabetic rats: quercetin as a potential antidiabetic agent acting via 11beta-hydroxysteroid dehydrogenase type 1 inhibition. 2010 , 45, 2606-12 | 89 |
| 2114 | MOLA: a bootable, self-configuring system for virtual screening using AutoDock4/Vina on computer clusters. 2010 , 2, 10 | 18 |
| 2113 | The measured and calculated affinity of methyl- and methoxy-substituted benzoquinones for the Q(A) site of bacterial reaction centers. 2010 , 78, 2638-54 | 11 |
| 2112 | Molecular dynamics simulations of 2-amino-6-arylsulphonylbenzonitriles analogues as HIV inhibitors: interaction modes and binding free energies. 2010 , 76, 518-26 | 35 |
| 2111 | The inducer maltotriose binds in the central cavity of the tetratricopeptide-like sensor domain of MalT, a bacterial STAND transcription factor. 2010 , 77, 628-41 | 11 |
| 2110 | Pharmacological inhibition of gut-derived serotonin synthesis is a potential bone anabolic treatment for osteoporosis. 2010 , 16, 308-12 | 234 |
| 2109 | Visualization of macromolecular structures. 2010 , 7, S42-55 | 107 |
| 2108 | Estimating affinities of calcium ions to proteins. 2010 , 3, 1-6 | 3 |
| 2107 | Defining specific lipid binding sites for a peripheral membrane protein in situ using subtesla field-cycling NMR. 2010 , 285, 26916-26922 | 28 |
| 2106 | Structure-based modeling of the functional HIV-1 intasome and its inhibition. 2010 , 107, 15910-5 | 169 |
| 2105 | Opal web services for biomedical applications. 2010 , 38, W724-31 | 36 |
| 2104 | Structural and kinetic analysis of Schwanniomyces occidentalis invertase reveals a new oligomerization pattern and the role of its supplementary domain in substrate binding. 2010 , 285, 13930-41 | 60 |
| 2103 | Ligand-binding site prediction of proteins based on known fragment-fragment interactions. 2010 , 26, 1493-9 | 14 |
| 2102 | Nitric-oxide synthase forms N-NO-pterin and S-NO-cys: implications for activity, allostery, and regulation. 2010 , 285, 31581-9 | 29 |
| 2101 | A flavin-dependent monooxygenase from Mycobacterium tuberculosis involved in cholesterol catabolism. 2010 , 285, 22264-75 | 86 |
| 2100 | The role of oligomerization and cooperative regulation in protein function: the case of tryptophan synthase. 2010 , 6, e1000994 | 31 |

| | | |
|------|---|-----|
| 2099 | Modulated photophysics of an ESIPT probe 1-hydroxy-2-naphthaldehyde within motionally restricted environments of liposome membranes having varying surface charges. 2010 , 114, 12528-40 | 75 |
| 2098 | Insights into the mechanism of binding of arachidonic acid to mammalian 15-lipoxygenases. 2010 , 114, 7037-46 | 30 |
| 2097 | Mapping spatial relationships between residues in the ligand-binding domain of the 5-HT ₃ receptor using a molecular ruler. 2010 , 98, 1847-55 | 9 |
| 2096 | Cured of "stickiness", poly-L α -hairpin is promoted with LL-to-DD mutation as a protein and a hydrolase mimic. 2010 , 114, 16887-93 | 14 |
| 2095 | Rapid context-dependent ligand desolvation in molecular docking. 2010 , 50, 1561-73 | 213 |
| 2094 | Sterol binding and membrane lipid attachment to the Osh4 protein of yeast. 2010 , 114, 13562-73 | 4 |
| 2093 | Mechanism of selective halogenation by SyrB2: a computational study. 2010 , 132, 12887-98 | 67 |
| 2092 | The ubiquitin-proteasome system and assays to determine responses to inhibitors. 2010 , 5, 1221-36 | 5 |
| 2091 | Virtual Screening with AutoDock: Theory and Practice. 2010 , 5, 597-607 | 326 |
| 2090 | The crystal structure of the novobiocin biosynthetic enzyme NovP: the first representative structure for the TylF O-methyltransferase superfamily. 2010 , 395, 390-407 | 26 |
| 2089 | A dynamic model of HIV integrase inhibition and drug resistance. 2010 , 397, 600-15 | 52 |
| 2088 | Molecular framework of steroid/retinoid discrimination in 17 β -hydroxysteroid dehydrogenase type 1 and photoreceptor-associated retinol dehydrogenase. 2010 , 399, 255-67 | 16 |
| 2087 | Interactions of the melanocortin-4 receptor with the peptide agonist NDP-MSH. 2010 , 401, 433-50 | 19 |
| 2086 | Three new powerful oseltamivir derivatives for inhibiting the neuraminidase of influenza virus. 2010 , 401, 188-91 | 33 |
| 2085 | A structural and functional perspective into the mechanism of Ca ²⁺ -sensitizers that target the cardiac troponin complex. 2010 , 49, 1031-41 | 47 |
| 2084 | Genome scanning of Amazonian Plasmodium falciparum shows subtelomeric instability and clindamycin-resistant parasites. 2010 , 20, 1534-44 | 48 |
| 2083 | Understanding the DNA binding of novel non-symmetrical guanidinium/2-aminoimidazolinium derivatives. 2010 , 8, 5558-67 | 32 |
| 2082 | VSDocker: a tool for parallel high-throughput virtual screening using AutoDock on Windows-based computer clusters. 2010 , 26, 1374-5 | 38 |

| | | |
|------|--|-----|
| 2081 | Binding of the hemopressin peptide to the cannabinoid CB1 receptor: structural insights. 2010 , 49, 10449-57 | 20 |
| 2080 | Synthetic triterpenoids target the Arp2/3 complex and inhibit branched actin polymerization. 2010 , 285, 27944-57 | 36 |
| 2079 | Assignment of UV-vis spectrum of (3,3')-diindolylmethane, a <i>Leishmania donovani</i> topoisomerase IB inhibitor and a candidate DNA minor groove binder. 2010 , 114, 7121-6 | 6 |
| 2078 | LPYFDa neutralizes amyloid-beta-induced memory impairment and toxicity. 2010 , 19, 991-1005 | 25 |
| 2077 | A Case of Meta-scheduling for Biological Researches in Cloud Environment. 2010 , | 0 |
| 2076 | Design of small molecules that target metal-A{beta} species and regulate metal-induced A{beta} aggregation and neurotoxicity. 2010 , 107, 21990-5 | 225 |
| 2075 | Identification and characterization of the first small molecule inhibitor of MDMX. 2010 , 285, 10786-96 | 152 |
| 2074 | Tackling the challenges posed by target flexibility in drug design. 2010 , 5, 347-59 | 35 |
| 2073 | Solvents derived from glycerol modify classical regioselectivity in the enzymatic synthesis of disaccharides with Biolacta β -galactosidase. 2011 , 13, 2810 | 23 |
| 2072 | Molecular modeling of the interaction between heparan sulfate and cellular growth factors: bringing pieces together. 2011 , 21, 1181-93 | 39 |
| 2071 | The Arabidopsis cell cycle F-box protein SKP2A binds to auxin. 2010 , 22, 3891-904 | 106 |
| 2070 | Binding modes of noncompetitive GABA-channel blockers revisited using engineered affinity-labeling reactions combined with new docking studies. 2011 , 59, 2803-7 | 9 |
| 2069 | Theonellasterols and conicasterols from <i>Theonella swinhoei</i> . Novel marine natural ligands for human nuclear receptors. 2011 , 54, 3065-75 | 55 |
| 2068 | Structural fine-tuning of a multifunctional cytochrome P450 monooxygenase. 2011 , 133, 2292-302 | 57 |
| 2067 | Significant enhancement of docking sensitivity using implicit ligand sampling. 2011 , 51, 693-706 | 23 |
| 2066 | Interaction between the biotin carboxyl carrier domain and the biotin carboxylase domain in pyruvate carboxylase from <i>Rhizobium etli</i> . 2011 , 50, 9708-23 | 35 |
| 2065 | L-arginine binding to human inducible nitric oxide synthase: an antisymmetric funnel route toward isoform-specific inhibitors?. 2011 , 51, 1325-35 | 2 |
| 2064 | Predicting the binding mode of known NCp7 inhibitors to facilitate the design of novel modulators. 2011 , 51, 446-54 | 32 |

| | | |
|------|---|----|
| 2063 | Theoretical studies on the interactions and interferences of HIV-1 glycoprotein gp120 and its coreceptor CCR5. 2011 , 51, 359-69 | 10 |
| 2062 | Discovery of a Potent Retinoid X Receptor Antagonist Structurally Closely Related to RXR Agonist NET-31B. 2011 , 2, 896-900 | 10 |
| 2061 | In vivo trypanosomicidal activity of imidazole- or pyrazole-based benzo[g]phthalazine derivatives against acute and chronic phases of Chagas disease. 2011 , 54, 970-9 | 37 |
| 2060 | Naproxen interferes with the assembly of A β oligomers implicated in Alzheimer's disease. 2011 , 100, 2024-32 | 20 |
| 2059 | Computational insight into small molecule inhibition of cyclophilins. 2011 , 51, 475-82 | 11 |
| 2058 | Computational modeling of the catalytic mechanism of human placental alkaline phosphatase (PLAP). 2011 , 51, 2538-48 | 11 |
| 2057 | In vitro polymerization of microtubules with a fullerene derivative. 2011 , 5, 6306-14 | 45 |
| 2056 | Virtual screening identification of nonfolate compounds, including a CNS drug, as antiparasitic agents inhibiting pteridine reductase. 2011 , 54, 211-21 | 52 |
| 2055 | Allosteric inhibition of the hepatitis C virus NS5B polymerase: in silico strategies for drug discovery and development. 2011 , 3, 1027-55 | 35 |
| 2054 | Paramagnetic relaxation assisted docking of a small indole compound in the HIV-1 gp41 hydrophobic pocket. 2011 , 6, 267-74 | 32 |
| 2053 | Structural analysis of mammalian cytochrome P450 2B4 covalently bound to the mechanism-based inactivator tert-butylphenylacetylene: insight into partial enzymatic activity. 2011 , 50, 4903-11 | 35 |
| 2052 | Role of Mg ²⁺ ions in protein kinase phosphorylation: insights from molecular dynamics simulations of ATP-kinase complexes. 2011 , 37, 1143-1150 | 12 |
| 2051 | Bioinformatics in crosslinking chemistry of collagen with selective cross linkers. 2011 , 4, 399 | 12 |
| 2050 | Using free energy of binding calculations to improve the accuracy of virtual screening predictions. 2011 , 51, 1648-55 | 20 |
| 2049 | A spectral deciphering of the binding interaction of an intramolecular charge transfer fluorescence probe with a cationic protein: thermodynamic analysis of the binding phenomenon combined with blind docking study. 2011 , 10, 980-91 | 88 |
| 2048 | Insight into the enzyme-inhibitor interactions of the first experimentally determined human aromatase. 2011 , 28, 759-71 | 4 |
| 2047 | Protein Flexibility in Structure-Based Virtual Screening: From Models to Algorithms. 2011 , 223-244 | 4 |
| 2046 | Accurate prediction of the bound form of the Akt pleckstrin homology domain using normal mode analysis to explore structural flexibility. 2011 , 51, 2352-60 | 8 |

| | | |
|------|--|----|
| 2045 | Development of bifunctional stilbene derivatives for targeting and modulating metal-amyloid- β species. 2011 , 50, 10724-34 | 71 |
| 2044 | NMR studies reveal an unexpected binding site for a redox inhibitor of AP endonuclease 1. 2011 , 50, 10540-9 | 18 |
| 2043 | Insight into the metabolism of 1,1,1-trichloro-2,2-bis(4-chlorophenyl)ethane (DDT) by biphenyl dioxygenases. 2011 , 516, 35-44 | 15 |
| 2042 | Synthetic biosensing systems. 2011 , 43, 310-9 | 18 |
| 2041 | Human aryl-hydrocarbon receptor and its interaction with dioxin and physiological ligands investigated by molecular modelling and docking simulations. 2011 , 413, 176-81 | 12 |
| 2040 | Dopamine release and molecular modeling study of some coumarin derivatives. 2011 , 59, 906-12 | 11 |
| 2039 | Juvenile hormone synthesis: "esterify then epoxidize" or "epoxidize then esterify"? Insights from the structural characterization of juvenile hormone acid methyltransferase. 2011 , 41, 228-35 | 35 |
| 2038 | Preparation and characterization of a thermostable and biodegradable biopolymers using natural cross-linker. 2011 , 48, 276-85 | 47 |
| 2037 | Structural analysis of chorismate synthase from Plasmodium falciparum: a novel target for antimalaria drug discovery. 2011 , 49, 767-77 | 11 |
| 2036 | A new Caucasian case of neonatal intrahepatic cholestasis caused by citrin deficiency (NICCD): a clinical, molecular, and functional study. 2011 , 104, 501-6 | 22 |
| 2035 | Progress in structure-based drug design against influenza A virus. 2011 , 6, 619-31 | 12 |
| 2034 | Application of NMR and molecular docking in structure-based drug discovery. 2012 , 326, 1-34 | 32 |
| 2033 | Towards new ligands of nuclear receptors. Discovery of malitasterol A, an unique bis-secosterol from marine sponge Theonella swinhoei. 2011 , 9, 4856-62 | 32 |
| 2032 | Identification of the Schistosoma mansoni molecular target for the antimalarial drug artemether. 2011 , 51, 3005-16 | 14 |
| 2031 | . 2011 , | 26 |
| 2030 | Complementary Use of NMR to X-Ray Crystallography for the Analysis of Protein Morphological Change in Solution. 2011 , | |
| 2029 | Molecular-docking study of capsular regulatory protein in Streptococcus pneumoniae portends the novel approach to its treatment. 2011 , 131 | |
| 2028 | Computational perspectives into plasmepsins structure-function relationship: implications to inhibitors design. 2011 , 2011, 657483 | 6 |

| | | |
|------|--|-----|
| 2027 | Schiff bases of indoline-2,3-dione: potential novel inhibitors of Mycobacterium tuberculosis (Mtb) DNA gyrase. 2011 , 16, 7864-79 | 33 |
| 2026 | Synthetic Biology & Bioinformatics Prospects in the Cancer Arena. 2011 , | |
| 2025 | Comparative analysis of species-specific ligand recognition in Toll-like receptor 8 signaling: a hypothesis. 2011 , 6, e25118 | 41 |
| 2024 | The bile acid receptor GPBAR-1 (TGR5) modulates integrity of intestinal barrier and immune response to experimental colitis. 2011 , 6, e25637 | 221 |
| 2023 | Elucidation of the ATP7B N-domain Mg ²⁺ -ATP coordination site and its allosteric regulation. 2011 , 6, e26245 | 1 |
| 2022 | Molecular recognition in the case of flexible targets. 2011 , 17, 1663-71 | 39 |
| 2021 | Bioinformatics. 2011 , 23-40 | |
| 2020 | Probing the S1 specificity pocket of the aminopeptidases that generate antigenic peptides. 2011 , 435, 411-20 | 74 |
| 2019 | Survey of public domain software for docking simulations and virtual screening. 2011 , 5, 497-505 | 30 |
| 2018 | Crystal structure of the N-terminal domain of linker L(R) and the assembly of cyanobacterial phycobilisome rods. 2011 , 82, 698-705 | 13 |
| 2017 | Predicting the substrate specificity of a glycosyltransferase implicated in the production of phenolic volatiles in tomato fruit. 2011 , 278, 390-400 | 18 |
| 2016 | The prion protein binds thiamine. 2011 , 278, 4002-14 | 25 |
| 2015 | Docking, synthesis, and in vitro evaluation of antimetabolic estrone analogs. 2011 , 77, 173-81 | 44 |
| 2014 | Drug-like leads for steric discrimination between substrate and inhibitors of human acetylcholinesterase. 2011 , 78, 495-504 | 20 |
| 2013 | Non-bisphosphonate inhibitors of isoprenoid biosynthesis identified via computer-aided drug design. 2011 , 78, 323-32 | 34 |
| 2012 | Status of GPCR modeling and docking as reflected by community-wide GPCR Dock 2010 assessment. 2011 , 19, 1108-26 | 243 |
| 2011 | Structural basis for μ opioid receptor binding and activation. 2011 , 19, 1683-90 | 23 |
| 2010 | Improved synthesis of disaccharides with Escherichia coli β -galactosidase using bio-solvents derived from glycerol. 2011 , 67, 7708-7712 | 20 |

| | | |
|------|--|-----|
| 2009 | Structural bases for understanding the stereoselectivity in ketone reductions with ADH from <i>Thermus thermophilus</i> : A quantitative model. 2011 , 70, 23-31 | 14 |
| 2008 | Multiscale methods for nanochemistry and biophysics in solution. 2011 , 164, 101-112 | 18 |
| 2007 | Role of the omega loop in specificity determination in subsite 2 of the D-alanine:D-alanine (D-lactate) ligase from <i>Leuconostoc mesenteroides</i> : a molecular docking study. 2011 , 30, 31-7 | 7 |
| 2006 | Molecular modeling and molecular dynamics simulation studies on pyrrolopyrimidine-based β -helix mimetic as dual inhibitors of MDM2 and MDMX. 2011 , 30, 167-78 | 16 |
| 2005 | Peptide binding prediction for the human class II MHC allele HLA-DP2: a molecular docking approach. 2011 , 11, 32 | 38 |
| 2004 | 6-Carboxyfluorescein and structurally similar molecules inhibit DNA binding and repair by O ⁶ -alkylguanine DNA alkyltransferase. 2011 , 10, 1193-202 | 11 |
| 2003 | Synthesis and evaluation of aliphatic-chain hydroxamates capped with osthole derivatives as histone deacetylase inhibitors. 2011 , 46, 4042-9 | 28 |
| 2002 | α (2)-Adrenergic activity of 6-methoxykaempferol-3-O-glucoside on rat uterus: in vitro and in silico studies. 2011 , 667, 348-54 | 6 |
| 2001 | Trypanocidal properties, structure-activity relationship and computational studies of quinoxaline 1,4-di-N-oxide derivatives. 2011 , 127, 745-51 | 29 |
| 2000 | Selective targeting of breast cancer cells through ROS-mediated mechanisms potentiates the lethality of paclitaxel by a novel diterpene, gelomulide K. 2011 , 51, 641-57 | 37 |
| 1999 | Structure-based design of dipeptide derivatives for the human neutral endopeptidase. 2011 , 19, 5935-47 | 5 |
| 1998 | Thiosemicarbazones derived from 1-indanones as new anti- <i>Trypanosoma cruzi</i> agents. 2011 , 19, 6818-26 | 43 |
| 1997 | New cholesterol esterase inhibitors based on rhodanine and thiazolidinedione scaffolds. 2011 , 19, 7453-63 | 36 |
| 1996 | Synthesis and bio-evaluation of human macrophage migration inhibitory factor inhibitor to develop anti-inflammatory agent. 2011 , 19, 7365-73 | 24 |
| 1995 | Targeting the proangiogenic VEGF-VEGFR protein-protein interface with drug-like compounds by in silico and in vitro screening. 2011 , 18, 1631-9 | 35 |
| 1994 | Activation and regulation of purinergic P2X receptor channels. 2011 , 63, 641-83 | 384 |
| 1993 | Luminescence resonance energy transfer in the cytoplasm of live <i>Escherichia coli</i> cells. 2011 , 50, 6789-96 | |
| 1992 | Efficient incorporation of protein flexibility and dynamics into molecular docking simulations. 2011 , 50, 6157-69 | 78 |

| | | |
|------|---|-----|
| 1991 | Monoamine neurotransmitters as substrates for novel tick sulfotransferases, homology modeling, molecular docking, and enzyme kinetics. 2011 , 6, 176-84 | 10 |
| 1990 | Protein-ligand docking. 2011 , 16, 2289-306 | 13 |
| 1989 | Outstanding challenges in protein-ligand docking and structure-based virtual screening. 2011 , 1, 229-259 | 67 |
| 1988 | Receptor flexibility in small-molecule docking calculations. 2011 , 1, 298-314 | 38 |
| 1987 | Computer-aided drug design platform using PyMOL. 2011 , 25, 13-9 | 278 |
| 1986 | Molecular motions in drug design: the coming age of the metadynamics method. 2011 , 25, 395-402 | 34 |
| 1985 | Structure-guided fragment-based in silico drug design of dengue protease inhibitors. 2011 , 25, 263-74 | 46 |
| 1984 | Substrate binding to mammalian 15-lipoxygenase. 2011 , 25, 825-35 | 15 |
| 1983 | A computational analysis of the binding mode of closantel as inhibitor of the <i>Onchocerca volvulus</i> chitinase: insights on macrofilaricidal drug design. 2011 , 25, 1107-19 | 16 |
| 1982 | Bonding interactions and stability assessment of biopolymer material prepared using type III collagen of avian intestine and anionic polysaccharides. 2011 , 22, 1419-29 | 8 |
| 1981 | Inhibition of snowshoe hare succinate dehydrogenase activity as a mechanism of deterrence for papyriferic acid in birch. 2011 , 37, 1285-93 | 19 |
| 1980 | Mixed-type inhibition of tyrosinase from <i>Agaricus bisporus</i> by terephthalic acid: computational simulations and kinetics. 2011 , 30, 273-80 | 29 |
| 1979 | Natural products as DNA methyltransferase inhibitors: a computer-aided discovery approach. 2011 , 15, 293-304 | 117 |
| 1978 | Molecular modeling of lanthionine synthetase component C-like protein 2: a potential target for the discovery of novel type 2 diabetes prophylactics and therapeutics. 2011 , 17, 543-53 | 29 |
| 1977 | Integrative computational protocol for the discovery of inhibitors of the <i>Helicobacter pylori</i> nickel response regulator (NikR). 2011 , 17, 3075-84 | 3 |
| 1976 | Modeling of ligand binding to G protein coupled receptors: cannabinoid CB1, CB2 and adrenergic β 2 AR. 2011 , 17, 2353-66 | 27 |
| 1975 | Prediction of a new surface binding pocket and evaluation of inhibitors against huntingtin interacting protein 14: an insight using docking studies. 2011 , 17, 3047-56 | 13 |
| 1974 | Thiabendazole inhibits ubiquinone reduction activity of mitochondrial respiratory complex II via a water molecule mediated binding feature. 2011 , 2, 531-42 | 29 |

| | | |
|------|---|--------|
| 1973 | Multilevel Parallelization of AutoDock 4.2. 2011 , 3, 12 | 92 |
| 1972 | Structure of the mouse galectin-4 N-terminal carbohydrate-recognition domain reveals the mechanism of oligosaccharide recognition. 2011 , 67, 204-11 | 17 |
| 1971 | Structural basis for drug and substrate specificity exhibited by FIV encoding a chimeric FIV/HIV protease. 2011 , 67, 540-8 | 9 |
| 1970 | Understanding the binding mode and function of BMS-488043 against HIV-1 viral entry. 2011 , 79, 1810-9 | 10 |
| 1969 | Conformational changes induced by ATP-hydrolysis in an ABC transporter: a molecular dynamics study of the Sav1866 exporter. 2011 , 79, 1977-90 | 53 |
| 1968 | Toward prediction of functional protein pockets using blind docking and pocket search algorithms. 2011 , 20, 880-93 | 50 |
| 1967 | vsLabAn implementation for virtual high-throughput screening using AutoDock and VMD. 2011 , 111, 1208-1212 | 25 |
| 1966 | Challenges and advances in computational docking: 2009 in review. 2011 , 24, 149-64 | 210 |
| 1965 | Hybrid Steered Molecular Dynamics-Docking: An Efficient Solution to the Problem of Ranking Inhibitor Affinities Against a Flexible Drug Target. 2011 , 30, 459-471 | 26 |
| 1964 | Blind Dockings of Benzothiazoles to Multiple Receptor Conformations of Triosephosphate Isomerase from <i>Trypanosoma cruzi</i> and Human. 2011 , 30, 986-95 | 7 |
| 1963 | Kinetic Resolution of β -Bromoamides: Experimental and Theoretical Investigation of Highly Enantioselective Reactions Catalyzed by Haloalkane Dehalogenases. 2011 , 353, 931-944 | 33 |
| 1962 | Species extrapolation for the 21st century. 2011 , 30, 52-63 | 54 |
| 1961 | Bicyclic substituted hydroxyphenylmethanone type inhibitors of 17 β -hydroxysteroid dehydrogenase Type 1 (17 β -HSD1): the role of the bicyclic moiety. 2011 , 6, 476-87 | 15 |
| 1960 | Design and synthesis of novel cyclooxygenase-1 inhibitors as analgesics: 5-amino-2-ethoxy-N-(substituted-phenyl)benzamides. 2011 , 6, 550-60 | 8 |
| 1959 | An amphiphilic conjugate approach toward the design and synthesis of betulinic acid-polyphenol conjugates as inhibitors of the HIV-1 gp41 fusion core formation. 2011 , 6, 1654-64 | 10 |
| 1958 | The inverse type II β -turn on D-Trp-Phe, a pharmacophoric motif for MOR agonists. 2011 , 6, 1640-53 | 22 |
| 1957 | Study on the effects of intermolecular interactions on firefly multicolor bioluminescence. 2011 , 12, 3002-8 | 33 |
| 1956 | Predicting the accuracy of protein-ligand docking on homology models. <i>Journal of Computational Chemistry</i> , 2011 , 32, 81-98 | 3-5 57 |

| | | | |
|------|--|-----|-----|
| 1955 | Role of bridging water molecules in GSK3 β -inhibitor complexes: insights from QM/MM, MD, and molecular docking studies. <i>Journal of Computational Chemistry</i> , 2011 , 32, 1907-18 | 3.5 | 31 |
| 1954 | Fast docking using the CHARMM force field with EADock DSS. <i>Journal of Computational Chemistry</i> , 2011 , 32, 2149-59 | 3.5 | 269 |
| 1953 | Theoretical modulation of the color of light emitted by firefly oxyluciferin. <i>Journal of Computational Chemistry</i> , 2011 , 32, 2654-63 | 3.5 | 28 |
| 1952 | DockoMatic: automated peptide analog creation for high throughput virtual screening. <i>Journal of Computational Chemistry</i> , 2011 , 32, 2936-41 | 3.5 | 13 |
| 1951 | Efficient inclusion of receptor flexibility in grid-based protein-ligand docking. <i>Journal of Computational Chemistry</i> , 2011 , 32, 3433-9 | 3.5 | 25 |
| 1950 | Redox-Based Probes for Protein Tyrosine Phosphatases. 2011 , 123, 4515-4519 | | 2 |
| 1949 | Redox-based probes for protein tyrosine phosphatases. 2011 , 50, 4423-7 | | 45 |
| 1948 | In-vitro antiproliferative activity of benzopyranone derivatives in comparison with standard chemotherapeutic drugs. 2011 , 344, 102-10 | | 7 |
| 1947 | The binding mode of cladocoran A to the human group IIA phospholipase A(2). 2011 , 12, 2686-91 | | 5 |
| 1946 | Chemical screens against a reconstituted multiprotein complex: myricetin blocks DnaJ regulation of DnaK through an allosteric mechanism. 2011 , 18, 210-21 | | 80 |
| 1945 | Vasorelaxant effect of flavonoids through calmodulin inhibition: Ex vivo, in vitro, and in silico approaches. 2011 , 19, 542-6 | | 21 |
| 1944 | Replacement of the hydrophobic part of 9-cis-retinoic acid with cyclic terpenoid moiety results in RXR-selective agonistic activity. 2011 , 19, 2939-49 | | 8 |
| 1943 | N-substituted homopiperazine barbiturates as gelatinase inhibitors. 2011 , 19, 4985-99 | | 43 |
| 1942 | Identification of a novel small molecule targeting UQCRB of mitochondrial complex III and its anti-angiogenic activity. 2011 , 21, 1052-6 | | 21 |
| 1941 | Dual function inhibitors of relevance to chronic obstructive pulmonary disease. 2011 , 21, 3177-80 | | 4 |
| 1940 | Virtual screening-driven identification of human carbonic anhydrase inhibitors incorporating an original, new pharmacophore. 2011 , 21, 2515-20 | | 7 |
| 1939 | Identification of chemicals to inhibit the kinase activity of leucine-rich repeat kinase 2 (LRRK2), a Parkinson's disease-associated protein. 2011 , 21, 2953-7 | | 21 |
| 1938 | 3-substitued indoles: one-pot synthesis and evaluation of anticancer and Src kinase inhibitory activities. 2011 , 21, 3511-4 | | 47 |

| | | |
|------|--|------|
| 1937 | Self-organizing molecular field analysis on human α -secretase nonpeptide inhibitors: 5, 5-disubstituted aminohydantoins. 2011 , 46, 58-64 | 15 |
| 1936 | Discovery, synthesis, and investigation of the antitumor activity of novel piperazinympyrimidine derivatives. 2011 , 46, 2043-57 | 33 |
| 1935 | In search of patterns over physicochemical properties and pharmacological activities for a set of [MCl ₂ (thiosemicarbazone)] complexes (M=Pt/Pd): support for multiple mechanisms of antichagasic action excluding DNA-bonding in vivo?. 2011 , 46, 2639-51 | 19 |
| 1934 | Insight into analysis of interactions of GW9508 to wild-type and H86F and H137F GPR40: a combined QM/MM study and pharmacophore modeling. 2011 , 29, 818-25 | 7 |
| 1933 | Prediction of the PPAR α agonism of fibrates by combined MM-docking approaches. 2011 , 29, 865-75 | 4 |
| 1932 | A new arginase enzymatic reactor: development and application for the research of plant-derived inhibitors. 2011 , 55, 48-53 | 13 |
| 1931 | Interaction of a ruthenium(II)halcone complex with double stranded DNA: Spectroscopic, molecular docking and nuclease properties. 2011 , 220, 145-152 | 53 |
| 1930 | Exploring possibility of promiscuity of amyloid inhibitor: Studies on effect of selected compounds on folding and amyloid formation of proteins. 2011 , 46, 1179-1185 | 14 |
| 1929 | Accounting for induced-fit effects in docking: what is possible and what is not?. 2011 , 11, 179-91 | 45 |
| 1928 | Molecular docking: a powerful approach for structure-based drug discovery. 2011 , 7, 146-57 | 1083 |
| 1927 | The significance of chirality in drug design and development. 2011 , 11, 760-70 | 284 |
| 1926 | A hierarchical framework for cross-domain MapReduce execution. 2011 , | 34 |
| 1925 | Epigallocatechin-3-gallate potently inhibits the in vitro activity of hydroxy-3-methyl-glutaryl-CoA reductase. 2011 , 52, 897-907 | 62 |
| 1924 | The statin class of HMG-CoA reductase inhibitors demonstrate differential activation of the nuclear receptors PXR, CAR and FXR, as well as their downstream target genes. 2011 , 41, 519-29 | 49 |
| 1923 | The role of long-range intermolecular interactions in discovery of new drugs. 2011 , 6, 1263-70 | 17 |
| 1922 | Structural insight into human variegate porphyria disease. 2011 , 25, 653-64 | 43 |
| 1921 | Atomic structure of salutaridine reductase from the opium poppy (<i>Papaver somniferum</i>). 2011 , 286, 6532-41 | 14 |
| 1920 | Structural aspects of binding of linked digalactosides to human galectin-1. 2011 , 21, 1627-41 | 40 |

| | | |
|------|--|-----|
| 1919 | Bimolecular Complementation to Visualize Filovirus VP40-Host Complexes in Live Mammalian Cells: Toward the Identification of Budding Inhibitors. 2011 , 2011, | 23 |
| 1918 | Inhibitory effect of phthalic Acid on tyrosinase: the mixed-type inhibition and docking simulations. 2011 , 2011, 294724 | 28 |
| 1917 | BEAR, a novel virtual screening methodology for drug discovery. 2011 , 16, 129-33 | 28 |
| 1916 | ATP independent type IB topoisomerase of <i>Leishmania donovani</i> is stimulated by ATP: an insight into the functional mechanism. 2011 , 39, 3295-309 | 3 |
| 1915 | Experimental and molecular simulation investigation of interaction between acyclovir and bovine serum albumin. 2011 , 37, 1239-1247 | 4 |
| 1914 | NADH oxidase activity of indoleamine 2,3-dioxygenase. 2011 , 286, 29273-29283 | 16 |
| 1913 | Correlating computational docking predictions with Raman spectroscopy for β -lactoglobulin-porphyrin complexes. 2011 , | |
| 1912 | CScore: a simple yet effective scoring function for protein-ligand binding affinity prediction using modified CMAC learning architecture. 2011 , 9 Suppl 1, 1-14 | 30 |
| 1911 | MD-2 as the target of nonlipid chalcone in the inhibition of endotoxin LPS-induced TLR4 activity. 2011 , 203, 1012-20 | 41 |
| 1910 | Selective phthalate activation of naturally occurring human constitutive androstane receptor splice variants and the pregnane X receptor. 2011 , 120, 381-91 | 72 |
| 1909 | Quantifying intramolecular binding in multivalent interactions: a structure-based synergistic study on Grb2-Sos1 complex. 2011 , 7, e1002192 | 15 |
| 1908 | Identification and characterization of novel small-molecule inhibitors against hepatitis delta virus replication by using docking strategies. 2011 , 11, 803-9 | 6 |
| 1907 | Preliminary structure-activity relationship on theonellasterol, a new chemotype of FXR antagonist, from the marine sponge <i>Theonella swinhoei</i> . 2012 , 10, 2448-66 | 14 |
| 1906 | Virtual screening of specific insulin-like growth factor 1 receptor (IGF1R) inhibitors from the National Cancer Institute (NCI) molecular database. 2012 , 13, 17185-209 | 4 |
| 1905 | BiP-mediated closing of the Sec61 channel limits Ca ²⁺ leakage from the ER. 2012 , 31, 3282-96 | 111 |
| 1904 | Structural basis for the substrate recognition and catalysis of peptidyl-tRNA hydrolase. 2012 , 40, 10521-31 | 23 |
| 1903 | A structural basis for the biosynthesis of the major chlorogenic acids found in coffee. 2012 , 160, 249-60 | 88 |
| 1902 | β Cyclodextrin dimer complexes of dopamine and levodopa derivatives to assess drug delivery to the central nervous system: ADME and molecular docking studies. 2012 , 7, 3211-9 | 32 |

| | | |
|------|---|-----|
| 1901 | Rotating wall vessel exposure alters protein secretion and global gene expression in <i>Staphylococcus aureus</i> . 2012 , 11, 71-81 | 7 |
| 1900 | Structural and functional characterization of the two phosphoinositide binding sites of PROPPINs, a β -propeller protein family. 2012 , 109, E2042-9 | 117 |
| 1899 | Rotavirus VP8*: phylogeny, host range, and interaction with histo-blood group antigens. 2012 , 86, 9899-910 | 128 |
| 1898 | An investigation of the catalytic activity of CYP2A13*4 with coumarin and polymorphisms of CYP2A13 in a Chinese Han population. 2012 , 40, 847-51 | 2 |
| 1897 | Pyrazine functionalized Ag(I) and Au(I)-NHC complexes are potential antibacterial agents. 2012 , 19, 4184-93 | 48 |
| 1896 | State-independent intracellular access of quaternary ammonium blockers to the pore of TREK-1. 2012 , 6, 473-8 | 34 |
| 1895 | Identification of SRC as a potent drug target for asthma, using an integrative approach of protein interactome analysis and in silico drug discovery. 2012 , 16, 513-26 | 12 |
| 1894 | High-throughput Molecular Docking Now in Reach for a Wider Biochemical Community. 2012 , | 2 |
| 1893 | Necrostatin-1 analogues: critical issues on the specificity, activity and in vivo use in experimental disease models. 2012 , 3, e437 | 290 |
| 1892 | Novel anti-inflammatory activity of epoxyazadiradione against macrophage migration inhibitory factor: inhibition of tautomerase and proinflammatory activities of macrophage migration inhibitory factor. 2012 , 287, 24844-61 | 63 |
| 1891 | Mechanistic insights into regulated cargo binding by ACAP1 protein. 2012 , 287, 28675-85 | 21 |
| 1890 | Rapid and accurate prediction and scoring of water molecules in protein binding sites. 2012 , 7, e32036 | 124 |
| 1889 | A salt-bridge controlled by ligand binding modulates the hydrolysis reaction in a GH5 endoglucanase. 2012 , 25, 223-33 | 8 |
| 1888 | Computational analyses of the catalytic and heparin-binding sites and their interactions with glycosaminoglycans in glycoside hydrolase family 79 endo- β -D-glucuronidase (heparanase). 2012 , 22, 35-55 | 40 |
| 1887 | Structural understanding of the glutathione-dependent reduction mechanism of glutathionyl-hydroquinone reductases. 2012 , 287, 35838-48 | 11 |
| 1886 | F18, a novel small-molecule nonnucleoside reverse transcriptase inhibitor, inhibits HIV-1 replication using distinct binding motifs as demonstrated by resistance selection and docking analysis. 2012 , 56, 341-51 | 20 |
| 1885 | Remarkable ability of <i>Pandoraea pnomenus</i> B356 biphenyl dioxygenase to metabolize simple flavonoids. 2012 , 78, 3560-70 | 31 |
| 1884 | The use of the R language for medicinal chemistry applications. 2012 , 12, 1957-64 | 20 |

| | | |
|------|---|----|
| 1883 | Identification of novel potent inhibitors against Bcl-xL anti-apoptotic protein using docking studies. 2012 , 19, 1302-17 | 5 |
| 1882 | WITHDRAWN: Baicalein protects against Staphylococcus aureus pneumonia in a mouse model via inhibition of the self-assembly of Hemolysin heptamers. 2012 , | |
| 1881 | Non-selective inhibition of cyclooxygenase enzymes by aminoacetylenic isoindoline 1,3-diones. 2012 , 11, 369-74 | 8 |
| 1880 | Targeting the epidermal growth factor receptor: exploring the potential of novel inhibitor N-(3-ethynylphenyl)-6, 7-bis (2-methoxyethoxy) quinolin- 4-amine using docking and molecular dynamics simulation. 2012 , 19, 955-68 | 9 |
| 1879 | Identification of HIV inhibitors guided by free energy perturbation calculations. 2012 , 18, 1199-216 | 19 |
| 1878 | Binding site and affinity prediction of general anesthetics to protein targets using docking. 2012 , 114, 947-55 | 23 |
| 1877 | Discovery, synthesis, and evaluation of small-molecule signal transducer and activator of transcription 3 inhibitors. 2012 , 60, 1574-80 | 6 |
| 1876 | Prediction of heparin binding sites in bone morphogenetic proteins (BMPs). 2012 , 1824, 1374-81 | 71 |
| 1875 | Identification of BfmR, a response regulator involved in biofilm development, as a target for a 2-Aminoimidazole-based antibiofilm agent. 2012 , 51, 9776-8 | 51 |
| 1874 | In silico implementation of synthetic gene networks. 2012 , 813, 3-21 | 7 |
| 1873 | Synthesis of new 7-oxycoumarin derivatives as potent and selective monoamine oxidase A inhibitors. 2012 , 55, 10424-36 | 30 |
| 1872 | Chemically modified tetracyclines as inhibitors of MMP-2 matrix metalloproteinase: a molecular and structural study. 2012 , 116, 13644-54 | 26 |
| 1871 | GalaxyDock: protein-ligand docking with flexible protein side-chains. 2012 , 52, 3225-32 | 44 |
| 1870 | Utilizing experimental data for reducing ensemble size in flexible-protein docking. 2012 , 52, 187-98 | 34 |
| 1869 | Cross-docking study on InhA inhibitors: a combination of Autodock Vina and PM6-DH2 simulations to retrieve bio-active conformations. 2012 , 10, 6341-9 | 47 |
| 1868 | Identification of pim kinases as novel targets for PJ34 with confounding effects in PARP biology. 2012 , 7, 1962-7 | 59 |
| 1867 | 3D-RISM-Dock: A New Fragment-Based Drug Design Protocol. 2012 , 8, 3356-72 | 34 |
| 1866 | Comparative analysis of Rac1 binding efficiency with different classes of ligands: morpholines, flavonoids and imidazoles. 2012 , 11, 181-7 | 3 |

| | | |
|------|---|----|
| 1865 | Virtual screening and biological evaluation of inhibitors targeting the XPA-ERCC1 interaction. 2012 , 7, e51329 | 48 |
| 1864 | On setting up and assessing docking simulations for virtual screening. 2012 , 928, 1-16 | 2 |
| 1863 | Dual-function triazole-pyridine derivatives as inhibitors of metal-induced amyloid- β aggregation. 2012 , 4, 910-20 | 53 |
| 1862 | Fluorescent stilbazolium dyes as probes of the norepinephrine transporter: structural insights into substrate binding. 2012 , 10, 8710-9 | 17 |
| 1861 | Modification in the side chain of solomonsterol A: discovery of cholestan disulfate as a potent pregnane-X-receptor agonist. 2012 , 10, 6350-62 | 15 |
| 1860 | In silico deconstruction of ATP-competitive inhibitors of glycogen synthase kinase-3 β . 2012 , 52, 3233-44 | 6 |
| 1859 | A naked-eye on-off-on molecular "light switch" based on a reversible "conformational switch" of G-quadruplex DNA. 2012 , 51, 12591-3 | 58 |
| 1858 | Identification and characterization of small molecule inhibitors of a plant homeodomain finger. 2012 , 51, 8293-306 | 70 |
| 1857 | Camptothecin-7-yl-methanthiole: semisynthesis and biological evaluation. 2012 , 7, 2134-43 | 18 |
| 1856 | Interaction between odorants and proteins involved in the perception of smell: the case of odorant-binding proteins probed by molecular modelling and biophysical data. 2012 , 27, 445-453 | 14 |
| 1855 | Identifying Activity Cliff Generators of PPAR Ligands Using SAS Maps. 2012 , 31, 837-46 | 26 |
| 1854 | Homology modeling and docking study of recent SHV type β -lactamses with traditional and novel inhibitors: an in silico approach to combat problem of multiple drug resistance in various infections. 2012 , 21, 2229-2237 | 1 |
| 1853 | Modeling, docking, simulation, and inhibitory activity of the benzimidazole analogue against β -tubulin protein from <i>Brugia malayi</i> for treating lymphatic filariasis. 2012 , 21, 2415-2427 | 19 |
| 1852 | Synthesis and anti-inflammatory evaluation of some new 3,6-disubstituted-1,2,4-triazolo-[3,4-b]-1,3,4-thiadiazoles bearing pyrazole moiety. 2012 , 21, 3272-3280 | 7 |
| 1851 | In silico and in vitro pharmacological investigations of a natural alkaloid. 2012 , 21, 4100-4107 | |
| 1850 | Virtual screening of phenylsulfonamido-3-morpholinopropan-2-yl dihydrogen phosphate derivatives as novel inhibitors of MurC/MurF ligases from <i>Mycobacterium leprae</i> . 2012 , 21, 4341-4351 | 4 |
| 1849 | Structural basis for modification of flavonol and naphthol glucoconjugates by <i>Nicotiana tabacum</i> malonyltransferase (NtMaT1). 2012 , 236, 781-93 | 17 |
| 1848 | Homology modeling of the human 5-HT _{1A} , 5-HT _{2A} , D ₁ , and D ₂ receptors: model refinement with molecular dynamics simulations and docking evaluation. 2012 , 18, 3639-55 | 24 |

| | | |
|------|--|-----|
| 1847 | Response surface methodology in docking study of small molecule BACE-1 inhibitors. 2012 , 18, 4567-76 | 11 |
| 1846 | Comprehensive model of wild-type and mutant HIV-1 reverse transcriptases. 2012 , 26, 907-19 | 12 |
| 1845 | MALT1 small molecule inhibitors specifically suppress ABC-DLBCL in vitro and in vivo. 2012 , 22, 812-24 | 182 |
| 1844 | Assessment of amsacrine binding with DNA using UV-visible, circular dichroism and Raman spectroscopic techniques. 2012 , 114, 38-43 | 54 |
| 1843 | In silico screening of quadruplex-binding ligands. 2012 , 57, 106-14 | 25 |
| 1842 | Ab initio molecular simulations on specific interactions between amyloid beta and monosaccharides. 2012 , 547, 89-96 | 2 |
| 1841 | Ligand selection from the analysis of protein conformational substates: new leads targeting the N-terminal domain of Hsp90. 2012 , 2, 4268 | 7 |
| 1840 | An improved LGA for protein-ligand docking prediction. 2012 , | 0 |
| 1839 | Suberic Acid Acts as a Dissolving Agent as Well as a Crosslinker for Natural Polymers (Carbohydrate and Protein): A Detailed Discussion on the Chemistry Behind the Interaction. 2012 , 49, 619-629 | 5 |
| 1838 | Differential interactions of a biological photosensitizer with liposome membranes having varying surface charges. 2012 , 11, 661-73 | 12 |
| 1837 | Efficient and selective enzymatic synthesis of N-acetyl-lactosamine in ionic liquid: a rational explanation. 2012 , 2, 6306 | 29 |
| 1836 | GPU Accelerated Molecular Docking with Parallel Genetic Algorithm. 2012 , | 1 |
| 1835 | Recent trends and applications in 3D virtual screening. 2012 , 15, 749-69 | 12 |
| 1834 | Peptide inhibitors of viral assembly: a novel route to broad-spectrum antivirals. 2012 , 52, 770-6 | 14 |
| 1833 | Combination of resonance Raman spectroscopy and docking simulations to study the nonspecific binding of a free-base porphyrin to a globular protein. 2012 , 116, 11032-40 | 10 |
| 1832 | Identification of binding specificity-determining features in protein families. 2012 , 55, 1926-39 | 4 |
| 1831 | Selective monocationic inhibitors of neuronal nitric oxide synthase. Binding mode insights from molecular dynamics simulations. 2012 , 134, 11559-72 | 20 |
| 1830 | Imaging Carbon Nanotube Interaction with Nucleobases in Water Using the Statistical Mechanical Theory of Molecular Liquids. 2012 , 116, 15087-15092 | 12 |

| | | |
|------|---|-----|
| 1829 | A bactericidal guanidinomethyl biaryl that alters the dynamics of bacterial FtsZ polymerization. 2012 , 55, 10160-76 | 45 |
| 1828 | Insights into the role of magnesium triad in myo-inositol monophosphatase: metal mechanism, substrate binding, and lithium therapy. 2012 , 52, 2398-409 | 21 |
| 1827 | A force field with discrete displaceable waters and desolvation entropy for hydrated ligand docking. 2012 , 55, 623-38 | 150 |
| 1826 | Conicasterol E, a small heterodimer partner sparing farnesoid X receptor modulator endowed with a pregnane X receptor agonistic activity, from the marine sponge Theonella swinhoei. 2012 , 55, 84-93 | 36 |
| 1825 | Structural insight into the mechanism of DNA-binding attenuation of the Neisserial adhesin repressor NadR by the small natural ligand 4-hydroxyphenylacetic acid. 2012 , 51, 6738-52 | 21 |
| 1824 | Potentiation of the Kv1 Family K(+) channel by cortisone analogues. 2012 , 7, 1641-6 | 5 |
| 1823 | New determinants in the catalytic mechanism of nucleoside hydrolases from the structures of two isozymes from Sulfolobus solfataricus. 2012 , 51, 4590-9 | 3 |
| 1822 | 4-Methylenesterols from Theonella swinhoei sponge are natural pregnane-X-receptor agonists and farnesoid-X-receptor antagonists that modulate innate immunity. 2012 , 77, 484-95 | 36 |
| 1821 | Isatin-3-N4-benzilthiosemicarbazone, a non-toxic thiosemicarbazone derivative, protects and reactivates rat and human cholinesterases inhibited by methamidophos in vitro and in silico. 2012 , 26, 1030-9 | 7 |
| 1820 | Synthesis, biological activity and docking study of some new isatin Schiff base derivatives. 2012 , 21, 3730-3740 | 35 |
| 1819 | The role of carboxymethyl substituents in the interaction of tetracationic porphyrins with DNA. 2012 , 41, 723-32 | 11 |
| 1818 | Anion activation site of insulin-degrading enzyme. 2012 , 287, 48-57 | 20 |
| 1817 | Mitochondrial F(0) F(1) -ATP synthase is a molecular target of 3-iodothyronamine, an endogenous metabolite of thyroid hormone. 2012 , 166, 2331-47 | 24 |
| 1816 | Cytotoxicity and slow release of the anti-cancer drug doxorubicin from ZIF-8. 2012 , 2, 9437 | 193 |
| 1815 | In silico mutagenesis and docking study of Ralstonia solanacearum RSL lectin: performance of docking software to predict saccharide binding. 2012 , 52, 1250-61 | 31 |
| 1814 | A strategy based on protein-protein interface motifs may help in identifying drug off-targets. 2012 , 52, 2273-86 | 22 |
| 1813 | Mechanism of taq DNA polymerase inhibition by fullerene derivatives: insight from computer simulations. 2012 , 116, 10676-83 | 22 |
| 1812 | The evolution of cefotaximase activity in the TEM β -lactamase. 2012 , 415, 205-20 | 11 |

| | | |
|------|---|--------|
| 1811 | Investigation on the interaction behavior between bisphenol A and pepsin by spectral and docking studies. 2012 , 1021, 34-39 | 42 |
| 1810 | Inhibition of forebrain μ opioid receptor signaling by low concentrations of rimonabant does not require cannabinoid receptors and directly involves μ opioid receptors. 2012 , 61, 378-88 | 15 |
| 1809 | Peptide screening to knockdown Bcl-2's anti-apoptotic activity: implications in cancer treatment. 2012 , 50, 796-814 | 17 |
| 1808 | Investigations on the binding of human hemoglobin with orange I and orange II. 2012 , 113, 14-21 | 17 |
| 1807 | Potent low toxicity inhibition of human melanogenesis by novel indole-containing octapeptides. 2012 , 1820, 1481-9 | 22 |
| 1806 | Investigation of a potential mechanism for the inhibition of SmTGR by Auranofin and its implications for Plasmodium falciparum inhibition. 2012 , 417, 576-81 | 24 |
| 1805 | Analysis of Fasciola cathepsin L5 by S2 subsite substitutions and determination of the P1-P4 specificity reveals an unusual preference. 2012 , 94, 1119-27 | 12 |
| 1804 | Design, synthesis and preliminary bioactivity studies of 1,3,4-thiadiazole hydroxamic acid derivatives as novel histone deacetylase inhibitors. 2012 , 20, 3865-72 | 39 |
| 1803 | Comparative theoretical study of the binding of luciferyl-adenylate and dehydroluciferyl-adenylate to firefly luciferase. 2012 , 543, 137-141 | 6 |
| 1802 | Pd-catalyzed direct C-H bond functionalization of spirocyclic π ligands: generation of a pharmacophore model and analysis of the reverse binding mode by docking into a 3D homology model of the μ receptor. 2012 , 55, 8047-65 | 46 |
| 1801 | The inactivation mechanism of human group IIA phospholipase A(2) by Scalaradial. 2012 , 13, 2259-64 | 5 |
| 1800 | Synthesis and biological evaluation of CTP synthetase inhibitors as potential agents for the treatment of African trypanosomiasis. 2012 , 7, 1623-34 | 26 |
| 1799 | Modeling, synthesis and biological evaluation of potential retinoid X receptor-selective agonists: novel halogenated analogues of 4-[1-(3,5,5,8,8-pentamethyl-5,6,7,8-tetrahydro-2-naphthyl)ethynyl]benzoic acid (bexarotene). 2012 , 7, 1551-66 | 8 |
| 1798 | SIMONA 1.0: an efficient and versatile framework for stochastic simulations of molecular and nanoscale systems. <i>Journal of Computational Chemistry</i> , 2012 , 33, 2602-13 | 3-5 21 |
| 1797 | Phylogenomics of the benzoxazinoid biosynthetic pathway of Poaceae: gene duplications and origin of the Bx cluster. 2012 , 12, 64 | 60 |
| 1796 | Peptide binding to HLA-DP proteins at pH 5.0 and pH 7.0: a quantitative molecular docking study. 2012 , 12, 20 | 15 |
| 1795 | Crystal structure of HutZ, a heme storage protein from <i>Vibrio cholerae</i> : A structural mismatch observed in the region of high sequence conservation. 2012 , 12, 23 | 14 |
| 1794 | Prospects of Modulating Protein-Protein Interactions. 2012 , 295-329 | 3 |

| | | |
|------|--|------|
| 1793 | Another brick in the wall. Validation of the β receptor 3D model by computer-assisted design, synthesis, and activity of new β ligands. 2012 , 9, 3107-26 | 41 |
| 1792 | Template-based protein structure modeling using the RaptorX web server. 2012 , 7, 1511-22 | 1115 |
| 1791 | Probing ligand-binding modes and binding mechanisms of benzoxazole-based amide inhibitors with soluble epoxide hydrolase by molecular docking and molecular dynamics simulation. 2012 , 116, 10219-33 | 19 |
| 1790 | Spectroscopic and molecular docking studies on chlorambucil interaction with DNA. 2012 , 51, 406-11 | 52 |
| 1789 | Molecular docking of bisphenol A and its nitrated and chlorinated metabolites onto human estrogen-related receptor-gamma. 2012 , 426, 215-20 | 24 |
| 1788 | Molecular characterization of a thermostable L-fucose isomerase from <i>Dictyoglomus turgidum</i> that isomerizes L-fucose and D-arabinose. 2012 , 94, 1926-34 | 13 |
| 1787 | Discovery of structurally-diverse inhibitor scaffolds by high-throughput screening of a fragment library with dimethylarginine dimethylaminohydrolase. 2012 , 20, 5550-8 | 12 |
| 1786 | Synthesis and SAR studies of 3-allyl-4-prenyloxylaniline amides as potent 15-lipoxygenase inhibitors. 2012 , 20, 5518-26 | 17 |
| 1785 | Benzimidazole inhibitors of the protein kinase CHK2: clarification of the binding mode by flexible side chain docking and protein-ligand crystallography. 2012 , 20, 6630-9 | 8 |
| 1784 | Discrete molecular dynamics distinguishes natively-like binding poses from decoys in difficult targets. 2012 , 102, 144-51 | 28 |
| 1783 | Design and synthesis of a second series of triazole-based compounds as potent dual mPGES-1 and 5-lipoxygenase inhibitors. 2012 , 54, 311-23 | 37 |
| 1782 | COX inhibitors Indomethacin and Sulindac derivatives as antiproliferative agents: synthesis, biological evaluation, and mechanism investigation. 2012 , 56, 17-29 | 25 |
| 1781 | Synthesis and SAR studies of mono O-prenylated coumarins as potent 15-lipoxygenase inhibitors. 2012 , 57, 134-42 | 44 |
| 1780 | Application of a post-docking procedure based on MM-PBSA and MM-GBSA on single and multiple protein conformations. 2012 , 58, 431-40 | 57 |
| 1779 | 3D models of human ER α and ER β complexed with 5-androsten-3 β ,17 β -diol. 2012 , 77, 1192-7 | 13 |
| 1778 | Synthesis of highly oxygenated decalins from sugar allyltins: an access to sulfur and phosphorus derivatives. 2012 , 23, 1501-1511 | 6 |
| 1777 | Targeting essential cell wall lipase Rv3802c for potential therapeutics against tuberculosis. 2012 , 38, 235-42 | 6 |
| 1776 | Inhibitory activities and possible anticancer targets of Ru(II)-based complexes using computational docking method. 2012 , 38, 60-9 | 11 |

| | | |
|------|--|-----|
| 1775 | Molecular modeling of the lipase-catalyzed hydrolysis of acetoxymethyl(i-propoxy)phenylphosphine oxide and its P-borane analogue. 2012 , 38, 290-7 | 11 |
| 1774 | A molecular model of the enantioselective liquid chromatographic separation of (R,S)-ifosfamide and its N-dechloroethylated metabolites on a teicoplanin aglycon chiral stationary phase. 2012 , 1269, 218-25 | 19 |
| 1773 | Structure-based and multiple potential three-dimensional quantitative structure-activity relationship (SB-MP-3D-QSAR) for inhibitor design. 2012 , 52, 996-1004 | 7 |
| 1772 | Molecular mechanism of direct proflavine-DNA intercalation: evidence for drug-induced minimum base-stacking penalty pathway. 2012 , 116, 12208-12 | 31 |
| 1771 | Promising antidiabetic potential of fucoxanthin isolated from the edible brown algae Eisenia bicyclis and Undaria pinnatifida. 2012 , 78, 1321-1329 | 46 |
| 1770 | Biotinylated Rh(III) complexes in engineered streptavidin for accelerated asymmetric C-H activation. 2012 , 338, 500-3 | 623 |
| 1769 | idock: A multithreaded virtual screening tool for flexible ligand docking. 2012 , | 44 |
| 1768 | Affinity binding-guided fluorescent nanobiosensor for acetylcholinesterase inhibitors via distance modulation between the fluorophore and metallic nanoparticle. 2012 , 84, 2830-6 | 36 |
| 1767 | TD-DFT/molecular mechanics study of the Photinus pyralis bioluminescence system. 2012 , 116, 2008-13 | 27 |
| 1766 | The alkenyl migration mechanism catalyzed by extradiol dioxygenases: a hybrid DFT study. 2012 , 17, 881-90 | 11 |
| 1765 | Plakilactones from the marine sponge Plakinastrella mamillaris. Discovery of a new class of marine ligands of peroxisome proliferator-activated receptor α . 2012 , 55, 8303-17 | 38 |
| 1764 | Protein-Ligand Interactions: Computational Docking. 2012 , | 4 |
| 1763 | A bacterial glycosidase enables mannose-6-phosphate modification and improved cellular uptake of yeast-produced recombinant human lysosomal enzymes. 2012 , 30, 1225-31 | 79 |
| 1762 | Discovery and synthesis of namalide reveals a new anabaenopeptin scaffold and peptidase inhibitor. 2012 , 55, 735-42 | 24 |
| 1761 | Computational Drug Discovery and Design. 2012 , | 19 |
| 1760 | Cuticular Proteins. 2012 , 134-166 | 23 |
| 1759 | Hierarchical MapReduce Programming Model and Scheduling Algorithms. 2012 , | 19 |
| 1758 | Auto dock-based incremental docking protocol to improve docking of large ligands. 2012 , | |

| | | |
|------|---|----|
| 1757 | In silico-aided design of a glycan ligand of sialoadhesin for in vivo targeting of macrophages. 2012 , 134, 15696-9 | 48 |
| 1756 | Discovery that theonellasterol a marine sponge sterol is a highly selective FXR antagonist that protects against liver injury in cholestasis. 2012 , 7, e30443 | 47 |
| 1755 | Natural iminosugar (+)-lentiginosine inhibits ATPase and chaperone activity of hsp90. 2012 , 7, e43316 | 35 |
| 1754 | 3D models of MBP, a biologically active metabolite of bisphenol A, in human estrogen receptor β and estrogen receptor α 2012 , 7, e46078 | 39 |
| 1753 | Understanding the basis of drug resistance of the mutants of β -tubulin dimer via molecular dynamics simulations. 2012 , 7, e42351 | 34 |
| 1752 | Binding modes of peptidomimetics designed to inhibit STAT3. 2012 , 7, e51603 | 20 |
| 1751 | Activation of hydrogen peroxide to peroxytetradecanoic acid is responsible for potent inhibition of protein tyrosine phosphatase CD45. 2012 , 7, e52495 | 7 |
| 1750 | The binding mode of second-generation sulfonamide inhibitors of MurD: clues for rational design of potent MurD inhibitors. 2012 , 7, e52817 | 10 |
| 1749 | Substituted derivatives of indole acetic acid as aldose reductase inhibitors with antioxidant activity: structure-activity relationship. 2011 , 30, 342-9 | 7 |
| 1748 | Interactomic and pharmacological insights on human sirt-1. 2012 , 3, 40 | 21 |
| 1747 | In Silico and In Vitro Comparison of HIV-1 Subtypes B and CRF02_AG Integrase Susceptibility to Integrase Strand Transfer Inhibitors. 2012 , 2012, 548657 | 5 |
| 1746 | Molecular docking studies of banana flower flavonoids as insulin receptor tyrosine kinase activators as a cure for diabetes mellitus. 2012 , 8, 216-20 | 23 |
| 1745 | Antifungal activity of homoaconitate and homoisocitrate analogs. 2012 , 17, 14022-36 | 7 |
| 1744 | In Silico Engineering of Proteins That Recognize Small Molecules. 2012 , | |
| 1743 | Comparative molecular docking analysis of essential oil constituents as elastase inhibitors. 2012 , 8, 457-60 | 14 |
| 1742 | Discovery of potential cyclooxygenase inhibitors using in silico docking studies. 2012 , 7, | 5 |
| 1741 | The use of glycoinformatics in glycochemistry. 2012 , 8, 915-29 | 20 |
| 1740 | Ab initio fragment molecular orbital calculations on specific interactions between aryl hydrocarbon receptor and dioxin. 2012 , 112, 289-299 | 2 |

| | | |
|------|--|----|
| 1739 | BSP-SLIM: a blind low-resolution ligand-protein docking approach using predicted protein structures. 2012 , 80, 93-110 | 65 |
| 1738 | Identifying continuous pores in protein structures with PROPORES by computational repositioning of gating residues. 2012 , 80, 421-32 | 9 |
| 1737 | In Silico study of carcinogenic o-Quinone metabolites derived from polycyclic aromatic hydrocarbons (PAHs). 2012 , 25, 720-728 | 1 |
| 1736 | Metal organic frameworks for drug delivery and environmental remediation: A molecular docking approach. 2012 , 112, 3346-3355 | 38 |
| 1735 | Protein flexibility in docking and surface mapping. 2012 , 45, 301-43 | 90 |
| 1734 | Bioinformatics and variability in drug response: a protein structural perspective. 2012 , 9, 1409-37 | 57 |
| 1733 | On the use of molecular dynamics receptor conformations for virtual screening. 2012 , 819, 93-103 | 16 |
| 1732 | FN-Linked glycopeptides: conformational analysis and bioactivity as lectin ligands. 2012 , 10, 5916-23 | 9 |
| 1731 | Chapter 7:Docking and Virtual Screening. 2012 , 171-194 | |
| 1730 | Chapter 13:Computational Strategies and Challenges for Targeting Protein-Protein Interactions with Small Molecules. 2012 , 319-359 | 2 |
| 1729 | Selective flexibility of side-chain residues improves VEGFR-2 docking score using AutoDock Vina. 2012 , 79, 530-4 | 24 |
| 1728 | X-ray structure of Salmonella typhimurium uridine phosphorylase complexed with 5-fluorouracil and molecular modelling of the complex of 5-fluorouracil with uridine phosphorylase from Vibrio cholerae. 2012 , 68, 968-74 | 6 |
| 1727 | CCR5 antagonist TD-0680 uses a novel mechanism for enhanced potency against HIV-1 entry, cell-mediated infection, and a resistant variant. 2012 , 287, 16499-509 | 14 |
| 1726 | Molecular docking studies of disubstituted diaryl diselenides as mammalian α -aminolevulinic acid dehydratase enzyme inhibitors. 2012 , 75, 1012-22 | 14 |
| 1725 | Determining the Role of the Aromatic Ring of N-Arylmethyl ent-conduramine F-1 in their Interactions with β -Glucosidases by Saturation Transfer Difference NMR Spectroscopy Experiments. 2012 , 1, 13-6 | 3 |
| 1724 | 3-D QSAutogrid/R: an alternative procedure to build 3-D QSAR models. Methodologies and applications. 2012 , 52, 1674-85 | 30 |
| 1723 | N-Octanoyldimethylglycine Trifluoroethyl Ester, an Acyl Donor Leading to Highly Enantioselective Protease-Catalysed Kinetic Resolution of Amines. 2012 , 354, 1759-1764 | 5 |
| 1722 | Adipic acid interaction enhances the mechanical and thermal stability of natural polymers. 2012 , 125, E490-E500 | 13 |

| | | |
|------|---|-----|
| 1721 | Functional analysis and molecular docking identify two active short-chain prenyltransferases in the green peach aphid, <i>Myzus persicae</i> . 2012 , 81, 63-76 | 10 |
| 1720 | Asymmetric synthesis of pochonin E and F, revision of their proposed structure, and their conversion to potent Hsp90 inhibitors. 2012 , 18, 8978-86 | 20 |
| 1719 | Comparative study of flavins binding with human serum albumin: a fluorometric, thermodynamic, and molecular dynamics approach. 2012 , 13, 2142-53 | 28 |
| 1718 | Combined QSAR studies of inhibitor properties of O-phosphorylated oximes toward serine esterases involved in neurotoxicity, drug metabolism and Alzheimer's disease. 2012 , 23, 627-47 | 34 |
| 1717 | Application of binding free energy calculations to prediction of binding modes and affinities of MDM2 and MDMX inhibitors. 2012 , 52, 1821-32 | 27 |
| 1716 | Validation of ITD mutations in FLT3 as a therapeutic target in human acute myeloid leukaemia. 2012 , 485, 260-3 | 525 |
| 1715 | Preparation and characterization of malonic acid cross-linked chitosan and collagen 3D scaffolds: an approach on non-covalent interactions. 2012 , 23, 1309-21 | 24 |
| 1714 | In silico and in vitro analyses identified three amino acid residues critical to the catalysis of two aphid farnesyl diphosphate synthase. 2012 , 31, 417-24 | 8 |
| 1713 | Effect of the electrostatic potential on the internalization mechanism of cell penetrating peptides derived from TIRAP. 2012 , 17, 485-499 | 3 |
| 1712 | New insights into the mechanism of the Schiff base formation catalyzed by type I dehydroquinase from <i>S. enterica</i> . 2012 , 131, 1 | 5 |
| 1711 | Hydroxybutyrate prevents protein aggregation in the halotolerant bacterium <i>Pseudomonas</i> sp. CT13 under abiotic stress. 2012 , 16, 455-62 | 34 |
| 1710 | Molecular dynamics simulations of the Bcl-2 protein to predict the structure of its unordered flexible loop domain. 2012 , 18, 1885-906 | 30 |
| 1709 | Transient pockets on XIAP-BIR2: toward the characterization of putative binding sites of small-molecule XIAP inhibitors. 2012 , 18, 2031-42 | 10 |
| 1708 | Mapping multiple potential ATP binding sites on the matrix side of the bovine ADP/ATP carrier by the combined use of MD simulation and docking. 2012 , 18, 2377-86 | 6 |
| 1707 | Homology modeling and docking analyses of <i>M. leprae</i> Mur ligases reveals the common binding residues for structure based drug designing to eradicate leprosy. 2012 , 18, 2659-72 | 10 |
| 1706 | Molecular dynamics simulation studies of betulinic acid with human serum albumin. 2012 , 18, 2589-97 | 41 |
| 1705 | Kinetics and molecular docking studies of kaempferol and its prenylated derivatives as aldose reductase inhibitors. 2012 , 197, 110-8 | 15 |
| 1704 | Rational design of small molecule inhibitors targeting the Rac GTPase-p67(phox) signaling axis in inflammation. 2012 , 19, 228-42 | 46 |

| | | |
|------|---|----|
| 1703 | Crystal structure of periplasmic catecholate-siderophore binding protein VctP from <i>Vibrio cholerae</i> at 1.7 Å resolution. 2012 , 586, 1240-4 | 14 |
| 1702 | Synthesis of novel benzofurocoumarin analogues and their anti-proliferative effect on human cancer cell lines. 2012 , 47, 370-6 | 20 |
| 1701 | Design, synthesis and biological evaluation of novel (E)- α -benzylsulfonyl chalcone derivatives as potential BRAF inhibitors. 2012 , 50, 288-95 | 27 |
| 1700 | Analysis of cause of failure of new targeting peptide in PEGylated liposome: molecular modeling as rational design tool for nanomedicine. 2012 , 46, 121-30 | 51 |
| 1699 | The exploration of thienothiazines as selective butyrylcholinesterase inhibitors. 2012 , 47, 190-205 | 30 |
| 1698 | Double-edged sword effect of biochanin to inhibit nuclear factor kappaB: suppression of serine/threonine and tyrosine kinases. 2012 , 83, 1383-92 | 24 |
| 1697 | Adamantane-substituted guanylhydrazones: novel inhibitors of butyrylcholinesterase. 2012 , 41-42, 28-34 | 13 |
| 1696 | Cell-based and in-silico studies on the high intrinsic activity of two boron-containing salbutamol derivatives at the human β_2 adrenoceptor. 2012 , 20, 933-41 | 14 |
| 1695 | Design, synthesis, biological evaluation, and comparative Cox1 and Cox2 docking of p-substituted benzylidenamino phenyl esters of ibuprofenic and mefenamic acids. 2012 , 20, 1259-70 | 35 |
| 1694 | Molecular modeling study of cyclic pentapeptide CXCR4 antagonists: new insight into CXCR4-FC131 interactions. 2012 , 22, 2146-50 | 33 |
| 1693 | A new series of N5 derivatives of the 1,1,5-trimethyl furo[3,4-c]pyridine-3,4-dione (cerpegin) selectively inhibits the post-acid activity of mammalian 20S proteasomes. 2012 , 22, 3822-7 | 12 |
| 1692 | Effect of different oximes on rat and human cholinesterases inhibited by methamidophos: a comparative in vitro and in silico study. 2012 , 111, 362-70 | 8 |
| 1691 | Revealing interaction mode between HIV-1 protease and mannitol analog inhibitor. 2012 , 79, 916-25 | |
| 1690 | Inhibition of bacterial virulence: drug-like molecules targeting the <i>Salmonella enterica</i> PhoP response regulator. 2012 , 79, 1007-17 | 39 |
| 1689 | Potential selective inhibitors against Rv0183 of <i>Mycobacterium tuberculosis</i> targeting host lipid metabolism. 2012 , 79, 1056-62 | 10 |
| 1688 | Molecular modeling of <i>T. rangeli</i> , <i>T. brucei gambiense</i> , and <i>T. evansi</i> sialidases in complex with the DANA inhibitor. 2012 , 80, 114-20 | 10 |
| 1687 | Interaction between phillygenin and human serum albumin based on spectroscopic and molecular docking. 2012 , 85, 120-6 | 19 |
| 1686 | Studies on the interaction between chromium(VI) and human serum albumin: spectroscopic approach. 2012 , 86, 381-6 | 11 |

| | | |
|------|---|----|
| 1685 | Reassignment of sense codons: Designing and docking of proline analogs for Escherichia coli prolyl-tRNA synthetase to expand the genetic code. 2012 , 78, 57-64 | 5 |
| 1684 | Development of metal-chelating inhibitors for the Class II fructose 1,6-bisphosphate (FBP) aldolase. 2012 , 112, 49-58 | 23 |
| 1683 | 1-Hydroxy-2-naphthaldehyde: A prospective excited-state intramolecular proton transfer (ESIPT) probe with multi-faceted applications. 2012 , 132, 2194-2208 | 43 |
| 1682 | Binding of TNT to amplifying fluorescent polymers: an ab initio and molecular dynamics study. 2012 , 33, 12-8 | 9 |
| 1681 | Molecular modeling and docking characterization of Dectin-1 (PAMP) receptor of Bubalus bubalis. 2012 , 92, 7-12 | 13 |
| 1680 | Enantioselective drug-protein interaction between mexiletine and plasma protein. 2012 , 64, 792-801 | 8 |
| 1679 | Computer-aided modeling of activity and selectivity of quinazolinones as noncompetitive NMDA receptor antagonists. 2012 , 443, 118-22 | 3 |
| 1678 | Directed evolution strategies for enantiocomplementary haloalkane dehalogenases: from chemical waste to enantiopure building blocks. 2012 , 13, 137-48 | 53 |
| 1677 | Inactivation of glucosamine-6-phosphate synthase by N3-oxoacyl derivatives of L-2,3-diaminopropanoic acid. 2012 , 13, 85-96 | 2 |
| 1676 | Active site analysis of cis-epoxysuccinate hydrolase from Nocardia tartaricans using homology modeling and site-directed mutagenesis. 2012 , 93, 2377-86 | 9 |
| 1675 | The interaction of blood proteins with brucine. 2012 , 39, 4937-47 | 16 |
| 1674 | Study of the interaction between 8-azaguanine and bovine serum albumin using optical spectroscopy and molecular modeling methods. 2012 , 18, 493-500 | 7 |
| 1673 | Analysis of surface cavity in serpin family reveals potential binding sites for chemical chaperone to reduce polymerization. 2012 , 18, 1143-51 | 2 |
| 1672 | Insight into the binding interactions of CYP450 aromatase inhibitors with their target enzyme: a combined molecular docking and molecular dynamics study. 2012 , 18, 1153-66 | 16 |
| 1671 | Oxidative inhibition of Hsp90 disrupts the super-chaperone complex and attenuates pancreatic adenocarcinoma in vitro and in vivo. 2013 , 132, 695-706 | 54 |
| 1670 | Fragment informatics and computational fragment-based drug design: an overview and update. 2013 , 33, 554-98 | 36 |
| 1669 | Emodin regulates apoptotic pathway in human liver cancer cells. 2013 , 27, 251-7 | 33 |
| 1668 | Epigallocatechin-3-gallate and penta-O-galloyl- β -D-glucose inhibit protein phosphatase-1. 2013 , 280, 612-26 | 25 |

| | | |
|------|---|-----|
| 1667 | Pyrazolo[3,4-d]pyrimidines as inhibitor of anti-coagulation and inflammation activities of phospholipase A 2 : insight from molecular docking studies. 2013 , 39, 419-38 | 27 |
| 1666 | Biophysical characterization and crystal structure of the Feline Immunodeficiency Virus p15 matrix protein. 2013 , 10, 64 | 8 |
| 1665 | Molecular insight into the inhibition mechanism of cyrtominetin to Hemolysin by molecular dynamics simulation. 2013 , 62, 320-8 | 33 |
| 1664 | Molecular dynamics simulation and molecular docking studies of Angiotensin converting enzyme with inhibitor lisinopril and amyloid Beta Peptide. 2013 , 32, 356-64 | 46 |
| 1663 | Exploring the dual role of di-carboxylic acids in the preparation of collagen based biomaterial. 2013 , 20, 647-661 | 3 |
| 1662 | Mechanism of the Humicola insolens Cel7B E197S mutant catalyzed flavonoid glycosides synthesis: a QM/MM metadynamics simulation study. 2013 , 132, 1 | 6 |
| 1661 | Binding of a natural anthocyanin inhibitor to influenza neuraminidase by mass spectrometry. 2013 , 405, 6563-72 | 34 |
| 1660 | Systems approaches and polypharmacology for drug discovery from herbal medicines: an example using licorice. 2013 , 146, 773-93 | 180 |
| 1659 | Quinazoline-based multi-tyrosine kinase inhibitors: synthesis, modeling, antitumor and antiangiogenic properties. 2013 , 67, 373-83 | 49 |
| 1658 | Antihyperglycemic and sub-chronic antidiabetic actions of morolic and moronic acids, in vitro and in silico inhibition of 11β-HSD 1. 2013 , 20, 571-6 | 17 |
| 1657 | Mimotope selection of blood group A antigen from a phage display 15-mer peptide library. 2013 , 29, 39-42 | 3 |
| 1656 | Evolution driven structural changes in CENP-E motor domain. 2013 , 5, 102-11 | 4 |
| 1655 | Assay platform for clinically relevant metallo-β-lactamases. 2013 , 56, 6945-53 | 80 |
| 1654 | Small-molecule modulation of Wnt signaling via modulating the Axin-LRP5/6 interaction. 2013 , 9, 579-85 | 58 |
| 1653 | Mutually exclusive binding of APPL(PH) to BAR domain and Reptin regulates β-catenin dependent transcriptional events. 2013 , 47, 48-55 | 1 |
| 1652 | Enterococcal and streptococcal resistance to PC190723 and related compounds: molecular insights from a FtsZ mutational analysis. 2013 , 95, 1880-7 | 16 |
| 1651 | Prospects in Bioscience: Addressing the Issues. 2013 , | 3 |
| 1650 | Inhibition of human alcohol and aldehyde dehydrogenases by cimetidine and assessment of its effects on ethanol metabolism. 2013 , 202, 275-82 | 15 |

| | | |
|------|---|-----|
| 1649 | Endocrine disrupting activities of the flavonoid nutraceuticals luteolin and quercetin. 2013 , 4, 293-300 | 41 |
| 1648 | Screening for enhanced triacetic acid lactone production by recombinant <i>Escherichia coli</i> expressing a designed triacetic acid lactone reporter. 2013 , 135, 10099-103 | 152 |
| 1647 | Asymmetric synthesis and molecular docking study of enantiomerically pure pyrrolidine derivatives with potential antithrombin activity. 2013 , 24, 838-843 | 13 |
| 1646 | Study of the enantioselective interaction of diclofop and human serum albumin by spectroscopic and molecular modeling approaches in vitro. 2013 , 25, 719-25 | 9 |
| 1645 | Mitochondrial glutamate carriers from <i>Drosophila melanogaster</i> : biochemical, evolutionary and modeling studies. 2013 , 1827, 1245-55 | 24 |
| 1644 | Application of the docking program SOL for CSAR benchmark. 2013 , 53, 1946-56 | 30 |
| 1643 | Insight into the binding model of new antagonists of kappa receptor using docking and molecular dynamics simulation. 2013 , 19, 3087-94 | 5 |
| 1642 | Potent anticancer activity of cystine-based dipeptides and their interaction with serum albumins. 2013 , 7, 91 | 10 |
| 1641 | CXI-benzo-84 reversibly binds to tubulin at colchicine site and induces apoptosis in cancer cells. 2013 , 86, 378-91 | 38 |
| 1640 | Pyridones as NNRTIs against HIV-1 mutants: 3D-QSAR and protein informatics. 2013 , 27, 637-54 | 12 |
| 1639 | On homology modeling of the M μ muscarinic acetylcholine receptor subtype. 2013 , 27, 525-38 | 16 |
| 1638 | Integrated analysis of cytochrome P450 gene superfamily in the red flour beetle, <i>Tribolium castaneum</i> . 2013 , 14, 174 | 71 |
| 1637 | Mapping of heparin/heparan sulfate binding sites on α 5 β 1 integrin by molecular docking. 2013 , 26, 76-85 | 28 |
| 1636 | Microbial Decolorization of an Azo Dye Reactive Black 5 Using White-Rot Fungus <i>Pleurotus eryngii</i> F032. 2013 , 224, 1 | 55 |
| 1635 | Potential role of cardiac calsequestrin in the lethal arrhythmic effects of cocaine. 2013 , 133, 344-51 | 10 |
| 1634 | Structural and functional insights on folate receptor (FR) by homology modeling, ligand docking and molecular dynamics. 2013 , 44, 197-207 | 19 |
| 1633 | The DNA binding mechanism of a SSB protein from <i>Lactococcus lactis</i> siphophage p2. 2013 , 1834, 1070-6 | 6 |
| 1632 | Computational analysis of protein-protein interfaces involving an alpha helix: insights for terphenyl-like molecules binding. 2013 , 14, 31 | 7 |

| | | |
|------|---|----|
| 1631 | Calix[4]arene methylenebisphosphonic acids as inhibitors of protein tyrosine phosphatase 1B. 2013 , 23, 5619-23 | 26 |
| 1630 | From natural products to drugs for epimutation computer-aided drug design. 2013 , 170, 164-75 | 20 |
| 1629 | Evidence of colorectal cancer-associated mutation in MCAK: a computational report. 2013 , 67, 837-51 | 29 |
| 1628 | Identification of protease inhibitors by a fast fluorimetric assay. 2013 , 54, 283-91 | 1 |
| 1627 | Synthesis of new series of pyrimido[4,5-b][1,4] benzothiazines as 15-lipoxygenase inhibitors and study of their inhibitory mechanism. 2013 , 22, 5036-5043 | 16 |
| 1626 | Pharmacophore model generation for microtubule-stabilizing anti-mitotic agents (MSAAs) against ovarian cancer. 2013 , 22, 4322-4330 | 2 |
| 1625 | Molecular modeling of the <i>Candida albicans</i> Vps4 and a virtual screening study for novel inhibitors. 2013 , 22, 4529-4536 | 1 |
| 1624 | Anti-melanogenic effect of (Z)-5-(2,4-dihydroxybenzylidene) thiazolidine-2,4-dione, a novel tyrosinase inhibitor. 2013 , 36, 1189-97 | 26 |
| 1623 | Antimicrobial dyes and mechanosensitive channels. 2013 , 104, 155-67 | 12 |
| 1622 | Conformational mobility of active and E-64-inhibited actinidin. 2013 , 1830, 4790-9 | 6 |
| 1621 | Kinetics and molecular docking studies of an anti-diabetic complication inhibitor fucosterol from edible brown algae <i>Eisenia bicyclis</i> and <i>Ecklonia stolonifera</i> . 2013 , 206, 55-62 | 45 |
| 1620 | Structure based design towards the identification of novel binding sites and inhibitors for the chikungunya virus envelope proteins. 2013 , 44, 241-52 | 25 |
| 1619 | Interaction between toxic azo dye C.I. Acid Red 88 and serum albumins. 2013 , 143, 715-722 | 16 |
| 1618 | Ligand-induced structural changes in TEM-1 probed by molecular dynamics and relative binding free energy calculations. 2013 , 53, 2648-58 | 6 |
| 1617 | In silico study on multidrug resistance conferred by I223R/H275Y double mutant neuraminidase. 2013 , 9, 2764-74 | 14 |
| 1616 | X-ray crystal structure of phosphodiesterase 2 in complex with a highly selective, nanomolar inhibitor reveals a binding-induced pocket important for selectivity. 2013 , 135, 11708-11 | 46 |
| 1615 | Interactions of acetylcholine binding site residues contributing to nicotinic acetylcholine receptor gating: role of residues Y93, Y190, K145 and D200. 2013 , 44, 145-54 | 7 |
| 1614 | Insights into the glycyl radical enzyme active site of benzylsuccinate synthase: a computational study. 2013 , 135, 12279-88 | 26 |

| | | |
|------|--|----|
| 1613 | [Ru(bpy) ₂ dppz-idzo] ²⁺ : a colorimetric molecular "light switch" and powerful stabilizer for G-quadruplex DNA. 2013 , 42, 5661-72 | 54 |
| 1612 | An in silico approach for understanding the molecular evolution of clinically important metallo-beta-lactamases. 2013 , 20, 39-47 | 13 |
| 1611 | Novel CYP2B6 enzyme variants in a Rwandese population: functional characterization and assessment of in silico prediction tools. 2013 , 34, 725-34 | 22 |
| 1610 | Syntheses and crystal structures of tetracopper(II) complexes bridged by asymmetric N,N'-bis(substituted)oxamides: Molecular docking, DNA-binding and in vitro anticancer activity. 2013 , 128, 97-107 | 12 |
| 1609 | Search for invisible binding sites of low-molecular-weight compounds on protein molecules and prediction of inhibitory activity. 2013 , 47, 592-598 | 2 |
| 1608 | Synthesis, hypoglycemic activity and molecular modeling studies of pyrazole-3-carbohydrazides designed by a CoMFA model. 2013 , 69, 10-21 | 33 |
| 1607 | Tuning the substrate selectivity of meta-cleavage product hydrolase by domain swapping. 2013 , 97, 5343-50 | 4 |
| 1606 | Three-component synthesis of pyrano[2,3-d]-pyrimidine dione derivatives facilitated by sulfonic acid nanoporous silica (SBA-Pr-SO ₃ H) and their docking and urease inhibitory activity. 2013 , 21, 3 | 62 |
| 1605 | Engineering of chitosan and collagen macromolecules using sebacic acid for clinical applications. 2013 , 2, 11 | 22 |
| 1604 | Extrapolating the effect of deleterious nsSNPs in the binding adaptability of flavopiridol with CDK7 protein: a molecular dynamics approach. 2013 , 7, 10 | 40 |
| 1603 | In silico analysis of the three-dimensional structures of the homodimer of uridine phosphorylase from <i>Yersinia Pseudotuberculosis</i> in the ligand-free state and in a complex with 5-fluorouracil. 2013 , 58, 280-286 | |
| 1602 | Chemical proteomics reveals HSP70 1A as a target for the anticancer diterpene oridonin in Jurkat cells. 2013 , 82, 14-26 | 48 |
| 1601 | Design and evaluation of substrate-based octapeptide and non substrate-based tetrapeptide inhibitors of dengue virus NS2B-NS3 proteases. 2013 , 434, 767-72 | 32 |
| 1600 | Synthesis and biological activity of new resveratrol derivative and molecular docking: dynamics studies on NFκB. 2013 , 171, 1639-57 | 9 |
| 1599 | Understanding molecular mechanisms of traditional Chinese medicine for the treatment of influenza viruses infection by computational approaches. 2013 , 9, 2696-700 | 22 |
| 1598 | Synthesis of novel psoralen analogues and their in vitro antitumor activity. 2013 , 21, 5047-53 | 15 |
| 1597 | The emerging role of cloud computing in molecular modelling. 2013 , 44, 177-87 | 23 |
| 1596 | In silico characterization of a novel α -1,3-glucanase gene from <i>Bacillus amyloliquefaciens</i> --a bacterial endophyte of <i>Hevea brasiliensis</i> antagonistic to <i>Phytophthora meadii</i> . 2013 , 19, 999-1007 | 9 |

| | | | | |
|------|--|-----|--|----|
| 1595 | Constant pH molecular dynamics (CpHMD) and molecular docking studies of CquiOBP1 pH-induced ligand releasing mechanism. 2013 , 19, 1301-9 | | | 10 |
| 1594 | Insights into the specific binding site of adenosine to the Stx2, the protein toxin from Escherichia coli O157:H7 using molecular dynamics simulations and free energy calculations. 2013 , 39, 199-205 | | | 5 |
| 1593 | Structural insights into the role of Bacillus subtilis YwfH (BacG) in tetrahydrotyrosine synthesis. 2013 , 69, 324-32 | | | 7 |
| 1592 | The effects of R683S (G) genetic mutations on the JAK2 activity, structure and stability. 2013 , 60, 186-95 | | | 14 |
| 1591 | Structural investigation of idarubicin-DNA interaction: spectroscopic and molecular docking study. 2013 , 60, 213-8 | | | 25 |
| 1590 | Discovery of a novel activator of 5-lipoxygenase from an anacardic acid derived compound collection. 2013 , 21, 7763-78 | | | 25 |
| 1589 | Identification of sulfonic acids as efficient ecto-5'-nucleotidase inhibitors. 2013 , 70, 685-91 | | | 27 |
| 1588 | Structural and energetic analyses of SNPs in drug targets and implications for drug therapy. 2013 , 53, 3343-51 | | | 22 |
| 1587 | Message passing interface and multithreading hybrid for parallel molecular docking of large databases on petascale high performance computing machines. <i>Journal of Computational Chemistry</i> , 2013 , 34, 915-27 | 3.5 | | 51 |
| 1586 | The impact of nitric oxide toxicity on the evolution of the glutathione transferase superfamily: a proposal for an evolutionary driving force. 2013 , 288, 24936-47 | | | 23 |
| 1585 | Exploring the chemical space of G-quadruplex binders: discovery of a novel chemotype targeting the human telomeric sequence. 2013 , 56, 9646-54 | | | 44 |
| 1584 | Design and synthesis of novel 3,5-bis-N-(aryl/heteroaryl) carbamoyl-4-aryl-1,4-dihydropyridines as small molecule BACE-1 inhibitors. 2013 , 21, 6893-909 | | | 17 |
| 1583 | Towards dual antithrombotic compounds - balancing thrombin inhibitory and fibrinogen GPIIb/IIIa binding inhibitory activities of 2,3-dihydro-1,4-benzodioxine derivatives through regio- and stereoisomerism. 2013 , 62, 329-40 | | | 16 |
| 1582 | GalaxyDock2: protein-ligand docking using beta-complex and global optimization. <i>Journal of Computational Chemistry</i> , 2013 , 34, 2647-56 | 3.5 | | 44 |
| 1581 | Rational design of chemical genetic probes of RNA function and lead therapeutics targeting repeating transcripts. 2013 , 18, 1228-36 | | | 22 |
| 1580 | Biodegradation of Lignin by Laccase for Conversion of Biomass to Fuel: Analysis of Substrate Binding. 2013 , 53, 1-10 | | | 1 |
| 1579 | Combining structure- and ligand-based approaches for studies of interactions between different conformations of the hERG K ⁺ channel pore and known ligands. 2013 , 46, 93-104 | | | 14 |
| 1578 | Similarity-based virtual screening for microtubule stabilizers reveals novel antimitotic scaffold. 2013 , 44, 188-96 | | | 14 |

| | | |
|------|---|-----|
| 1577 | Virtual screening for alpha7 nicotinic acetylcholine receptor for treatment of Alzheimer's disease. 2013 , 39, 98-107 | 10 |
| 1576 | Modeling iron-catecholates binding to NGAL protein. 2013 , 45, 111-21 | 13 |
| 1575 | Computation of binding energies including their enthalpy and entropy components for protein-ligand complexes using support vector machines. 2013 , 53, 2559-70 | 17 |
| 1574 | How conformational flexibility stabilizes the hyperthermophilic elongation factor G-domain. 2013 , 117, 13775-85 | 29 |
| 1573 | Investigation of aryl halides as ketone bioisosteres: refinement of potent and selective inhibitors of human cytochrome P450 19A1 (aromatase). 2013 , 23, 6060-3 | 7 |
| 1572 | Lessons learned in empirical scoring with smina from the CSAR 2011 benchmarking exercise. 2013 , 53, 1893-904 | 354 |
| 1571 | New 7-methylguanine derivatives targeting the influenza polymerase PB2 cap-binding domain. 2013 , 56, 8915-30 | 57 |
| 1570 | 6-Cyclohexylmethoxy-5-(cyano-NNO-azoxy)pyrimidine-4-amine: a new scaffold endowed with potent CDK2 inhibitory activity. 2013 , 68, 333-8 | 14 |
| 1569 | Binding mechanism and synergetic effects of xanthone derivatives as noncompetitive β -glucosidase inhibitors: a theoretical and experimental study. 2013 , 117, 13464-71 | 33 |
| 1568 | 17(E)-picolinylidene androstane derivatives as potential inhibitors of prostate cancer cell growth: antiproliferative activity and molecular docking studies. 2013 , 21, 7257-66 | 24 |
| 1567 | Novel anti-plasmodial hits identified by virtual screening of the ZINC database. 2013 , 27, 859-71 | 13 |
| 1566 | Phenobarbital indirectly activates the constitutive active androstane receptor (CAR) by inhibition of epidermal growth factor receptor signaling. 2013 , 6, ra31 | 138 |
| 1565 | Inhibition of dipeptidyl peptidase IV and xanthine oxidase by amino acids and dipeptides. 2013 , 141, 644-53 | 91 |
| 1564 | Optical control of an ion channel gate. 2013 , 110, 20813-8 | 50 |
| 1563 | In Silico Models for Drug Discovery. 2013 , | 4 |
| 1562 | Some insights into the binding mechanism of the GABAA receptor: a combined docking and MM-GBSA study. 2013 , 19, 5489-500 | 6 |
| 1561 | Enthalpic signature of methonium desolvation revealed in a synthetic host-guest system based on cucurbit[7]uril. 2013 , 135, 6084-91 | 8 |
| 1560 | Molecular basis for benzimidazole resistance from a novel β -tubulin binding site model. 2013 , 45, 26-37 | 40 |

| | | |
|------|--|-----|
| 1559 | Computational methods of studying the binding of toxins from venomous animals to biological ion channels: theory and applications. 2013 , 93, 767-802 | 42 |
| 1558 | Controlling the Fate of Protein Corona by Tuning Surface Properties of Nanoparticles. 2013 , 4, 3747-3752 | 41 |
| 1557 | Design of novel potent inhibitors of human uridine phosphorylase-1: synthesis, inhibition studies, thermodynamics, and in vitro influence on 5-fluorouracil cytotoxicity. 2013 , 56, 8892-902 | 14 |
| 1556 | EpiDOCK: a molecular docking-based tool for MHC class II binding prediction. 2013 , 26, 631-4 | 36 |
| 1555 | Inhibition of human alcohol and aldehyde dehydrogenases by acetaminophen: Assessment of the effects on first-pass metabolism of ethanol. 2013 , 47, 559-65 | 25 |
| 1554 | Shaping the interaction landscape of bioactive molecules. 2013 , 29, 3073-9 | 185 |
| 1553 | Nonlinear scoring functions for similarity-based ligand docking and binding affinity prediction. 2013 , 53, 3097-112 | 38 |
| 1552 | Mn ²⁺ and Mg ²⁺ improved sphingomyelinase production by <i>Lactobacillus rhamnosus</i> FTDC 8313 and binding affinity to sphingomyelin for generation of ceramides. 2013 , 48, 1815-1821 | 4 |
| 1551 | Mutational analysis of the binding pockets of the diketo acid inhibitor L-742,001 in the influenza virus PA endonuclease. 2013 , 87, 10524-38 | 56 |
| 1550 | Theoretical studies on the binding of rhenium(II) complexes to inducible nitric oxide synthase. 2013 , 45, 13-25 | 12 |
| 1549 | Collaborative Drug Design of Plasmodium Kinase Inhibitors. 2013 , 375-416 | |
| 1548 | Homology modeling, molecular docking and MD simulation studies to investigate role of cysteine protease from <i>Xanthomonas campestris</i> in degradation of A β peptide. 2013 , 43, 2063-70 | 35 |
| 1547 | Structure-activity relationship analysis of curcumin analogues on anti-influenza virus activity. 2013 , 280, 5829-40 | 61 |
| 1546 | Isoform selectivity of adenylyl cyclase inhibitors: characterization of known and novel compounds. 2013 , 347, 265-75 | 49 |
| 1545 | Optimized CGenFF force-field parameters for acylphosphate and N-phosphonosulfonimidoyl functional groups. 2013 , 19, 5075-87 | 1 |
| 1544 | A knowledge-based halogen bonding scoring function for predicting protein-ligand interactions. 2013 , 19, 5015-30 | 32 |
| 1543 | Structural analysis of the novel influenza A (H7N9) viral Neuraminidase interactions with current approved neuraminidase inhibitors Oseltamivir, Zanamivir, and Peramivir in the presence of mutation R289K. 2013 , 14 Suppl 16, S7 | 24 |
| 1542 | Detection of persistent organic pollutants binding modes with androgen receptor ligand binding domain by docking and molecular dynamics. 2013 , 13, 16 | 6 |

| | | |
|------|---|----|
| 1541 | Identifying ligand binding sites and poses using GPU-accelerated Hamiltonian replica exchange molecular dynamics. 2013 , 27, 989-1007 | 86 |
| 1540 | Drug-target and disease networks: polypharmacology in the post-genomic era. 2013 , 1, 17 | 31 |
| 1539 | Effect of intracellular loop 3 on intrinsic dynamics of human β_2 -adrenergic receptor. 2013 , 13, 29 | 21 |
| 1538 | Molecular docking and simulation studies towards exploring antiviral compounds against envelope protein of Japanese encephalitis virus. 2013 , 2, 231-243 | 7 |
| 1537 | Characterization of PfTrxR inhibitors using antimalarial assays and in silico techniques. 2013 , 7, 175 | 9 |
| 1536 | Analysis of bortezomib inhibitor docked within the catalytic subunits of the Plasmodium falciparum 20S proteasome. 2013 , 2, 566 | 2 |
| 1535 | Experimental and computational studies on newly synthesized resveratrol derivative: a new method for cancer chemoprevention and therapeutics?. 2013 , 17, 568-83 | 11 |
| 1534 | Raltegravir flexibility and its impact on recognition by the HIV-1 IN targets. 2013 , 26, 383-401 | 1 |
| 1533 | Melatonin inhibits matrix metalloproteinase-9 activity by binding to its active site. 2013 , 54, 398-405 | 74 |
| 1532 | Drug-DNA intercalation: from discovery to the molecular mechanism. 2013 , 92, 1-62 | 47 |
| 1531 | Unusual binding of a potential biomarker with human serum albumin. 2013 , 8, 728-35 | 10 |
| 1530 | Binding of the repressor complex REST-mSIN3b by small molecules restores neuronal gene transcription in Huntington's disease models. 2013 , 127, 22-35 | 33 |
| 1529 | Inhibitors of the aminoglycoside 6'-N-acetyltransferase type Ib [AAC(6')-Ib] identified by in silico molecular docking. 2013 , 23, 5694-8 | 21 |
| 1528 | The enzyme 3-hydroxykynurenine transaminase as potential target for 1,2,4-oxadiazoles with larvicidal activity against the dengue vector Aedes aegypti. 2013 , 21, 6996-7003 | 14 |
| 1527 | Quantification of Solvent Contribution to the Stability of Noncovalent Complexes. 2013 , 9, 4542-51 | 33 |
| 1526 | 2-(2-Hydrazinyl)thiazole derivatives: design, synthesis and in vitro antimycobacterial studies. 2013 , 69, 564-76 | 72 |
| 1525 | Monoamine oxidase A and B inhibiting effect and molecular modeling of some synthesized coumarin derivatives. 2013 , 62, 198-209 | 22 |
| 1524 | Combining molecular docking and 3-D pharmacophore generation to enclose the in vivo antigenotoxic activity of naturally occurring aromatic compounds: myricetin, quercetin, rutin, and rosmarinic acid. 2013 , 86, 1376-96 | 20 |

| | | |
|------|--|----|
| 1523 | Discovery of triazine mimetics as potent antileishmanial agents. 2013 , 4, 1108-13 | 26 |
| 1522 | Molecular modeling of butyrylcholinesterase inhibition by cresyl saligenin phosphate. 2013 , 62, 2527-2537 | 17 |
| 1521 | Diversity oriented approach to triazole based peptidomimetics as mammalian sterile 20 kinase inhibitors. 2013 , 3, 24447 | 11 |
| 1520 | Highly efficient and regioselective enzymatic synthesis of β -(1 \rightarrow 3) galactosides in biosolvents. 2013 , 3, 12155 | 15 |
| 1519 | Immunosuppression by co-stimulatory molecules: inhibition of CD2-CD48/CD58 interaction by peptides from CD2 to suppress progression of collagen-induced arthritis in mice. 2013 , 82, 106-18 | 19 |
| 1518 | Analysis of decapping scavenger cap complex using modified cap analogs reveals molecular determinants for efficient cap binding. 2013 , 280, 6508-27 | 13 |
| 1517 | Modulation of excimer formation of 9-(dicyano-vinyl)julolidine by the macrocyclic hosts. 2013 , 15, 330-40 | 19 |
| 1516 | How Do Perfluorinated Alkanoic Acids Elicit Cytochrome P450 to Catalyze Methane Hydroxylation? An MD and QM/MM Study. 2013 , 3, 2995-3005 | 18 |
| 1515 | Identifying key amino acid residues that affect α -conotoxin AulB inhibition of α 4 nicotinic acetylcholine receptors. 2013 , 288, 34428-42 | 40 |
| 1514 | The Effect of Pimelic Acid Interaction on the Mechanical and Thermal Properties of Chitosan and Collagen. 2013 , 62, 572-582 | 17 |
| 1513 | Improving Escherichia coli FucO for furfural tolerance by saturation mutagenesis of individual amino acid positions. 2013 , 79, 3202-8 | 25 |
| 1512 | 3-Aryl-N-aminoylsulfonylphenyl-1H-pyrazole-5-carboxamides: a new class of selective Rac inhibitors. 2013 , 4, 537 | 21 |
| 1511 | Structure and function of an insect α -carboxylesterase (α esterase7) associated with insecticide resistance. 2013 , 110, 10177-82 | 84 |
| 1510 | Aspergillus niger β -glucosidase has a cellulase-like tadpole molecular shape: insights into glycoside hydrolase family 3 (GH3) β -glucosidase structure and function. 2013 , 288, 32991-3005 | 49 |
| 1509 | Probing the steric requirements of the β -aminobutyric acid aminotransferase active site with fluorinated analogues of vigabatrin. 2013 , 21, 903-11 | 11 |
| 1508 | G4LDB: a database for discovering and studying G-quadruplex ligands. 2013 , 41, D1115-23 | 99 |
| 1507 | Unraveling the binding interaction and kinetics of a prospective anti-HIV drug with a model transport protein: results and challenges. 2013 , 15, 1275-87 | 90 |
| 1506 | Anticoagulant activity of a unique sulfated pyranosic (1 \rightarrow 3)- β -L-arabinan through direct interaction with thrombin. 2013 , 288, 223-33 | 33 |

| | | | |
|------|---|-----|----|
| 1505 | Comprehensive classification and diversity assessment of atomic contacts in protein-small ligand interactions. 2013 , 53, 241-8 | | 10 |
| 1504 | In silico identification of novel and selective monoamine oxidase B inhibitors. 2013 , 120, 853-8 | | 8 |
| 1503 | Chemiosostericism in the proteome. 2013 , 53, 279-92 | | 15 |
| 1502 | Interactions of vasopressin and oxytocin receptors with vasopressin analogues substituted in position 2 with 3,3'-diphenylalanine--a molecular docking study. 2013 , 19, 118-26 | | 6 |
| 1501 | Increased hydrophobicity and estrogenic activity of simple phenols with silicon and germanium-containing substituents. 2013 , 56, 160-6 | | 43 |
| 1500 | Molecular docking and NMR binding studies to identify novel inhibitors of human phosphomevalonate kinase. 2013 , 430, 313-9 | | 4 |
| 1499 | Solvent structure improves docking prediction in lectin-carbohydrate complexes. 2013 , 23, 241-58 | | 32 |
| 1498 | Selective antitumour activity and ER β -molecular docking studies of newly synthesized D-homo fused steroidal tetrazoles. 2013 , 4, 317-323 | | 22 |
| 1497 | Studies on ornithine decarboxylase of <i>Leishmania donovani</i> : structure modeling and inhibitor docking. 2013 , 22, 466-478 | | 10 |
| 1496 | QM study and conformational analysis of an isatin Schiff base as a potential cytotoxic agent. 2013 , 19, 727-35 | | 66 |
| 1495 | Theoretical study of zeatin [A plant hormone and potential drug for neural diseases] [On the basis of DFT, MP2 and target docking. 2013 , 557, 140-144 | | 4 |
| 1494 | Functional analysis of a mosquito short-chain dehydrogenase cluster. 2013 , 82, 96-115 | | 13 |
| 1493 | CovalentDock: automated covalent docking with parameterized covalent linkage energy estimation and molecular geometry constraints. <i>Journal of Computational Chemistry</i> , 2013 , 34, 326-36 | 3-5 | 98 |
| 1492 | Biaryl tetrazolyl ureas as inhibitors of endocannabinoid metabolism: modulation at the N-portion and distal phenyl ring. 2013 , 63, 118-32 | | 12 |
| 1491 | Conserved water mediated H-bonding dynamics of Ser117 and Thr119 residues in human transthyretin-thyroxin complexation: inhibitor modeling study through docking and molecular dynamics simulation. 2013 , 44, 70-80 | | 7 |
| 1490 | Targeting cathepsin E in pancreatic cancer by a small molecule allows in vivo detection. 2013 , 15, 684-93 | | 29 |
| 1489 | The peroxisomal enzyme L-PBE is required to prevent the dietary toxicity of medium-chain fatty acids. 2013 , 5, 248-58 | | 32 |
| 1488 | Mechanochemical activation of vincamine mediated by linear polymers: assessment of some "critical" steps. 2013 , 50, 56-68 | | 10 |

| | | |
|------|--|----|
| 1487 | Differential interactions of the broad spectrum drugs artemisinin, dihydroartemisinin and artesunate with serum albumin. 2013 , 20, 969-74 | 12 |
| 1486 | The second messenger phosphatidylinositol-5-phosphate facilitates antiviral innate immune signaling. 2013 , 14, 148-58 | 32 |
| 1485 | In silico metabolism studies of dietary flavonoids by CYP1A2 and CYP2C9. 2013 , 50, 102-110 | 20 |
| 1484 | Development of novel molecular probes of the Rio1 atypical protein kinase. 2013 , 1834, 1292-301 | 8 |
| 1483 | Trp266 determines the binding specificity of a porcine aflatoxin B ₁ aldehyde reductase for aflatoxin B ₁ aldehyde. 2013 , 86, 1357-65 | 11 |
| 1482 | Highly efficient Michael-type addition of acetaldehyde to α -nitrostyrenes by whole resting cells of Escherichia coli expressing 4-oxalocrotonate tautomerase. 2013 , 142, 462-8 | 20 |
| 1481 | Static and dynamic interactions between GALK enzyme and known inhibitors: guidelines to design new drugs for galactosemic patients. 2013 , 63, 423-34 | 23 |
| 1480 | Molecular dynamics simulations reveal insight into key structural elements of aaptamines as sortase inhibitors with free energy calculations. 2013 , 585, 171-177 | 8 |
| 1479 | Design of novel Geldanamycin analogue hsp90 alpha-inhibitor in silico for breast cancer therapy. 2013 , 81, 463-9 | 7 |
| 1478 | DINC: a new AutoDock-based protocol for docking large ligands. 2013 , 13 Suppl 1, S11 | 32 |
| 1477 | Modeling, synthesis, and biological evaluation of potential retinoid X receptor (RXR) selective agonists: novel analogues of 4-[1-(3,5,5,8,8-pentamethyl-5,6,7,8-tetrahydro-2-naphthyl)ethynyl]benzoic acid (bexarotene) and (E)-3-(3-(1,2,3,4-tetrahydro-1,1,4,4,6-pentamethylnaphthalen-7-yl)-4-hydroxyphenyl)acrylic acid | 17 |
| 1476 | Synthesis, chemical characterization, DNA binding and antioxidant studies of ferrocene incorporated selenoure. 2013 , 1048, 367-374 | 25 |
| 1475 | Structure-dependent activities of hydroxylated polybrominated diphenyl ethers on human estrogen receptor. 2013 , 309, 15-22 | 33 |
| 1474 | Diversity and analysis of sequences encoded by arcelin genes from Indian wild pulses resistant to bruchids. 2013 , 48, 1697-1705 | 3 |
| 1473 | Evaluation of molecular model-based discovery of ecto-5'-nucleotidase inhibitors on the basis of X-ray structures. 2013 , 21, 6616-22 | 7 |
| 1472 | Computational evaluation of small molecule inhibitors of RGS4 to regulate the dopaminergic control of striatal LTD. 2013 , 14, 135-142 | 1 |
| 1471 | Molecular determinants for nuclear receptors selectivity: chemometric analysis, dockings and site-directed mutagenesis of dual peroxisome proliferator-activated receptors α agonists. 2013 , 63, 321-32 | 16 |
| 1470 | Carbonyl reduction of triadimefon by human and rodent 11 β -hydroxysteroid dehydrogenase 1. 2013 , 85, 1370-8 | 10 |

| | | |
|------|---|----|
| 1469 | Identification of RC-33 as a potent and selective α receptor agonist potentiating NGF-induced neurite outgrowth in PC12 cells. Part 2: g-scale synthesis, physicochemical characterization and in vitro metabolic stability. 2013 , 21, 2577-86 | 35 |
| 1468 | Design and synthesis of silicon-containing tubulin polymerization inhibitors: replacement of the ethylene moiety of combretastatin A-4 with a silicon linker. 2013 , 21, 7381-91 | 27 |
| 1467 | Parametrisation of the free energy of ATP binding to wild-type and mutant Kir6.2 potassium channels. 2013 , 171, 76-83 | 2 |
| 1466 | Insight into the dynamic interaction between different flavonoids and bovine serum albumin using molecular dynamics simulations and free energy calculations. 2013 , 19, 1039-47 | 20 |
| 1465 | Collaborative virtual organisation and infrastructure for drug discovery. 2013 , 18, 681-6 | 3 |
| 1464 | Molecular Dynamics Investigation on the Inhibition of MDM2-p53 Interaction by Polyphenols. 2013 , 32, 203-12 | 4 |
| 1463 | Farnesyl diphosphate synthase inhibitors from in silico screening. 2013 , 81, 742-8 | 39 |
| 1462 | An automated docking protocol for hERG channel blockers. 2013 , 53, 159-75 | 16 |
| 1461 | Accounting for receptor flexibility and enhanced sampling methods in computer-aided drug design. 2013 , 81, 41-9 | 90 |
| 1460 | Structure-based identification of aporphines with selective 5-HT(2A) receptor-binding activity. 2013 , 81, 250-6 | 21 |
| 1459 | Quantitative analysis of pheromone-binding protein specificity. 2013 , 22, 31-40 | 13 |
| 1458 | Enrichment factor analyses on G-protein coupled receptors with known crystal structure. 2013 , 53, 739-43 | 19 |
| 1457 | Substrate specificity and the effect of calcium on Trypanosoma brucei metacaspase 2. 2013 , 280, 2608-21 | 17 |
| 1456 | Tuning the Bioactivity of Tensioactive Deoxy Glycosides to Structure: Antibacterial Activity Versus Selective Cholinesterase Inhibition Rationalized by Molecular Docking. 2013 , 2013, 1448-1459 | 6 |
| 1455 | The impact of introducing a histidine into an apolar cavity site on docking and ligand recognition. 2013 , 56, 2874-84 | 7 |
| 1454 | Psoralen derivatives as inhibitors of NF- κ B/DNA interaction: synthesis, molecular modeling, 3D-QSAR, and biological evaluation. 2013 , 56, 1830-42 | 28 |
| 1453 | Ligand interaction, binding site and G protein activation of the mu opioid receptor. 2013 , 702, 309-15 | 25 |
| 1452 | Antitumor effects of a tetradentate amido-carboxylate ligands and corresponding square-planar palladium(II) complexes toward some cancer cells. Crystal structure, DFT modeling and ligand to DNA probe docking simulation. 2013 , 121, 134-44 | 29 |

| | | |
|------|--|-----|
| 1451 | Synthesis and biological evaluation of guanidino analogues of roscovitine. 2013 , 62, 443-52 | 9 |
| 1450 | Conformationally restricted short peptides inhibit human islet amyloid polypeptide (hIAPP) fibrillization. 2013 , 49, 2688-90 | 39 |
| 1449 | Polyhydroxylated pyrrolidine and 2-oxapyrrolizidine as glycosidase inhibitors. 2013 , 4, 783 | 6 |
| 1448 | X-ray crystallographic and molecular docking studies on a unique chloromuconolactone dehalogenase from <i>Rhodococcus opacus</i> 1CP. 2013 , 182, 44-50 | 2 |
| 1447 | Intercalation and de-intercalation pathway of proflavine through the minor and major grooves of DNA: roles of water and entropy. 2013 , 15, 6446-55 | 32 |
| 1446 | Quercetin and taxifolin completely break MDM2p53 association: molecular dynamics simulation study. 2013 , 22, 2778-2787 | 4 |
| 1445 | Small molecule regulation of protein conformation by binding in the Flap of HIV protease. 2013 , 8, 1223-31 | 27 |
| 1444 | Solution structures of <i>Mycobacterium tuberculosis</i> thioredoxin C and models of intact thioredoxin system suggest new approaches to inhibitor and drug design. 2013 , 81, 675-89 | 9 |
| 1443 | Novel analgesic/anti-inflammatory agents: 1,5-diarylpyrrole nitrooxyalkyl ethers and related compounds as cyclooxygenase-2 inhibiting nitric oxide donors. 2013 , 56, 3191-206 | 37 |
| 1442 | Constructing manmade enzymes for oxygen activation. 2013 , 42, 3136-50 | 19 |
| 1441 | Homoisocitrate dehydrogenase from <i>Candida albicans</i> : properties, inhibition, and targeting by an antifungal pro-drug. 2013 , 13, 143-55 | 9 |
| 1440 | Design and synthesis of dual-action inhibitors targeting histone deacetylases and 3-hydroxy-3-methylglutaryl coenzyme A reductase for cancer treatment. 2013 , 56, 3645-55 | 61 |
| 1439 | Rational discovery of dengue type 2 non-competitive inhibitors. 2013 , 82, 1-11 | 28 |
| 1438 | Protein engineering for metabolic engineering: current and next-generation tools. 2013 , 8, 545-55 | 33 |
| 1437 | Antimicrobial activity of phenolic compounds identified in wild mushrooms, SAR analysis and docking studies. 2013 , 115, 346-57 | 222 |
| 1436 | Genome comparisons as a tool for antimicrobial target discovery. 2013 , 993, 31-8 | |
| 1435 | Conjugate symbiotic populations part II: Analysis of nfr5 receptor gene polymorphisms using molecular docking. 2013 , 3, 146-151 | 0 |
| 1434 | Molecular recognition of rosmarinic acid from <i>Salvia sclareoides</i> extracts by acetylcholinesterase: a new binding site detected by NMR spectroscopy. 2013 , 19, 6641-9 | 29 |

| | | |
|------|--|----|
| 1433 | Molecular dynamics simulations in drug design. 2013 , 993, 95-113 | 43 |
| 1432 | Structural characterization of 2,6-dichloro-p-hydroquinone 1,2-dioxygenase (PcpA) from <i>Sphingobium chlorophenicum</i> , a new type of aromatic ring-cleavage enzyme. 2013 , 88, 523-36 | 22 |
| 1431 | A chemical-biological study reveals C9-type iridoids as novel heat shock protein 90 (Hsp90) inhibitors. 2013 , 56, 1583-95 | 45 |
| 1430 | Investigation on the interaction behavior between curcumin and PAMAM dendrimer by spectral and docking studies. 2013 , 108, 251-5 | 22 |
| 1429 | Synthesis and evaluation of 7-chloro-4-(piperazin-1-yl)quinoline-sulfonamide as hybrid antiprotozoal agents. 2013 , 21, 3080-9 | 46 |
| 1428 | Correlation between sequence, structure and function for trisporoid processing proteins in the model zygomycete <i>Mucor mucedo</i> . 2013 , 320, 66-75 | 6 |
| 1427 | Design, development and evaluation of novel dual PPAR α /PPAR γ agonists. 2013 , 23, 873-9 | 12 |
| 1426 | A derivative of the natural compound kakuol affects DNA relaxation of topoisomerase IB inhibiting the cleavage reaction. 2013 , 530, 7-12 | 13 |
| 1425 | Structure-function analyses and molecular modeling of caffeic acid-O-methyltransferase and caffeoyl-CoA-O-methyltransferase: revisiting the basis of alternate methylation pathways during monolignol biosynthesis. 2013 , 60, 170-89 | 6 |
| 1424 | How to improve docking accuracy of AutoDock4.2: a case study using different electrostatic potentials. 2013 , 53, 188-200 | 70 |
| 1423 | New molecular scaffolds for the design of Alzheimer's acetylcholinesterase inhibitors identified using ligand- and receptor-based virtual screening. 2013 , 22, 2328-2345 | 28 |
| 1422 | Rational design of small-molecule inhibitors for β -catenin/T-cell factor protein-protein interactions by bioisostere replacement. 2013 , 8, 524-9 | 39 |
| 1421 | Pharmacophore assessment through 3-D QSAR: evaluation of the predictive ability on new derivatives by the application on a series of antitubercular agents. 2013 , 53, 1463-74 | 7 |
| 1420 | Designing novel inhibitors of <i>Trypanosoma brucei</i> . 2013 , 993, 231-43 | 1 |
| 1419 | Peptide inhibitors of the Keap1-Nrf2 protein-protein interaction with improved binding and cellular activity. 2013 , 11, 3553-7 | 69 |
| 1418 | Synthesis and characterization of a novel inhibitor of C-reactive protein-mediated proinflammatory effects. 2013 , 11, 177-84 | 6 |
| 1417 | Identification of sumoylation activating enzyme 1 inhibitors by structure-based virtual screening. 2013 , 53, 809-20 | 37 |
| 1416 | Kinetics and molecular docking studies of pimarane-type diterpenes as protein tyrosine phosphatase (PTP1B) inhibitors from <i>Aralia continentalis</i> roots. 2013 , 36, 957-65 | 17 |

| | | |
|------|--|----|
| 1415 | Highly efficient enzymatic synthesis of Gal β -(1 \rightarrow 3)-GalNAc and Gal β -(1 \rightarrow 3)-GlcNAc in ionic liquids. 2013 , 69, 4973-4978 | 24 |
| 1414 | Thermodynamic computational approach to capture molecular recognition in the binding of different inhibitors to the DNA gyrase B subunit from Escherichia coli. 2013 , 19, 3187-200 | 2 |
| 1413 | Computationally efficient methodology for atomic-level characterization of dendrimer-drug complexes: a comparison of amine- and acetyl-terminated PAMAM. 2013 , 117, 6801-13 | 67 |
| 1412 | Interaction of bovine serum albumin with a psychotropic drug alprazolam: Physicochemical, photophysical and molecular docking studies. 2013 , 142, 220-230 | 61 |
| 1411 | Inhibition of CK2 Activity by TCDD via binding to ATP-competitive binding site of catalytic subunit: Insight from computational studies. 2013 , 29, 299-306 | 1 |
| 1410 | Stepwise identification of potent antimicrobial peptides from human genome. 2013 , 113, 1-8 | 11 |
| 1409 | Binding of bis-ANS to Bacillus subtilis lipase: a combined computational and experimental investigation. 2013 , 1834, 1501-9 | 5 |
| 1408 | Homology modeling of dihydrofolate reductase from T. gondii bonded to antagonists: molecular docking and molecular dynamics simulations. 2013 , 9, 1308-15 | 10 |
| 1407 | Cytotoxic activity assessment, QSAR and docking study of novel bis-carboxamide derivatives of 4-pyrones synthesized by Ugi four-component reaction. 2013 , 66, 388-99 | 11 |
| 1406 | 2-Aminobenzothiazole derivatives: search for new antifungal agents. 2013 , 64, 357-64 | 54 |
| 1405 | Understanding traditional Chinese medicine anti-inflammatory herbal formulae by simulating their regulatory functions in the human arachidonic acid metabolic network. 2013 , 9, 1931-8 | 21 |
| 1404 | Insight into substrate preference of two chimeric esterases by combining experiment and molecular simulation. 2013 , 29, 533-537 | |
| 1403 | Compressed images for affinity prediction (CIFAP): a study on predicting binding affinities for checkpoint kinase 1 protein inhibitors. 2013 , 27, 155-164 | 4 |
| 1402 | Predicting the impact of single-nucleotide polymorphisms in CDK2-flavopiridol complex by molecular dynamics analysis. 2013 , 66, 681-95 | 12 |
| 1401 | Discovery and mechanism study of SIRT1 activators that promote the deacetylation of fluorophore-labeled substrate. 2013 , 56, 761-80 | 28 |
| 1400 | Phenylimino-2H-chromen-3-carboxamide derivatives as novel small molecule inhibitors of β -secretase (BACE1). 2013 , 21, 2396-2412 | 45 |
| 1399 | Synthesis, molecular modeling, and in vitro screening of monoamine oxidase inhibitory activities of some novel hydrazone derivatives. 2013 , 120, 883-91 | 10 |
| 1398 | Insights into the structural determinants for selective inhibition of nitric oxide synthase isoforms. 2013 , 19, 1537-51 | 12 |

| | | |
|------|---|-----|
| 1397 | Drug target prioritization in Plasmodium falciparum through metabolic network analysis, and inhibitor designing using virtual screening and docking approach. 2013 , 11, 1350003 | 2 |
| 1396 | Identification of function and mechanistic insights of guanine deaminase from Nitrosomonas europaea: role of the C-terminal loop in catalysis. 2013 , 52, 3512-22 | 9 |
| 1395 | Novel VEGFR-2 kinase inhibitors identified by the back-to-front approach. 2013 , 23, 2962-7 | 33 |
| 1394 | A new fluorescence "switch on" assay for heparin detection by using a functional ruthenium polypyridyl complex. 2013 , 138, 3483-9 | 23 |
| 1393 | Different molecular mechanisms of inhibition of bovine viral diarrhoea virus and hepatitis C virus RNA-dependent RNA polymerases by a novel benzimidazole. 2013 , 52, 3752-64 | 32 |
| 1392 | Prediction of cytochrome P450 xenobiotic metabolism: tethered docking and reactivity derived from ligand molecular orbital analysis. 2013 , 53, 1294-305 | 36 |
| 1391 | Discovery of thiazolidine-2,4-dione/biphenylcarbonitrile hybrid as dual PPAR α modulator with antidiabetic effect: in vitro, in silico and in vivo approaches. 2013 , 81, 474-83 | 42 |
| 1390 | Identification of key nodes of type 2 diabetes mellitus protein interactome and study of their interactions with phloridzin. 2013 , 17, 302-17 | 11 |
| 1389 | The evolutionary portrait of metazoan NAD salvage. 2013 , 8, e64674 | 8 |
| 1388 | Proposal for an inhibitor of Alzheimer's disease blocking aggregation of amyloid- β peptides: ab initio molecular simulations. 2013 , 433, 012033 | 1 |
| 1387 | Identification of benzofuran-3-yl(phenyl)methanones as novel SIRT1 inhibitors: binding mode, inhibitory mechanism and biological action. 2013 , 60, 441-50 | 17 |
| 1386 | Antituberculosis thiophenes define a requirement for Pks13 in mycolic acid biosynthesis. 2013 , 9, 499-506 | 95 |
| 1385 | A mechanistic hypothesis for the aspirin-induced switch in lipid mediator production by cyclooxygenase-2. 2013 , 135, 10404-10 | 16 |
| 1384 | Study of the interaction of Huperzia saururus Lycopodium alkaloids with the acetylcholinesterase enzyme. 2013 , 44, 136-44 | 12 |
| 1383 | Small-molecule ligand docking into comparative models with Rosetta. 2013 , 8, 1277-98 | 116 |
| 1382 | Experimental and computational correlation and prediction on herbicide resistance for acetoxyacid synthase mutants to Bispyribac. 2013 , 56, 286-295 | 8 |
| 1381 | Mechanism of retinoid X receptor partial agonistic action of 1-(3,5,5,8,8-pentamethyl-5,6,7,8-tetrahydro-2-naphthyl)-1H-benzotriazole-5-carboxylic acid and structural development to increase potency. 2013 , 56, 1865-77 | 28 |
| 1380 | Comparing the acceptor promiscuity of a Rosa hybrida glucosyltransferase RhGT1 and an engineered microbial glucosyltransferase OleD(PSA) toward a small flavonoid library. 2013 , 368, 73-7 | 11 |

| | | |
|------|---|-----|
| 1379 | Molecular docking characterizes substrate-binding sites and efflux modulation mechanisms within P-glycoprotein. 2013 , 53, 1747-60 | 111 |
| 1378 | Mechanism-based corrector combination restores β 508-CFTR folding and function. 2013 , 9, 444-54 | 275 |
| 1377 | Mahanine, a DNA minor groove binding agent exerts cellular cytotoxicity with involvement of C-7-OH and -NH functional groups. 2013 , 56, 5709-21 | 38 |
| 1376 | The dynamics of camphor in the cytochrome P450 CYP101D2. 2013 , 22, 1218-29 | 12 |
| 1375 | Virtual screening of low molecular weight mushrooms compounds as potential Mdm2 inhibitors. 2013 , 28, 569-75 | 10 |
| 1374 | Synthesis and crystal structure of a new copper(II) complex with N,N'-(4,4'-bithiazole-2,2'-diyl)diacetimidamide as ligand: Molecular docking, DNA-binding and cytotoxicity activity studies. 2013 , 1037, 15-22 | 17 |
| 1373 | Discovery of a New Class of Potent MMP Inhibitors by Structure-Based Optimization of the Arylsulfonamide Scaffold. 2013 , 4, 565-9 | 15 |
| 1372 | Substrate versus inhibitor dynamics of P-glycoprotein. 2013 , 81, 1653-68 | 30 |
| 1371 | Insights into the binding mode of new N-substituted pyrazoline derivatives to MAO-A: docking and quantum chemical calculations. 2013 , 120, 859-62 | 5 |
| 1370 | Evaluation of selective human MAO inhibitory activities of some novel pyrazoline derivatives. 2013 , 120, 863-73 | 11 |
| 1369 | Mechanism of the irreversible inhibition of human cyclooxygenase-1 by aspirin as predicted by QM/MM calculations. 2013 , 40, 99-109 | 38 |
| 1368 | Structural properties of the truncated and wild types of Taka-amylase: A molecular dynamics simulation and docking study. 2013 , 95, 36-40 | 11 |
| 1367 | Lipase from <i>Pseudomonas stutzeri</i> : Purification, homology modelling and rational explanation of the substrate binding mode. 2013 , 87, 88-98 | 26 |
| 1366 | Interactions of bacterial cell division protein FtsZ with C8-substituted guanine nucleotide inhibitors. A combined NMR, biochemical and molecular modeling perspective. 2013 , 135, 16418-28 | 26 |
| 1365 | Flooding enzymes: quantifying the contributions of interstitial water and cavity shape to ligand binding using extended linear response free energy calculations. 2013 , 53, 2349-59 | 8 |
| 1364 | Exploring the multifunctionality of thioflavin- and deferiprone-based molecules as acetylcholinesterase inhibitors for potential application in Alzheimer's disease. 2013 , 9, 792-805 | 22 |
| 1363 | CovalentDock Cloud: a web server for automated covalent docking. 2013 , 41, W329-32 | 23 |
| 1362 | Three-dimensional mapping of differential amino acids of human, murine, canine and equine TLR4/MD-2 receptor complexes conferring endotoxic activation by lipid A, antagonism by Eritoran and species-dependent activities of Lipid IVA in the mammalian LPS sensor system. 2013 , 7, e201305003 | 17 |

| | | |
|------|--|--------|
| 1361 | Crystal and solution studies of the "Plus-C" odorant-binding protein 48 from <i>Anopheles gambiae</i> : control of binding specificity through three-dimensional domain swapping. 2013 , 288, 33427-38 | 29 |
| 1360 | Molecular modelling and docking studies of an α -1,4-amylase from endophytic <i>Bacillus amyloliquefaciens</i> . 2013 , 7, 140-147 | |
| 1359 | A computational model for overcoming drug resistance using selective dual-inhibitors for aurora kinase A and its T217D variant. 2013 , 10, 4572-89 | 9 |
| 1358 | Assessment of estrogenic activity of perfluoroalkyl acids based on ligand-induced conformation state of human estrogen receptor. 2013 , 47, 634-41 | 36 |
| 1357 | DR-predictor: incorporating flexible docking with specialized electronic reactivity and machine learning techniques to predict CYP-mediated sites of metabolism. 2013 , 53, 3352-66 | 25 |
| 1356 | Prototropical and photophysical properties of ellipticine inside the nanocavities of molecular containers. 2013 , 117, 14099-107 | 17 |
| 1355 | Agensis of corpus callosum and optic nerve hypoplasia due to mutations in SLC25A1 encoding the mitochondrial citrate transporter. 2013 , 50, 240-5 | 45 |
| 1354 | Study on the interaction of the drug mesalamine with calf thymus DNA using molecular docking and spectroscopic techniques. 2013 , 128, 20-6 | 74 |
| 1353 | Virtual screening of novel reversible inhibitors for marine alkaline protease MP. 2013 , 46, 125-31 | 8 |
| 1352 | Solution structure and dynamics of human hemoglobin in the carbonmonoxy form. 2013 , 52, 5809-20 | 13 |
| 1351 | Rapid calculation of accurate atomic charges for proteins via the electronegativity equalization method. 2013 , 53, 2548-58 | 17 |
| 1350 | Physicochemical and thermodynamic characterization of the encapsulation of methyl jasmonate by natural and modified cyclodextrins using reversed-phase high-pressure liquid chromatography. 2013 , 61, 11347-54 | 20 |
| 1349 | Structural basis for enzyme I inhibition by β -ketoglutarate. 2013 , 8, 1232-40 | 24 |
| 1348 | FIPSDock: a new molecular docking technique driven by fully informed swarm optimization algorithm. <i>Journal of Computational Chemistry</i> , 2013 , 34, 67-75 | 3-5 50 |
| 1347 | Conformational analysis of furanoside-containing mono- and oligosaccharides. 2013 , 113, 1851-76 | 95 |
| 1346 | New dihydropyridine derivatives: anti-inflammatory, analgesic and docking studies. 2013 , 22, 1549-1562 | 6 |
| 1345 | Multidrug resistance protein P-gp interaction with nanoparticles (fullerenes and carbon nanotube) to assess their drug delivery potential: a theoretical molecular docking study. 2013 , 6, 343-57 | 27 |
| 1344 | Functional assignment of KEOPS/EKC complex subunits in the biosynthesis of the universal t6A tRNA modification. 2013 , 41, 9484-99 | 45 |

| | | |
|------|--|----|
| 1343 | Secondary anionic phospholipid binding site and gating mechanism in Kir2.1 inward rectifier channels. 2013 , 4, 2786 | 51 |
| 1342 | coREF: A COMBINED REFINEMENT APPROACH FOR GPCRS BACKBONE FLEXIBILITY. 2013 , 25, 1350047 | |
| 1341 | A two-hybrid approach to identify inhibitors of the RAS-RAF interaction. 2013 , 33 Pt A, 213-48 | 3 |
| 1340 | Berry and Citrus Phenolic Compounds Inhibit Dipeptidyl Peptidase IV: Implications in Diabetes Management. 2013 , 2013, 479505 | 76 |
| 1339 | Parallel strategies for an inverse docking method. 2013 , | 1 |
| 1338 | A data warehouse as an infrastructure to mine molecular descriptors for virtual screening. 2013 , | |
| 1337 | Ligand-Based Pharmacophore Modeling and Virtual Screening of RAD9 Inhibitors. 2013 , 2013, 1-7 | 4 |
| 1336 | A high performance cloud-based protein-ligand docking prediction algorithm. 2013 , 2013, 909717 | 3 |
| 1335 | Structural insights into the inhibition of type VI effector Tae3 by its immunity protein Tai3. 2013 , 454, 59-68 | 23 |
| 1334 | A small organic compound enhances the religation reaction of human topoisomerase I and identifies crucial elements for the religation mechanism. 2013 , 33, e00025 | 6 |
| 1333 | Ligand clouds around protein clouds: a scenario of ligand binding with intrinsically disordered proteins. 2013 , 9, e1003249 | 60 |
| 1332 | Antiangiogenic activity and pharmacogenomics of medicinal plants from traditional Korean medicine. 2013 , 2013, 131306 | 19 |
| 1331 | Discovery of anthelmintic drug targets and drugs using chokepoints in nematode metabolic pathways. 2013 , 9, e1003505 | 57 |
| 1330 | Structures of trans-2-enoyl-CoA reductases from <i>Clostridium acetobutylicum</i> and <i>Treponema denticola</i> : insights into the substrate specificity and the catalytic mechanism. 2013 , 449, 79-89 | 5 |
| 1329 | LigandRNA: computational predictor of RNA-ligand interactions. 2013 , 19, 1605-16 | 46 |
| 1328 | Transcriptional regulation of the <i>nadA</i> gene in <i>Neisseria meningitidis</i> impacts the prediction of coverage of a multicomponent meningococcal serogroup B vaccine. 2013 , 81, 560-9 | 44 |
| 1327 | The Effect of CYP2B6, CYP2D6, and CYP3A4 Alleles on Methadone Binding: A Molecular Docking Study. 2013 , 2013, 1-7 | 4 |
| 1326 | A Drug-Target Network-Based Approach to Evaluate the Efficacy of Medicinal Plants for Type II Diabetes Mellitus. 2013 , 2013, 203614 | 7 |

| | | |
|------|--|-----|
| 1325 | Sialyloxenitols as precursors for new analogues of sialidase inhibitors. 2013 , 85, 1803-1811 | 3 |
| 1324 | wFREDoW: a cloud-based web environment to handle molecular docking simulations of a fully flexible receptor model. 2013 , 2013, 469363 | 22 |
| 1323 | Computational Simulation on the Material Basis of Sini Decoction in Treating Coronary Heart Disease. 2013 , 278-280, 535-538 | 0 |
| 1322 | Fungicides Inhibition Analysis by Molecular Docking and Sensitivity Testing of <i>Penicillium italicum</i> . 2013 , 380-384, 4170-4174 | |
| 1321 | Predicting drug-target interactions using restricted Boltzmann machines. 2013 , 29, i126-34 | 133 |
| 1320 | Molecular characterization of an rsmD-like rRNA methyltransferase from the <i>Wolbachia</i> endosymbiont of <i>Brugia malayi</i> and antifilarial activity of specific inhibitors of the enzyme. 2013 , 57, 3843-56 | 10 |
| 1319 | Menthol inhibits 5-HT ₃ receptor-mediated currents. 2013 , 347, 398-409 | 35 |
| 1318 | Structural evidence for the involvement of the residues Ser187 and Tyr422 in substrate recognition in the 3-methylcrotonyl-coenzyme A carboxylase from <i>Pseudomonas aeruginosa</i> . 2013 , 154, 291-7 | 3 |
| 1317 | Crystal structure of the DdrB/ssDNA complex from <i>Deinococcus radiodurans</i> reveals a DNA binding surface involving higher-order oligomeric states. 2013 , 41, 9934-44 | 9 |
| 1316 | Tricyclic GyrB/ParE (TriBE) inhibitors: a new class of broad-spectrum dual-targeting antibacterial agents. 2013 , 8, e84409 | 71 |
| 1315 | Stabilization of the conductive conformation of a voltage-gated K ⁺ (Kv) channel: the lid mechanism. 2013 , 288, 16619-16628 | 5 |
| 1314 | Target prediction for an open access set of compounds active against <i>Mycobacterium tuberculosis</i> . 2013 , 9, e1003253 | 42 |
| 1313 | Has the bacterial biphenyl catabolic pathway evolved primarily to degrade biphenyl? The diphenylmethane case. 2013 , 195, 3563-74 | 13 |
| 1312 | The structure of substrate-free 1,5-anhydro-D-fructose reductase from <i>Sinorhizobium meliloti</i> 1021 reveals an open enzyme conformation. 2013 , 69, 844-9 | 7 |
| 1311 | Cloud infrastructures for in silico drug discovery: economic and practical aspects. 2013 , 2013, 138012 | 18 |
| 1310 | 1-aryl-3-[4-(thieno[3,2-d]pyrimidin-4-yloxy)phenyl]ureas as VEGFR-2 tyrosine kinase inhibitors: synthesis, biological evaluation, and molecular modelling studies. 2013 , 2013, 154856 | 3 |
| 1309 | Frataxin mRNA isoforms in FRDA patients and normal subjects: effect of tocotrienol supplementation. 2013 , 2013, 276808 | 12 |
| 1308 | LigandBox: A database for 3D structures of chemical compounds. 2013 , 9, 113-21 | 20 |

| | | |
|------|---|----|
| 1307 | Potent anti-diabetic effects of MHY908, a newly synthesized PPAR α dual agonist in db/db mice. 2013 , 8, e78815 | 22 |
| 1306 | Structure of the type VI effector-immunity complex (Tae4-Tai4) provides novel insights into the inhibition mechanism of the effector by its immunity protein. 2013 , 288, 5928-39 | 46 |
| 1305 | Disruption of E627 and R683 interaction is responsible for B-cell acute lymphoblastic leukemia caused by JAK2 R683G(S) mutations. 2013 , 54, 2693-700 | 3 |
| 1304 | Retinoid binding to human serum albumin. 2013 , 39, 294-303 | 29 |
| 1303 | PELE web server: atomistic study of biomolecular systems at your fingertips. 2013 , 41, W322-8 | 59 |
| 1302 | 9-Fluorenone-2-Carboxylic Acid as a Scaffold for Tubulin Interacting Compounds. 2013 , 78, 663-669 | 6 |
| 1301 | In silico optimization of a fragment-based hit yields biologically active, high-efficiency inhibitors for glutamate racemase. 2013 , 8, 1681-9 | 9 |
| 1300 | Flexible Refinement of Protein-Ligand Docking on Manifolds. 2013 , 1392-1397 | 2 |
| 1299 | N-substituted piperazinopyridylsteroid derivatives as abiraterone analogues inhibit growth and induce pro-apoptosis in human hormone-independent prostate cancer cell lines. 2013 , 82, 620-9 | 11 |
| 1298 | Chemical, pharmacological, and in vitro metabolic stability studies on enantiomerically pure RC-33 compounds: promising neuroprotective agents acting as μ receptor agonists. 2013 , 8, 1514-27 | 34 |
| 1297 | A bioinformatics pipeline for sequence to structure: A case study with a Cml patient undergoing treatment with imatinib. 2013 , | |
| 1296 | Improved flexible refinement of protein docking in CAPRI rounds 22-27. 2013 , 81, 2129-36 | 14 |
| 1295 | Structure and function of nucleoside hydrolases from <i>Physcomitrella patens</i> and maize catalyzing the hydrolysis of purine, pyrimidine, and cytokinin ribosides. 2013 , 163, 1568-83 | 25 |
| 1294 | . 2013 , | |
| 1293 | C-terminus of ETA/ETB receptors regulate endothelin-1 signal transmission. 2013 , 19, 257-62 | 8 |
| 1292 | Putative cholesterol-binding sites in human immunodeficiency virus (HIV) coreceptors CXCR4 and CCR5. 2013 , 81, 555-67 | 18 |
| 1291 | Identification of novel benzimidazole derivatives as anti- <i>Trypanosoma cruzi</i> agents: solid-phase synthesis, structure-activity relationships and molecular docking studies. 2013 , 5, 1719-32 | 7 |
| 1290 | Structural analysis and insights into the glycon specificity of the rice GH1 Os7BGlu26 α -D-mannosidase. 2013 , 69, 2124-35 | 9 |

| | | |
|------|--|-----|
| 1289 | Self-adaptive differential evolution algorithm incorporating local search for protein-ligand docking. 2013 , 410, 012030 | 3 |
| 1288 | Biochemical characterization of IMP-30, a metallo- β -lactamase with enhanced activity toward ceftazidime. 2013 , 57, 5122-6 | 9 |
| 1287 | Rottlerin-induced autophagy leads to the apoptosis in breast cancer stem cells: molecular mechanisms. 2013 , 12, 171 | 99 |
| 1286 | PfRIO-2 Kinase is a Potential Therapeutic Target of Antimalarial Protein Kinase Inhibitors. 2013 , 10, 85-91 | 1 |
| 1285 | Modulation of photophysics and pKa shift of the anti-cancer drug camptothecin in the nanocavities of supramolecular hosts. 2013 , 14, 532-42 | 39 |
| 1284 | Crystal structure of SsfS6, the putative C-glycosyltransferase involved in SF2575 biosynthesis. 2013 , 81, 1277-82 | 23 |
| 1283 | A New Binding Site Involving the C-terminal Domain to Design Specific Inhibitors of PepX. 2013 , 20, 45-53 | 1 |
| 1282 | A proposal of potent inhibitor for cancer metastasis blocking the pocket of urokinase receptor: ab initio molecular simulations. 2013 , 433, 012034 | 0 |
| 1281 | Swarm intelligence for molecular docking. 2013 , 18, 357 | 1 |
| 1280 | Comparing the suitability of autodock, gold and glide for the docking and predicting the possible targets of Ru(II)-based complexes as anticancer agents. 2013 , 18, 3760-78 | 42 |
| 1279 | An insight into the anticancer activities of Ru(II)-based metallocomounds using docking methods. 2013 , 18, 10829-56 | 5 |
| 1278 | Studies on Cross-linking of succinic acid with chitosan/collagen. 2013 , 16, 755-765 | 59 |
| 1277 | Design of multiligand inhibitors for the swine flu H1N1 neuraminidase binding site. 2013 , 6, 47-53 | 6 |
| 1276 | Structural elucidation of unique inhibitory activities of two thiazolo[4,5-d]pyrimidines against epidermal growth factor receptor (EGFR): implications for successful drug design. 2014 , 10, 46-58 | 6 |
| 1275 | Drug metabolite generation using a laboratory evolved NADPH independent cytochrome P450: application of in vitro and in silico approaches. 2013 , 7, 68-77 | |
| 1274 | Discovery of potential cholesterol esterase inhibitors using in silico docking studies. 2013 , 8, | 4 |
| 1273 | 2,2'-diphenyl-3,3'-diindolylmethane: a potent compound induces apoptosis in breast cancer cells by inhibiting EGFR pathway. 2013 , 8, e59798 | 27 |
| 1272 | Use of natural products as chemical library for drug discovery and network pharmacology. 2013 , 8, e62839 | 224 |

| | | |
|------|--|----|
| 1271 | Elucidation of the binding mechanism of coumarin derivatives with human serum albumin. 2013 , 8, e63805 | 46 |
| 1270 | Physicochemical characterization of a thermostable alcohol dehydrogenase from <i>Pyrobaculum aerophilum</i> . 2013 , 8, e63828 | 3 |
| 1269 | Cancer associated E17K mutation causes rapid conformational drift in AKT1 pleckstrin homology (PH) domain. 2013 , 8, e64364 | 41 |
| 1268 | Menthol binding and inhibition of α -nicotinic acetylcholine receptors. 2013 , 8, e67674 | 56 |
| 1267 | Identification of potential Plk1 targets in a cell-cycle specific proteome through structural dynamics of kinase and Polo box-mediated interactions. 2013 , 8, e70843 | 16 |
| 1266 | Ansamitocin P3 depolymerizes microtubules and induces apoptosis by binding to tubulin at the vinblastine site. 2013 , 8, e75182 | 29 |
| 1265 | Structure-activity relationships of FMRF-NH ₂ peptides demonstrate A role for the conserved C terminus and unique N-terminal extension in modulating cardiac contractility. 2013 , 8, e75502 | 12 |
| 1264 | Ligand pose and orientational sampling in molecular docking. 2013 , 8, e75992 | 83 |
| 1263 | Norisoboldine suppresses VEGF-induced endothelial cell migration via the cAMP-PKA-NF- κ B/Notch1 pathway. 2013 , 8, e81220 | 19 |
| 1262 | Structural insights into the mechanism for recognizing substrate of the cytochrome P450 enzyme TxtE. 2013 , 8, e81526 | 16 |
| 1261 | Exploration of virtual candidates for human HMG-CoA reductase inhibitors using pharmacophore modeling and molecular dynamics simulations. 2013 , 8, e83496 | 12 |
| 1260 | Antiviral potential of 4-hydroxypanduratin A, secondary metabolite of Fingerroot, <i>Boesenbergia pandurata</i> (Schult.), towards Japanese Encephalitis virus NS2B/NS3 protease. 2013 , 9, 54-60 | 14 |
| 1259 | Tetrodotoxin blockade on canine cardiac L-type Ca ²⁺ channels depends on pH and redox potential. 2013 , 11, 2140-53 | 9 |
| 1258 | Computational medicinal chemistry for rational drug design: Identification of novel chemical structures with potential anti-tuberculosis activity. 2014 , 14, 176-88 | 14 |
| 1257 | Structure-activity relationships of 3,3'-phenylmethylene-bis-4-hydroxycoumarins: selective and potent inhibitors of gram-positive bacteria. 2013 , 2013, 178649 | 7 |
| 1256 | Identification of PPAR α ligands with One-dimensional Drug Profile Matching. 2013 , 7, 917-28 | 4 |
| 1255 | Xanthene and xanthone derivatives as G-quadruplex stabilizing ligands. 2013 , 18, 13446-70 | 11 |
| 1254 | How computational studies of mosquito repellents contribute to the control of vector Borne Diseases. 2013 , 9, 300-7 | 6 |

| | | |
|------|--|----|
| 1253 | PfPRIO-2 Kinase is a Potential Therapeutic Target of Antimalarial Protein Kinase Inhibitors. 2013 , 10, 85-91 | 4 |
| 1252 | . 2013 , | |
| 1251 | In Silico Molecular Docking of Lavandula Angustifolia Mill [®] Compounds Along with a Number of Antianxiety Drugs with GABAA Receptor for Reduce Stress. 2014 , 30, 889-894 | |
| 1250 | istar: a web platform for large-scale protein-ligand docking. 2014 , 9, e85678 | 74 |
| 1249 | Anti-enterovirus 71 effects of chrysin and its phosphate ester. 2014 , 9, e89668 | 43 |
| 1248 | The curious case of benzbromarone: insight into super-inhibition of cytochrome P450. 2014 , 9, e89967 | 18 |
| 1247 | The bitter barricading of prostaglandin biosynthesis pathway: understanding the molecular mechanism of selective cyclooxygenase-2 inhibition by amarogentin, a secoiridoid glycoside from Swertia chirayita. 2014 , 9, e90637 | 14 |
| 1246 | Inhibition of Paracoccidioides lutzii Pb01 isocitrate lyase by the natural compound argenilactone and its semi-synthetic derivatives. 2014 , 9, e94832 | 18 |
| 1245 | Computational design of a pH stable enzyme: understanding molecular mechanism of penicillin acylase's adaptation to alkaline conditions. 2014 , 9, e100643 | 42 |
| 1244 | Rational modification of estrogen receptor by combination of computational and experimental analysis. 2014 , 9, e102658 | 8 |
| 1243 | In silico analysis of missense mutations in LPAR6 reveals abnormal phospholipid signaling pathway leading to hypotrichosis. 2014 , 9, e104756 | 18 |
| 1242 | The major cow milk allergen Bos d 5 manipulates T-helper cells depending on its load with siderophore-bound iron. 2014 , 9, e104803 | 37 |
| 1241 | Molecular investigations of protriptyline as a multi-target directed ligand in Alzheimer's disease. 2014 , 9, e105196 | 20 |
| 1240 | Metabolomics analysis and modeling suggest a lysophosphocholines-PAF receptor interaction in fibromyalgia. 2014 , 9, e107626 | 34 |
| 1239 | Search for α adrenergic receptor ligands by virtual screening via grid computing and investigation of binding modes by docking and molecular dynamics simulations. 2014 , 9, e107837 | 10 |
| 1238 | Break CDK2/Cyclin E1 interface allosterically with small peptides. 2014 , 9, e109154 | 15 |
| 1237 | Sequential application of ligand and structure based modeling approaches to index chemicals for their hH4R antagonism. 2014 , 9, e109340 | 16 |
| 1236 | Operating mechanism and molecular dynamics of pheromone-binding protein ASP1 as influenced by pH. 2014 , 9, e110565 | 5 |

| | | |
|------|---|-----|
| 1235 | Rational design of small-molecule stabilizers of spermine synthase dimer by virtual screening and free energy-based approach. 2014 , 9, e110884 | 19 |
| 1234 | In Silico Screening of Mutated K-Ras Inhibitors from Malaysian Typhonium flagelliforme for Non-Small Cell Lung Cancer. 2014 , 2014, 431696 | 4 |
| 1233 | Evaluation of 11 scoring functions performance on matrix metalloproteinases. 2014 , 2014, 162150 | 8 |
| 1232 | Prediction of epitope-based peptides for the utility of vaccine development from fusion and glycoprotein of nipah virus using in silico approach. 2014 , 2014, 402492 | 31 |
| 1231 | Glycyrrhetic acid and E.resveratrolsoid act as potential plant derived compounds against dopamine receptor D3 for Parkinson's disease: a pharmacoinformatics study. 2015 , 9, 187-98 | 17 |
| 1230 | In silico discovery of potential VEGFR-2 inhibitors from natural derivatives for anti-angiogenesis therapy. 2014 , 15, 15994-6011 | 35 |
| 1229 | Structure-based virtual screening for drug discovery: principles, applications and recent advances. 2014 , 14, 1923-38 | 481 |
| 1228 | Bis-indole derivatives with antitumor activity turn out to be specific ligands of human telomeric G-quadruplex. 2014 , 2, 54 | 20 |
| 1227 | Multipose binding in molecular docking. 2014 , 15, 2622-45 | 38 |
| 1226 | Inhibition of GlcNAc-processing glycosidases by C-6-azido-NAG-thiazoline and its derivatives. 2014 , 19, 3471-88 | 11 |
| 1225 | Computational redesign of bacterial biotin carboxylase inhibitors using structure-based virtual screening of combinatorial libraries. 2014 , 19, 4021-45 | 8 |
| 1224 | Conformational characterization of ipomotaosides and their recognition by COX-1 and 2. 2014 , 19, 5421-33 | 3 |
| 1223 | In silico docking, molecular dynamics and binding energy insights into the bolinaquinone-clathrin terminal domain binding site. 2014 , 19, 6609-22 | 27 |
| 1222 | Flavonoids with M1 muscarinic acetylcholine receptor binding activity. 2014 , 19, 8933-48 | 15 |
| 1221 | Potential mechanism of action of meso-dihydroguaiaretic acid on Mycobacterium tuberculosis H37Rv. 2014 , 19, 20170-82 | 13 |
| 1220 | Use of Bioinformatics Tools in Different Spheres of Life Sciences. 2014 , 05, | 7 |
| 1219 | A Review of Computational Tools for Designing Drugs Used by General Practitioners. 2014 , 02, | |
| 1218 | GABAA receptor binding molecules from traditional Chinese Medicine: An in silico approach. 2014 , 9, | |

| | | |
|------|--|-----|
| 1217 | In silico targeting of interleukin-6 by natural compounds. 2014 , 9, | 2 |
| 1216 | Receptor Chemoprint Derived Pharmacophore Model for Development of CAIX Inhibitors. 2014 , s8, | 2 |
| 1215 | Molecular docking studies of anti-cancerous candidates in Hippophae rhamnoides and Hippophae salicifolia. 2014 , 28, 406-15 | 22 |
| 1214 | QSAR-based models for designing quinazoline/imidazothiazoles/pyrazolopyrimidines based inhibitors against wild and mutant EGFR. 2014 , 9, e101079 | 27 |
| 1213 | Theoretical Research into Anticancer Activity of Diterpenes Isolated from the Paraiban Flora. 2014 , 9, 1934578X1400900 | |
| 1212 | Molecular dynamics simulations reveal initial structural and dynamic features for the A2AR as a result of ligand binding. 2014 , 40, 996-1014 | 4 |
| 1211 | 4-Amino-2-arylamino-6-(2,6-dichlorophenyl)-pyrido[2,3-d]pyrimidin-7-(8H)-ones as BCR kinase inhibitors for B lymphoid malignancies. 2014 , 86, 664-75 | 17 |
| 1210 | p63 threonine phosphorylation signals the interaction with the WW domain of the E3 ligase Itch. 2014 , 13, 3207-17 | 9 |
| 1209 | NADH-dependent lactate dehydrogenase from Alcaligenes eutrophus H16 reduces 2-oxoadipate to 2-hydroxyadipate. 2014 , 19, 1048-1057 | 3 |
| 1208 | Structural requirements for imidazo[1,2-a]pyrazine derivatives as Aurora A kinase inhibitors and validation of the model. 2014 , 23, 5215-5223 | 5 |
| 1207 | SwissTargetPrediction: a web server for target prediction of bioactive small molecules. 2014 , 42, W32-8 | 557 |
| 1206 | Molecular cloning and in-silico characterization of high temperature stress responsive pAPX gene isolated from heat tolerant Indian wheat cv. Raj 3765. 2014 , 7, 713 | 12 |
| 1205 | Residue Leu940 has a crucial role in the linkage and reaction specificity of the glucansucrase GTF180 of the probiotic bacterium Lactobacillus reuteri 180. 2014 , 289, 32773-82 | 28 |
| 1204 | Evidence of ternary complex formation in Trypanosoma cruzi trans-sialidase catalysis. 2014 , 289, 423-36 | 13 |
| 1203 | Molecular dynamics simulations to explore the active/inactive conformers of guinea pig β 2 adrenoceptor for the selective design of agonists or antagonists. 2014 , 40, 1244-1254 | 4 |
| 1202 | Overexpression of the pp32r1 (ANP32C) oncogene or its functional mutant pp32r1Y140H confers enhanced resistance to FTY720 (Fingolimod). 2014 , 15, 289-96 | 7 |
| 1201 | In silico analysis of different generation β -lactams antibiotics with penicillin binding protein-2 of Neisseria meningitidis for curing meningococcal disease. 2014 , 6, 259-70 | 2 |
| 1200 | Targeting Btk/Etk of prostate cancer cells by a novel dual inhibitor. 2014 , 5, e1409 | 29 |

| | | |
|------|--|----|
| 1199 | Design and synthesis of nonsteroidal progesterone receptor antagonists based on C,C'-diphenylcarborane scaffold as a hydrophobic pharmacophore. 2014 , 84, 264-77 | 28 |
| 1198 | Analysis of hepatitis B virus drug-resistant mutation (M204V) using molecular dynamics simulation techniques. 2014 , 69, 1445-1452 | 1 |
| 1197 | Novel computational study on π -stacking to understand mechanistic interactions of Tryptanthrin analogues with DNA. 2014 , 24, 73-9 | 6 |
| 1196 | A computational model for non-conserved mature miRNAs from the rice genome. 2014 , 25, 205-20 | 3 |
| 1195 | In silico study of binding motifs in squalene synthase enzyme of secondary metabolic pathway of solanaceae [corrected]. 2014 , 41, 7201-8 | 5 |
| 1194 | Theoretical evaluation and improvement on the potency of the rhodanine-based inhibitors for human serotonin N-acetyltransferase. 2014 , 40, 1201-1208 | 1 |
| 1193 | The interaction pattern between a homology model of 40S ribosomal S9 protein of Rhizoctonia solani and 1-hydroxyphenazine by docking study. 2014 , 2014, 682946 | 6 |
| 1192 | Inhibition of Enterovirus 71 replication by 7-hydroxyflavone and diisopropyl-flavon7-yl Phosphate. 2014 , 9, e92565 | 10 |
| 1191 | Cellular target recognition of perfluoroalkyl acids: in vitro evaluation of inhibitory effects on lysine decarboxylase. 2014 , 496, 381-388 | 4 |
| 1190 | Molecular tetris. 2014 , | 0 |
| 1189 | Photophysics of a solvent sensitive keto-tetrahydrocarbazole based fluorophore and its interaction with triethylamine: A spectroscopic inquest under surfactant and β -CD confinement. 2014 , 1074, 617-628 | 5 |
| 1188 | The structural basis of ATP as an allosteric modulator. 2014 , 10, e1003831 | 59 |
| 1187 | Integrating in silico prediction methods, molecular docking, and molecular dynamics simulation to predict the impact of ALK missense mutations in structural perspective. 2014 , 2014, 895831 | 29 |
| 1186 | Common recognition principles across diverse sequence and structural families of sialic acid binding proteins. 2014 , 24, 5-16 | 5 |
| 1185 | Elucidating the specificity of non-heparin-based conformational activators of antithrombin for factor Xa inhibition. 2014 , 5, 36-42 | 8 |
| 1184 | An infrastructure to mine molecular descriptors for ligand selection on virtual screening. 2014 , 2014, 325959 | |
| 1183 | Maximum Entropy in Drug Discovery. 2014 , 16, 3754-3768 | 6 |
| 1182 | Predicting the Origins of Anti-Blood Group Antibody Specificity: A Case Study of the ABO A- and B-Antigens. 2014 , 5, 397 | 8 |

| | | |
|------|---|----|
| 1181 | Differentiated adaptive evolution, episodic relaxation of selective constraints, and pseudogenization of umami and sweet taste genes TAS1Rs in catarrhine primates. 2014 , 11, 79 | 10 |
| 1180 | Phytochemicals increase the antibacterial activity of antibiotics by acting on a drug efflux pump. 2014 , 3, 885-96 | 51 |
| 1179 | The MID-PIWI module of Piwi proteins specifies nucleotide- and strand-biases of piRNAs. 2014 , 20, 773-81 | 59 |
| 1178 | A Cloud-Based Data Farming Platform for Molecular Dynamics Simulations. 2014 , | 6 |
| 1177 | Structural insights into the catalytic mechanism of <i>Synechocystis</i> magnesium protoporphyrin IX O-methyltransferase (ChLM). 2014 , 289, 25690-8 | 16 |
| 1176 | Anticonvulsant effects of bis-1,4-dihydropyridines and the probable role of L-type calcium channels suggested by docking simulations. 2014 , 23, 5149-5159 | 6 |
| 1175 | Protein-ligand structure guided by backbone and side-chain proton chemical shift perturbations. 2014 , 60, 147-56 | 14 |
| 1174 | Understanding the effects on constitutive activation and drug binding of a D130N mutation in the β_2 adrenergic receptor via molecular dynamics simulation. 2014 , 20, 2491 | 2 |
| 1173 | In silico screening for inhibitors of p-glycoprotein that target the nucleotide binding domains. 2014 , 86, 716-26 | 17 |
| 1172 | Comparative kinetics of Qi site inhibitors of cytochrome bc1 complex: picomolar antimycin and micromolar cyazofamid. 2014 , 83, 71-80 | 20 |
| 1171 | Catalytic activity of human placental alkaline phosphatase (PLAP): insights from a computational study. 2014 , 118, 14302-13 | 6 |
| 1170 | A zebrafish in vivo phenotypic assay to identify 3-aminothiophene-2-carboxylic acid-based angiogenesis inhibitors. 2014 , 12, 527-35 | 14 |
| 1169 | Crystal structure of DszC from <i>Rhodococcus</i> sp. XP at 1.79 Å 2014 , 82, 1708-20 | 20 |
| 1168 | Rational design of small molecule inhibitors targeting the Ras GEF, SOS1. 2014 , 21, 1618-28 | 37 |
| 1167 | Design and synthesis of imidazo[2,1-b]thiazole-chalcone conjugates: microtubule-destabilizing agents. 2014 , 9, 2766-80 | 26 |
| 1166 | Standards-based metadata management for molecular simulations. 2014 , 26, 1744-1759 | 15 |
| 1165 | <i>Anadenanthera colubrina</i> (Vell.) Brenan produces steroidal substances that are active against <i>Alternaria alternata</i> (Fr.) Keissler and that may bind to oxysterol-binding proteins. 2014 , 70, 1815-22 | 9 |
| 1164 | Alkyl chain substituted 1,9-pyrazoloanthrones exhibit prominent inhibitory effect on c-Jun N-terminal kinase (JNK). 2014 , 12, 4656-62 | 5 |

| | | |
|------|--|----|
| 1163 | Mutational analysis of the quorum-sensing receptor LasR reveals interactions that govern activation and inhibition by nonlactone ligands. 2014 , 21, 1361-1369 | 31 |
| 1162 | HIV-1 group O integrase displays lower enzymatic efficiency and higher susceptibility to raltegravir than HIV-1 group M subtype B integrase. 2014 , 58, 7141-50 | 6 |
| 1161 | Structure of pneumococcal peptidoglycan hydrolase LytB reveals insights into the bacterial cell wall remodeling and pathogenesis. 2014 , 289, 23403-16 | 37 |
| 1160 | HLA-B*13:01 is associated with salazosulfapyridine-induced drug rash with eosinophilia and systemic symptoms in Chinese Han population. 2014 , 15, 1461-9 | 30 |
| 1159 | Structure-based discovery of the first allosteric inhibitors of cyclin-dependent kinase 2. 2014 , 13, 2296-305 | 41 |
| 1158 | Structure-activity relationships of peptidomimetics that inhibit PPI of HER2-HER3. 2014 , 101, 693-702 | 17 |
| 1157 | Oriental switching of protein conformation as a function of nanoparticle curvature and their geometrical fitting. 2014 , 141, 084707 | 17 |
| 1156 | Structures of lipoyl synthase reveal a compact active site for controlling sequential sulfur insertion reactions. 2014 , 464, 123-33 | 57 |
| 1155 | Crystal structure of a bacterial unsaturated glucuronyl hydrolase with specificity for heparin. 2014 , 289, 4787-97 | 6 |
| 1154 | A review of metabolic and enzymatic engineering strategies for designing and optimizing performance of microbial cell factories. 2014 , 11, 91-9 | 42 |
| 1153 | Genetic encoding of caged cysteine and caged homocysteine in bacterial and mammalian cells. 2014 , 15, 1793-9 | 43 |
| 1152 | Toward on-the-fly quantum mechanical/molecular mechanical (QM/MM) docking: development and benchmark of a scoring function. 2014 , 54, 3137-52 | 51 |
| 1151 | Identification of toxin inhibitors using a magnetic nanosensor-based assay. 2014 , 10, 1202-11 | 5 |
| 1150 | Targeting the hydrophobic pocket of autotaxin with virtual screening of inhibitors identifies a common aromatic sulfonamide structural motif. 2014 , 281, 1017-28 | 17 |
| 1149 | Novel mutations in CYP51B from <i>Penicillium digitatum</i> involved in prochloraz resistance. 2014 , 52, 762-70 | 30 |
| 1148 | Mechanistic insight into ligand binding to G-quadruplex DNA. 2014 , 42, 5447-55 | 65 |
| 1147 | Molecular insight into the role of the N-terminal extension in the maturation, substrate recognition, and catalysis of a bacterial alginate lyase from polysaccharide lyase family 18. 2014 , 289, 29558-69 | 37 |
| 1146 | Optimizing the underlying parameters for protein-nanoparticle interaction: advancement in theoretical simulation. 2014 , 3, | 7 |

| | | |
|------|--|-----|
| 1145 | Integration of binding peptide selection and multifunctional particles as tool-box for capture of soluble proteins in serum. 2014 , 11, | 13 |
| 1144 | Switches of hydrogen bonds during ligand-protein association processes determine binding kinetics. 2014 , 27, 537-48 | 14 |
| 1143 | Binding of polarity-sensitive hydrophobic ligands to erythroid and nonerythroid spectrin: fluorescence and molecular modeling studies. 2014 , 32, 852-65 | 12 |
| 1142 | Protocol for rational design of covalently interacting inhibitors. 2014 , 15, 3226-35 | 16 |
| 1141 | A PIK3C3-ankyrin-B-dynactin pathway promotes axonal growth and multiorganelle transport. 2014 , 207, 735-52 | 58 |
| 1140 | Practical Chemoinformatics. 2014 , | 10 |
| 1139 | Exploring the free-energy landscape of carbohydrate-protein complexes: development and validation of scoring functions considering the binding-site topology. 2014 , 28, 1191-204 | 1 |
| 1138 | Mathematical and computational modeling in biology at multiple scales. 2014 , 11, 52 | 7 |
| 1137 | Some gating potentiators, including VX-770, diminish B508-CFTR functional expression. 2014 , 6, 246ra97 | 223 |
| 1136 | An in silico structure-based approach to anti-infective drug discovery. 2014 , 141, 17-27 | 2 |
| 1135 | 2- and 3-substituted imidazo[1,2-a]pyrazines as inhibitors of bacterial type IV secretion. 2014 , 22, 6459-70 | 22 |
| 1134 | Crystal structure of tRNA m1G9 methyltransferase Trm10: insight into the catalytic mechanism and recognition of tRNA substrate. 2014 , 42, 509-25 | 31 |
| 1133 | Design and synthesis of chalcone derivatives as inhibitors of the ferredoxin - ferredoxin-NADP+ reductase interaction of Plasmodium falciparum: pursuing new antimalarial agents. 2014 , 19, 21473-88 | 27 |
| 1132 | A quantitative chemical proteomics approach to profile the specific cellular targets of andrographolide, a promising anticancer agent that suppresses tumor metastasis. 2014 , 13, 876-86 | 72 |
| 1131 | Differential coupling of KLF10 to Sin3-HDAC and PCAF regulates the inducibility of the FOXP3 gene. 2014 , 307, R608-20 | 20 |
| 1130 | Isoxazolotacrine as non-toxic and selective butyrylcholinesterase inhibitors for Alzheimer's disease. 2014 , 6, 1883-91 | 9 |
| 1129 | Molecular docking studies of flavonoids for their inhibition pattern against β -catenin and pharmacophore model generation from experimentally known flavonoids to fabricate more potent inhibitors for Wnt signaling pathway. 2014 , 10, S264-71 | 17 |
| 1128 | Resistance mechanisms of cancer cells to the novel vacuolar H(+)-ATPase inhibitor archazolid B. 2014 , 32, 893-903 | 8 |

| | | |
|------|--|----|
| 1127 | Exploring Large Scale Receptor-Ligand Pairs in Molecular Docking Workflows in HPC Clouds. 2014 , | 5 |
| 1126 | Identification of potential herbal inhibitor of acetylcholinesterase associated Alzheimer's disorders using molecular docking and molecular dynamics simulation. 2014 , 2014, 705451 | 17 |
| 1125 | Exploring the molecular mechanisms of electron shuttling across the microbe/metal space. 2014 , 5, 318 | 48 |
| 1124 | Does the cis/trans configuration of peptide bonds in bioactive tripeptides play a role in ACE-1 enzyme inhibition?. 2014 , 8, 59-65 | 13 |
| 1123 | Reduced ribosomes of the apicoplast and mitochondrion of Plasmodium spp. and predicted interactions with antibiotics. 2014 , 4, 140045 | 26 |
| 1122 | Structure and properties of complexes of Ecthyotrypsin with hydroxyl-containing gemini dicationic surfactants with a spacer moiety of varying length. 2014 , 55, 1556-1564 | 2 |
| 1121 | MORT: a powerful foundational library for computational biology and CADD. 2014 , 6, | 78 |
| 1120 | Novel 9-(alkylthio)-Acenaphtho[1,2-e]-1,2,4-triazine derivatives: synthesis, cytotoxic activity and molecular docking studies on B-cell lymphoma 2 (Bcl-2). 2014 , 22, 2 | 12 |
| 1119 | Structural and biochemical basis for the inhibition of cell death by APIP, a methionine salvage enzyme. 2014 , 111, E54-61 | 23 |
| 1118 | Functional architecture of MFS D-glucose transporters. 2014 , 111, E719-27 | 67 |
| 1117 | Structure determination of human Fas apoptosis inhibitory molecule and identification of the critical residues linking the interdomain interaction to the anti-apoptotic activity. 2014 , 70, 1812-22 | 5 |
| 1116 | Crystal structures and catalytic mechanism of the C-methyltransferase Coq5 provide insights into a key step of the yeast coenzyme Q synthesis pathway. 2014 , 70, 2085-92 | 15 |
| 1115 | Structural and functional characterization of an arylamine N-acetyltransferase from the pathogen Mycobacterium abscessus: differences from other mycobacterial isoforms and implications for selective inhibition. 2014 , 70, 3066-79 | 6 |
| 1114 | Structure of 4-pyridoxolactonase from Mesorhizobium loti. 2014 , 70, 424-32 | |
| 1113 | Hierarchical MapReduce: towards simplified cross-domain data processing. 2014 , 26, 878-893 | 3 |
| 1112 | Integrated approach to structure-based enzymatic drug design: molecular modeling, spectroscopy, and experimental bioactivity. 2014 , 114, 493-537 | 81 |
| 1111 | Identification and characterization of a new reversible MAGL inhibitor. 2014 , 22, 3285-91 | 36 |
| 1110 | Fullerene derivatives as a new class of inhibitors of protein tyrosine phosphatases. 2014 , 24, 3175-9 | 22 |

| | | |
|------|--|-----|
| 1109 | Synthesis and characterization of 1H-phenanthro[9,10-d]imidazole derivatives as multifunctional agents for treatment of Alzheimer's disease. 2014 , 1840, 2886-903 | 20 |
| 1108 | Study of the binding interaction between fluorinated matrix metalloproteinase inhibitors and Human Serum Albumin. 2014 , 79, 13-23 | 7 |
| 1107 | Expression, characterization and homology modeling of a novel eukaryotic GH84 β -N-acetylglucosaminidase from <i>Penicillium chrysogenum</i> . 2014 , 95, 204-10 | 14 |
| 1106 | Nitensidine A, a guanidine alkaloid from <i>Pterogyne nitens</i> , is a novel substrate for human ABC transporter ABCB1. 2014 , 21, 323-32 | 28 |
| 1105 | Exploring unsymmetrical dyads as efficient inhibitors against the insect β -N-acetyl-D-hexosaminidase OfHex2. 2014 , 97, 152-62 | 8 |
| 1104 | Unraveling the binding mechanism of asiatic acid with human serum albumin and its biological implications. 2014 , 32, 1290-302 | 34 |
| 1103 | Advances in Computational Biology. 2014 , | |
| 1102 | In silico study on Penicillin derivatives and Cephalosporins for upper respiratory tract bacterial pathogens. 2014 , 4, 241-251 | 4 |
| 1101 | 3-Hydroxy-4-pyridinone derivatives as metal ion and amyloid binding agents. 2014 , 6, 249-62 | 17 |
| 1100 | New chemotypes as <i>Trypanosoma cruzi</i> triosephosphate isomerase inhibitors: a deeper insight into the mechanism of inhibition. 2014 , 29, 198-204 | 17 |
| 1099 | Analysis of essential amino acid residues for catalytic activity of cis-epoxysuccinate hydrolase from <i>Bordetella</i> sp. BK-52. 2014 , 98, 1641-9 | 6 |
| 1098 | Integrating molecular docking, CoMFA analysis, and machine-learning classification with virtual screening toward identification of novel scaffolds as <i>Plasmodium falciparum</i> enoyl acyl carrier protein reductase inhibitor. 2014 , 23, 3308-3326 | 5 |
| 1097 | Targeting VEGFR1 on endothelial progenitors modulates their differentiation potential. 2014 , 17, 603-16 | 11 |
| 1096 | In silico study on indole derivatives as anti HIV-1 agents: a combined docking, molecular dynamics and 3D-QSAR study. 2014 , 37, 1001-15 | 14 |
| 1095 | Homology modeling of lanosterol 14 α -demethylase of <i>Candida albicans</i> and insights into azole binding. 2014 , 23, 2890-2899 | 6 |
| 1094 | Lassomycin, a ribosomally synthesized cyclic peptide, kills mycobacterium tuberculosis by targeting the ATP-dependent protease ClpC1P1P2. 2014 , 21, 509-518 | 255 |
| 1093 | Molecular Cloning, Modelling and Docking with Curcumin of the Dengue Virus 2 NS5 Polymerase Domain. 2014 , 273-278 | 2 |
| 1092 | Protein Conformational Dynamics. 2014 , | 9 |

| | | |
|------|--|-----|
| 1091 | Targeting of Smoothed for therapeutic gain. 2014 , 35, 237-46 | 81 |
| 1090 | Synthesis and biological evaluation of imidazo[2,1-b][1,3,4]thiadiazole-linked oxindoles as potent tubulin polymerization inhibitors. 2014 , 9, 1463-75 | 14 |
| 1089 | Use of Freely Available and Open Source Tools for In Silico Screening in Chemical Biology. 2014 , 91, 602-604 | 11 |
| 1088 | Genome of the human hookworm <i>Necator americanus</i> . 2014 , 46, 261-269 | 139 |
| 1087 | Chemical inhibition of prometastatic lysyl-tRNA synthetase-laminin receptor interaction. 2014 , 10, 29-34 | 42 |
| 1086 | Anti- <i>Trypanosoma cruzi</i> and anti-leishmanial activity by quinoxaline-7-carboxylate 1,4-di-N-oxide derivatives. 2014 , 113, 2027-35 | 30 |
| 1085 | Mn(2+) and Mg(2+) synergistically enhanced lactic acid production by <i>Lactobacillus rhamnosus</i> FTDC 8313 via affecting different stages of the hexose monophosphate pathway. 2014 , 116, 644-53 | 8 |
| 1084 | Virtual screening with AutoDock Vina and the common pharmacophore engine of a low diversity library of fragments and hits against the three allosteric sites of HIV integrase: participation in the SAMPL4 protein-ligand binding challenge. 2014 , 28, 429-441 | 38 |
| 1083 | A novel allosteric mechanism in the cysteine peptidase cathepsin K discovered by computational methods. 2014 , 5, 3287 | 59 |
| 1082 | Computational approaches and resources in single amino acid substitutions analysis toward clinical research. 2014 , 94, 365-423 | 19 |
| 1081 | Structural Genomics and Drug Discovery. 2014 , | 3 |
| 1080 | Structural and energetic insights into the selective interactions of monoacylglycerol lipase with its natural substrate and small-molecule inhibitors. 2014 , 23, 2391-2404 | 8 |
| 1079 | Molecular docking and 3D-QSAR studies on the MAPKAP-K2 inhibitors. 2014 , 23, 2252-2263 | 7 |
| 1078 | iview: an interactive WebGL visualizer for protein-ligand complex. 2014 , 15, 56 | 42 |
| 1077 | Engineering of <i>Thermomyces lanuginosus</i> lipase Lip: creation of novel biocatalyst for efficient biosynthesis of chiral intermediate of Pregabalin. 2014 , 98, 2473-83 | 23 |
| 1076 | Structure and dynamics studies of sterol 24-C-methyltransferase with mechanism based inactivators for the disruption of ergosterol biosynthesis. 2014 , 41, 4279-93 | 14 |
| 1075 | Cloning, expression, and characterization of the β -glucosidase hydrolyzing secoisolariciresinol diglucoside to secoisolariciresinol from <i>Bacteroides uniformis</i> ZL1. 2014 , 98, 2519-31 | 18 |
| 1074 | Could glutaric acid (GA) replace glutaraldehyde in the preparation of biocompatible biopolymers with high mechanical and thermal properties?. 2014 , 126, 127-140 | 23 |

| | | |
|------|--|----|
| 1073 | Virtual screening of potential inhibitors from TCM for the CPSF30 binding site on the NS1A protein of influenza A virus. 2014 , 20, 2142 | 12 |
| 1072 | Insights into the influence of 5-HT _{2c} aminoacidic variants with the inhibitory action of serotonin inverse agonists and antagonists. 2014 , 20, 2120 | 14 |
| 1071 | Interaction of phospholipase A of the E. coli outer membrane with the inhibitors of eucaryotic phospholipases A and their effect on the Ca ²⁺ -induced permeabilization of the bacterial membrane. 2014 , 247, 281-8 | 3 |
| 1070 | Molecular modeling study on the structural basis of binding mechanism of C6-substituted phthalides with monoamine oxidases. 2014 , 23, 3624-3631 | |
| 1069 | Structural evaluation of an alternative Protein A biomimetic ligand for antibody purification. 2014 , 28, 25-34 | 13 |
| 1068 | Insights into the binding of GABA to the insect RDL receptor from atomistic simulations: a comparison of models. 2014 , 28, 35-48 | 14 |
| 1067 | Probing the binding of Syzygium-derived α-glucosidase inhibitors with N- and C-terminal human maltase glucoamylase by docking and molecular dynamics simulation. 2014 , 172, 102-14 | 9 |
| 1066 | Virtual screening of integrase inhibitors by large scale binding free energy calculations: the SAMPL4 challenge. 2014 , 28, 475-90 | 45 |
| 1065 | Directed evolution of nitrobenzene dioxygenase for the synthesis of the antioxidant hydroxytyrosol. 2014 , 98, 4975-85 | 15 |
| 1064 | Novel structural hybrids of pyrazolobenzothiazines with benzimidazoles as cholinesterase inhibitors. 2014 , 78, 106-17 | 27 |
| 1063 | Naphthoquinone-mediated inhibition of lysine acetyltransferase KAT3B/p300, basis for non-toxic inhibitor synthesis. 2014 , 289, 7702-17 | 26 |
| 1062 | Interaction of anticancer reduced Schiff base coumarin derivatives with human serum albumin investigated by fluorescence quenching and molecular modeling. 2014 , 52, 16-23 | 40 |
| 1061 | Toward drug repurposing in epigenetics: olsalazine as a hypomethylating compound active in a cellular context. 2014 , 9, 560-5 | 60 |
| 1060 | Synthesis and evaluation of (-)-Massoialactone and analogues as potential anticancer and anti-inflammatory agents. 2014 , 76, 291-300 | 26 |
| 1059 | Molecular modeling of 5HT _{2A} receptor - arylpiperazine ligands interactions. 2014 , 83, 462-71 | 9 |
| 1058 | Decreasing acidity in a series of aldose reductase inhibitors: 2-Fluoro-4-(1H-pyrrol-1-yl)phenol as a scaffold for improved membrane permeation. 2014 , 22, 2194-207 | 15 |
| 1057 | Beyond topoisomerase inhibition: antitumor 1,4-naphthoquinones as potential inhibitors of human monoamine oxidase. 2014 , 83, 401-10 | 18 |
| 1056 | Regioselective Enzymatic Halogenation of Substituted Tryptophan Derivatives using the FAD-Dependent Halogenase RebH. 2014 , 6, n/a-n/a | 25 |

| | | |
|------|--|-----|
| 1055 | Structural insights into the MDP binding and CARD-CARD interaction in zebrafish (<i>Danio rerio</i>) NOD2: a molecular dynamics approach. 2014 , 27, 260-75 | 35 |
| 1054 | Oncogenic protein interfaces: small molecules, big challenges. 2014 , 14, 248-62 | 196 |
| 1053 | Synthesis, pharmacological evaluation, and μ receptor interaction analysis of hydroxyethyl substituted piperazines. 2014 , 57, 2884-94 | 13 |
| 1052 | Homology and molecular dynamics models of toll-like receptor 7 protein and its dimerization. 2014 , 83, 656-65 | 10 |
| 1051 | Enantioselective synthesis of benzomorphan analogues by intramolecular oxa-Pictet-Spengler cyclization. 2014 , 25, 489-496 | 7 |
| 1050 | Sulfur rich 2-mercaptobenzothiazole and 1,2,3-triazole conjugates as novel antitubercular agents. 2014 , 76, 274-83 | 65 |
| 1049 | Peptide inhibitors against dengue virus infection. 2014 , 84, 148-57 | 27 |
| 1048 | Synthesis and structure of a new mononuclear copper(II) complex with 2,2'-bipyridine and picrate: molecular docking, DNA-binding, and in vitro anticancer activity. 2014 , 67, 630-648 | 24 |
| 1047 | A molecular model to explain the controlled release from SBA-15 functionalized with APTES. 2014 , 195, 43-49 | 30 |
| 1046 | Molecular Characterization and In Silico Analysis of the Pheromone-Binding Protein of the European Grapevine Moth <i>Lobesia botrana</i> (Denis & Schiffermüller) (Lepidoptera, Tortricidae). 2014 , 43, 266-75 | 5 |
| 1045 | Enabling Technologies to Facilitate Natural Product-Based Drug Discovery from African Biodiversity. 2014 , 57-67 | |
| 1044 | Small-molecule probes targeting the viral PPxY-host Nedd4 interface block egress of a broad range of RNA viruses. 2014 , 88, 7294-306 | 77 |
| 1043 | 2-[N-Alkyl(R-phenyl)-aminomethyl]-3-phenyl-7-trifluoromethylquinoxalines as anticancer agents inhibitors of folate enzymes. 2014 , 75, 169-83 | 7 |
| 1042 | Multivariate analysis in the identification of biological targets for designed molecular structures: the BIOTA protocol. 2014 , 75, 106-10 | 18 |
| 1041 | Molecular Dynamics simulations of Inhibitor of Apoptosis Proteins and identification of potential small molecule inhibitors. 2014 , 24, 2098-104 | 6 |
| 1040 | Synthesis of 2-{2-[(1-naphthalen-1-ylsulfonyl)amino]-1,3-thiazol-4-yl} acetamides with 11 β -hydroxysteroid dehydrogenase inhibition and in combo antidiabetic activities. 2014 , 74, 179-86 | 17 |
| 1039 | Discovery of novel anti-leishmanial agents targeting LdLip3 lipase. 2014 , 49, 68-79 | 6 |
| 1038 | Stable polyglutamine dimers can contain β -hairpins with interdigitated side chains-but not β -helices, β -nanotubes, β -pseudohelices, or steric zippers. 2014 , 106, 1721-8 | 6 |

| | | |
|------|---|--------|
| 1037 | 5,6,7,9-Tetrahydro-[1,3]dioxolo[4,5-h]carbazol-8-one: A solvatochromic PET-acceptor fluorescent probe. 2014 , 153, 296-303 | 4 |
| 1036 | In silico approaches to predict the potential of milk protein-derived peptides as dipeptidyl peptidase IV (DPP-IV) inhibitors. 2014 , 57, 43-51 | 90 |
| 1035 | In-silico modelling and identification of a possible inhibitor of H1N1 virus. 2014 , 4, S467-S476 | 4 |
| 1034 | Structural modifications of 4-aryl-4-oxo-2-aminybutanamides and their acetyl- and butyrylcholinesterase inhibitory activity. Investigation of AChE-ligand interactions by docking calculations and molecular dynamics simulations. 2014 , 81, 158-75 | 17 |
| 1033 | Protein-drug interactome analysis of SSRI-mediated neurorecovery following stroke. 2014 , 120, 1-9 | 13 |
| 1032 | Structural elucidation of specific noncovalent association of folic acid with native cyclodextrins using an ion mobility mass spectrometry and theoretical approach. 2014 , 86, 4249-55 | 21 |
| 1031 | Genetically induced dysfunctions of Kir2.1 channels: implications for short QT3 syndrome and autism-epilepsy phenotype. 2014 , 23, 4875-86 | 52 |
| 1030 | Synthesis and biological evaluation of benzo[b]furans as inhibitors of tubulin polymerization and inducers of apoptosis. 2014 , 9, 117-28 | 28 |
| 1029 | Structure modeling and hybrid virtual screening study of Alzheimer's associated protease kallikrein 8 for the identification of novel inhibitors. 2014 , 23, 3516-3527 | 3 |
| 1028 | Cheminformatic and Chemogenomic Approach to ADMET. 2014 , 125-143 | |
| 1027 | Imidazolylchromanones containing non-benzylic oxime ethers: synthesis and molecular modeling study of newazole antifungals selective against <i>Cryptococcus gattii</i> . 2014 , 76, 264-73 | 15 |
| 1026 | Modeling in vitro inhibition of butyrylcholinesterase using molecular docking, multi-linear regression and artificial neural network approaches. 2014 , 22, 538-49 | 21 |
| 1025 | Bifunctional ADP-dependent phosphofructokinase/glucokinase activity in the order Methanococcales--biochemical characterization of the mesophilic enzyme from <i>Methanococcus maripaludis</i> . 2014 , 281, 2017-29 | 15 |
| 1024 | Carbonic anhydrase inhibition by 1-aryl-3-(4-aminosulfonylphenyl)thioureas. 2014 , 29, 901-5 | 13 |
| 1023 | Studies of (-)-pironetin binding to tubulin: conformation, docking, and molecular dynamics. 2014 , 79, 3752-64 | 18 |
| 1022 | Improvement of virtual screening results by docking data feature analysis. 2014 , 54, 1401-11 | 21 |
| 1021 | Importance of ligand conformational energies in carbohydrate docking: Sorting the wheat from the chaff. <i>Journal of Computational Chemistry</i> , 2014 , 35, 526-39 | 3.5 57 |
| 1020 | Molecular dynamics simulations reveal a novel mechanism for ATP inhibition of insulin degrading enzyme. 2014 , 54, 1380-90 | 10 |

| | | |
|------|--|----|
| 1019 | Recent advances in engineering proteins for biocatalysis. 2014 , 111, 1273-87 | 71 |
| 1018 | LIMD2 is a small LIM-only protein overexpressed in metastatic lesions that regulates cell motility and tumor progression by directly binding to and activating the integrin-linked kinase. 2014 , 74, 1390-1403 | 19 |
| 1017 | Docking and Pharmacophore Modelling for Virtual Screening. 2014 , 195-269 | 1 |
| 1016 | Structure-activity relationships and molecular modelling of new 5-arylidene-4-thiazolidinone derivatives as aldose reductase inhibitors and potential anti-inflammatory agents. 2014 , 81, 1-14 | 53 |
| 1015 | Hsp90 inhibitors, part 2: combining ligand-based and structure-based approaches for virtual screening application. 2014 , 54, 970-7 | 21 |
| 1014 | Novel and versatile methodology for synthesis of α -aryl- β -mercapto ketone derivatives as potential urease inhibitors. 2014 , 11, 1113-1119 | 8 |
| 1013 | Electrospray ionization mass spectrometry probing of binding affinity of berbamine, a flexible cyclic alkaloid from traditional Chinese medicine, with G-quadruplex DNA. 2014 , 28, 143-7 | 19 |
| 1012 | jMetalCpp: optimizing molecular docking problems with a C++ metaheuristic framework. 2014 , 30, 437-8 | 24 |
| 1011 | Molecular modeling evidence for His438 flip in the mechanism of butyrylcholinesterase hysteretic behavior. 2014 , 52, 434-45 | 13 |
| 1010 | Ligand binding mode prediction by docking: mdm2/mdmx inhibitors as a case study. 2014 , 54, 648-59 | 30 |
| 1009 | Insight into the binding modes and inhibition mechanisms of adamantyl-based 1,3-disubstituted urea inhibitors in the active site of the human soluble epoxide hydrolase. 2014 , 32, 1231-47 | 2 |
| 1008 | Unraveling the mode of binding of the anticancer drug topotecan with dsDNA. 2014 , 4, 1015-1024 | 20 |
| 1007 | The versatile nature of the 6-aminoquinolone scaffold: identification of submicromolar hepatitis C virus NS5B inhibitors. 2014 , 57, 1952-63 | 35 |
| 1006 | Synthesis and crystal structure of a ternary copper(II) complex of 2,2'-bipyridine and picrate: Molecular docking, reactivity towards DNA and in vitro anticancer activity. 2014 , 1058, 97-105 | 13 |
| 1005 | A conformational analysis of mouse Nalp3 domain structures by molecular dynamics simulations, and binding site analysis. 2014 , 10, 1104-16 | 22 |
| 1004 | BP-Dock: a flexible docking scheme for exploring protein-ligand interactions based on unbound structures. 2014 , 54, 913-25 | 30 |
| 1003 | Preliminary studies on trigonelline as potential anti-Alzheimer disease agent: determination by hydrophilic interaction liquid chromatography and modeling of interactions with beta-amyloid. 2014 , 968, 101-4 | 22 |
| 1002 | Multi-spectroscopic and molecular modeling studies on the interaction of antihypertensive drug; methyldopa with calf thymus DNA. 2014 , 10, 338-47 | 88 |

| | | |
|------|---|-----|
| 1001 | Homology modeling of T. cruzi and L. major NADH-dependent fumarate reductases: ligand docking, molecular dynamics validation, and insights on their binding modes. 2014 , 48, 47-59 | 13 |
| 1000 | Docking studies in target proteins involved in antibacterial action mechanisms: extending the knowledge on standard antibiotics to antimicrobial mushroom compounds. 2014 , 19, 1672-84 | 25 |
| 999 | LIBSA--a method for the determination of ligand-binding preference to allosteric sites on receptor ensembles. 2014 , 54, 530-8 | 12 |
| 998 | Virtual screening of the SAMPL4 blinded HIV integrase inhibitors dataset. 2014 , 28, 455-62 | 11 |
| 997 | Methods to enable the design of bioactive small molecules targeting RNA. 2014 , 12, 1029-39 | 52 |
| 996 | Receptor-ligand molecular docking. 2014 , 6, 75-87 | 218 |
| 995 | Exploration of the binding mode between (-)-zampanolide and tubulin using docking and molecular dynamics simulation. 2014 , 20, 2070 | 13 |
| 994 | Binding and molecular dynamics studies of 7-hydroxycoumarin derivatives with human serum albumin and its pharmacological importance. 2014 , 11, 1117-31 | 90 |
| 993 | Rational design of a structural framework with potential use to develop chemical reagents that target and modulate multiple facets of Alzheimer's disease. 2014 , 136, 299-310 | 142 |
| 992 | 3D-QSAR and docking studies of piperidine carboxamide derivatives as ALK inhibitors. 2014 , 23, 2576-2583 | 6 |
| 991 | Descriptors requirement for QSAR analysis of pyrazolo-triazolo-pyrimidine derivative as human A3 receptor antagonists: design of novel furan derivatives and validation by docking. 2014 , 23, 2554-2563 | 1 |
| 990 | Related genes and potential biomarkers for early diagnosis of Alzheimer's disease: a preliminary study based on DNA microarray. 2014 , 29, 90-5 | 3 |
| 989 | DiSCuS: an open platform for (not only) virtual screening results management. 2014 , 54, 347-54 | 5 |
| 988 | In situ and in silico evaluation of amine- and folate-terminated dendrimers as nanocarriers of anesthetics. 2014 , 73, 250-7 | 15 |
| 987 | A systematic study of chemogenomics of carbohydrates. 2014 , 10, 391-7 | 8 |
| 986 | A QM/MM study of the catalytic mechanism of nicotinamidase. 2014 , 12, 1265-77 | 10 |
| 985 | Effect of oxindolimine copper(II) and zinc(II) complexes on human topoisomerase I activity. 2014 , 6, 117-25 | 33 |
| 984 | Spectroscopic and theoretical investigation of oxali-palladium interactions with β -lactoglobulin. 2014 , 118, 1038-46 | 95 |

| | | |
|-----|---|-----|
| 983 | Design, synthesis and characterization of novel inhibitors against mycobacterial β -ketoacyl CoA reductase FabG4. 2014 , 12, 73-85 | 11 |
| 982 | iSyn: WebGL-Based Interactive De Novo Drug Design. 2014 , | 4 |
| 981 | An introduction to biomolecular simulations and docking. 2014 , 40, 732-764 | 18 |
| 980 | Genetically encoded optochemical probes for simultaneous fluorescence reporting and light activation of protein function with two-photon excitation. 2014 , 136, 15551-8 | 102 |
| 979 | Repurposing of Kinase Inhibitors to Target c-Abl as Potential Therapeutics for Alzheimer's Disease. 2014 , 9, 331-340 | 5 |
| 978 | Synthesis of ethyl (R)-4-cyano-3-hydroxybutyrate in high concentration using a novel halohydrin dehalogenase HHDH-PL from <i>Parvibaculum lavamentivorans</i> DS-1. 2014 , 4, 64027-64031 | 16 |
| 977 | Ion mobility and Top-down MS complementary approaches for the structural analysis of protein models bound to anticancer metallodrugs. 2014 , 423, 60-69 | 5 |
| 976 | Computational approach to understanding the mechanism of action of isoniazid, an anti-TB drug. 2014 , 3, 276-82 | 32 |
| 975 | Protein Ligand Docking in Drug Discovery. 2014 , 249-286 | 5 |
| 974 | In vitro inhibition of lysine decarboxylase activity by organophosphate esters. 2014 , 92, 506-16 | 16 |
| 973 | Molecular interaction of 2-mercaptobenzimidazole with catalase reveals a potentially toxic mechanism of the inhibitor. 2014 , 141, 241-6 | 8 |
| 972 | VEGFR tyrosine kinase inhibitor II (VRI) induced vascular insufficiency in zebrafish as a model for studying vascular toxicity and vascular preservation. 2014 , 280, 408-20 | 33 |
| 971 | Computational Docking as a Tool for the Rational Design of Carbohydrate-Based Drugs. 2014 , 53-72 | 3 |
| 970 | Role of 3D Structures in Understanding, Predicting, and Designing Molecular Interactions in the Chemokine Receptor Family. 2014 , 41-85 | 1 |
| 969 | Synthesis, characterization, antibacterial, DNA binding and cleavage studies of mixed ligand Cu(II), Co(II) complexes. 2014 , 24, 1687-99 | 8 |
| 968 | A β -1-6/ β -1-3 galactosidase from <i>Bifidobacterium animalis</i> subsp. <i>lactis</i> BI-04 gives insight into sub-specificities of β -galactoside catabolism within <i>Bifidobacterium</i> . 2014 , 94, 1024 | 26 |
| 967 | Spectroscopic and molecular modeling studies of the interaction between morin and polyamidoamine dendrimer. 2014 , 29, 573-8 | 1 |
| 966 | Understanding the guanidine-like cationic moiety for optimal binding into the DNA minor groove. 2014 , 9, 2065-73 | 13 |

| | | |
|-----|--|----|
| 965 | Withanone-rich combination of Ashwagandha withanolides restricts metastasis and angiogenesis through hnRNP-K. 2014 , 13, 2930-40 | 45 |
| 964 | Computational library design for increasing haloalkane dehalogenase stability. 2014 , 15, 1660-72 | 52 |
| 963 | Synthesis of 2-anilinopyridine dimers as microtubule targeting and apoptosis inducing agents. 2014 , 22, 6755-67 | 10 |
| 962 | Structure and catalytic activity of β -chymotrypsin in solutions of gemini surfactants. 2014 , 63, 273-279 | 5 |
| 961 | Docking-based 3D-QSAR study of pyridyl aminothiazole derivatives as checkpoint kinase 1 inhibitors. 2014 , 25, 651-71 | 11 |
| 960 | Interactions of the aquated forms of ruthenium(III) anticancer drugs with protein: a detailed molecular docking and QM/MM investigation. 2014 , 4, 60548-60556 | 9 |
| 959 | Molecular Modeling and Evaluation of Novel Dibenzopyrrole Derivatives as Telomerase Inhibitors and Potential Drug for Cancer Therapy. 2014 , 11, 1196-207 | 8 |
| 958 | Binding interaction of a newly developed bisindole drug molecule with β -cyclodextrin: face to face shielding of indole hoops. 2014 , 4, 38206 | 1 |
| 957 | A spectroscopic investigation and molecular docking study on the interaction of hen egg white lysozyme with liposomes of saturated and unsaturated phosphocholines probed by an anticancer drug ellipticine. 2014 , 16, 5368-81 | 43 |
| 956 | Development of regioselective deacylation of peracetylated β -D-monosaccharides using lipase from <i>Pseudomonas stutzeri</i> under sustainable conditions. 2014 , 4, 55495-55502 | 13 |
| 955 | Rational design and synthesis of novel 2-(substituted-2H-chromen-3-yl)-5-aryl-1H-imidazole derivatives as an anti-angiogenesis and anti-cancer agent. 2014 , 4, 56489-56501 | 17 |
| 954 | Application of molecular docking and ONIOM methods for the description of interactions between anti-quorum sensing active (AHL) analogues and the <i>Pseudomonas aeruginosa</i> LasR binding site. 2014 , 10, 1162-71 | 19 |
| 953 | Femtosecond to nanosecond dynamics of 2,2'-bipyridine-3,3'-diol inside the nano-cavities of molecular containers. 2014 , 16, 933-9 | 15 |
| 952 | Sequence context induced antimicrobial activity: insight into lipopolysaccharide permeabilization. 2014 , 10, 1596-612 | 27 |
| 951 | Docking and molecular dynamics studies of the binding between Peloruside A and tubulin. 2014 , 29, 702-9 | 3 |
| 950 | Growth arrest by the antitumor steroidal lactone withaferin A in human breast cancer cells is associated with down-regulation and covalent binding at cysteine 303 of β -tubulin. 2014 , 289, 1852-65 | 81 |
| 949 | β -Hydroxyethyl piperidine iminosugar and N-alkylated derivatives: a study of their activity as glycosidase inhibitors and as immunosuppressive agents. 2014 , 22, 5776-82 | 11 |
| 948 | Active compounds from a diverse library of triazolothiadiazole and triazolothiadiazine scaffolds: synthesis, crystal structure determination, cytotoxicity, cholinesterase inhibitory activity, and binding mode analysis. 2014 , 22, 6163-73 | 46 |

| | | |
|-----|--|-----|
| 947 | Elucidation of mechanisms of interaction of a multifunctional peptide Pa-MAP with lipid membranes. 2014 , 1838, 2899-909 | 9 |
| 946 | HLA-DQA1-HLA-DRB1 variants confer susceptibility to pancreatitis induced by thiopurine immunosuppressants. 2014 , 46, 1131-4 | 130 |
| 945 | Development of p-carborane-based nonsteroidal progesterone receptor antagonists. 2014 , 22, 5329-37 | 15 |
| 944 | Palladium-catalyzed direct addition of arylboronic acids to 2-aminobenzonitrile derivatives: synthesis, biological evaluation and in silico analysis of 2-aminobenzophenones, 7-benzoyl-2-oxindolines, and 7-benzoylindoles. 2014 , 12, 8204-11 | 27 |
| 943 | Lead Discovery and Lead Modification. 2014 , 19-122 | 5 |
| 942 | Systems pharmacology strategies for anticancer drug discovery based on natural products. 2014 , 10, 1912-7 | 36 |
| 941 | Microwave-assisted synthesis of arene ruthenium(II) complexes [(RCH) ₃ Ru(m-MOPIP)Cl]Cl (R = -H and -CH ₃) as groove binder to c-myc G4 DNA. 2014 , 43, 9216-25 | 26 |
| 940 | Discovery of pentacyclic triterpenoids as potential entry inhibitors of influenza viruses. 2014 , 57, 10058-71 | 77 |
| 939 | Effects of bisphenol S on the structures and activities of trypsin and pepsin. 2014 , 62, 11303-11 | 36 |
| 938 | Discovery of a benzofuran derivative (MBPTA) as a novel ROCK inhibitor that protects against MPP ⁺ -induced oxidative stress and cell death in SH-SY5Y cells. 2014 , 74, 283-93 | 27 |
| 937 | Molecular mechanism of viral resistance to a potent non-nucleoside inhibitor unveiled by molecular simulations. 2014 , 53, 6941-53 | 28 |
| 936 | Design of glycosyltransferase inhibitors targeting human O-GlcNAc transferase (OGT). 2014 , 5, 1172-1178 | 15 |
| 935 | Protein binding for detection of small changes on a nanoparticle surface. 2014 , 139, 1364-71 | 10 |
| 934 | Ligand induced change of α_2 adrenergic receptor from active to inactive conformation and its implication for the closed/open state of the water channel: insight from molecular dynamics simulation, free energy calculation and Markov state model analysis. 2014 , 16, 15874-85 | 33 |
| 933 | Validation of a computational docking methodology to identify the non-covalent binding site of ligands to DNA. 2014 , 10, 2106-25 | 11 |
| 932 | Extensive consensus docking evaluation for ligand pose prediction and virtual screening studies. 2014 , 54, 2980-6 | 66 |
| 931 | Preferential DNA photocleavage potency of Zn(II) over Ni(II) derivatives of carboxymethyl tetracationic porphyrin: the role of the mode of binding to DNA. 2014 , 43, 545-54 | 12 |
| 930 | DNA Recognition with Polycyclic-Aromatic-Hydrocarbon-Presenting Calixarene Conjugates. 2014 , 2014, 7605-7613 | 12 |

| | | |
|-----|---|-----|
| 929 | Potential anti-bacterial agents: montmorillonite clay-catalyzed synthesis of novel 2-(3,5-substituted-1H-pyrazol-1-yl)-3-substituted quinolines and their in silico molecular docking studies. 2014 , 4, 58011-58018 | 3 |
| 928 | Rice cytochrome P450 MAX1 homologs catalyze distinct steps in strigolactone biosynthesis. 2014 , 10, 1028-33 | 230 |
| 927 | Structural and functional investigation of zebrafish (<i>Danio rerio</i>) NOD1 leucine rich repeat domain and its interaction with iE-DAP. 2014 , 10, 2942-53 | 22 |
| 926 | Acetylcholine promotes binding of α -conotoxin MII at $\alpha 5 \beta 2$ nicotinic acetylcholine receptors. 2014 , 15, 413-24 | 12 |
| 925 | Novel benzopsoresalen analogues: synthesis, biological activity and molecular docking studies. 2014 , 87, 298-305 | 6 |
| 924 | Stereoselective Enzyme Cascades: An Efficient Synthesis of Chiral β -Butyrolactones. 2014 , 4, 1321-1331 | 68 |
| 923 | Diaryl-substituted azolythioacetamides: Inhibitor discovery of New Delhi metallo- β -lactamase-1 (NDM-1). 2014 , 9, 2445-8 | 53 |
| 922 | Analyzing the interaction of a herbal compound Andrographolide from as a folklore against swine flu (H1N1). 2014 , 4, S624-S630 | 7 |
| 921 | Endogenous polyamines reduce the toxicity of soluble α -peptide aggregates associated with Alzheimer's disease. 2014 , 15, 1985-91 | 22 |
| 920 | Single water entropy: hydrophobic crossover and application to drug binding. 2014 , 118, 10553-64 | 19 |
| 919 | Celastrol inhibits Plasmodium falciparum enoyl-acyl carrier protein reductase. 2014 , 22, 6053-6061 | 13 |
| 918 | Structural basis of valmerins as dual inhibitors of GSK3 β /CDK5. 2014 , 20, 2407 | 20 |
| 917 | Parametrization of an Orbital-Based Linear-Scaling Quantum Force Field for Noncovalent Interactions. 2014 , 10, 1086-1098 | 27 |
| 916 | Implementation of the Hungarian algorithm to account for ligand symmetry and similarity in structure-based design. 2014 , 54, 518-29 | 49 |
| 915 | TRPV4 channel activity is modulated by direct interaction of the ankyrin domain to PI(4,5)P $_2$ 2014 , 5, 4994 | 69 |
| 914 | Synthesis and biological evaluation of tetrahydro[1,4]diazepino[1,2-a]indol-1-ones as cyclin-dependent kinase inhibitors. 2014 , 83, 617-29 | 11 |
| 913 | Structure prediction of GPCRs using piecewise homologs and application to the human CCR5 chemokine receptor: validation through agonist and antagonist docking. 2014 , 32, 1274-89 | 10 |
| 912 | Identification and characterization of a tyramine-glutamate ligase (MfnD) involved in methanofuran biosynthesis. 2014 , 53, 6220-30 | 13 |

| | | |
|-----|---|-----|
| 911 | Discovery of high affinity ligands for α_2 -adrenergic receptor through pharmacophore-based high-throughput virtual screening and docking. 2014 , 53, 148-160 | 5 |
| 910 | Role of the flavan-3-ol and galloyl moieties in the interaction of (-)-epigallocatechin gallate with serum albumin. 2014 , 62, 3768-75 | 58 |
| 909 | SERS and MD simulation studies of a kinase inhibitor demonstrate the emergence of a potential drug discovery tool. 2014 , 111, 10416-21 | 31 |
| 908 | Exploration of the antagonist CP-376395 escape pathway for the corticotropin-releasing factor receptor 1 by random acceleration molecular dynamics simulations. 2014 , 10, 1958-67 | 20 |
| 907 | Design and synthesis of a new dimeric xanthone derivative: enhancement of G-quadruplex selectivity and telomere damage. 2014 , 12, 9572-82 | 10 |
| 906 | Crenolanib is a selective type I pan-FLT3 inhibitor. 2014 , 111, 5319-24 | 155 |
| 905 | Design, synthesis and biological evaluation of 4-anilinothieno[2,3-d]pyrimidine-based hydroxamic acid derivatives as novel histone deacetylase inhibitors. 2014 , 22, 6146-55 | 18 |
| 904 | Synthesis of fused tetrahydropyrido[2,3-c]coumarin derivatives as potential inhibitors for dopamine d3 receptors, catalyzed by hydrated ferric sulfate. 2014 , 4, 3581-3590 | 10 |
| 903 | Synthesis, biological evaluation, and molecular modeling of new 3-(cyclopentyloxy)-4-methoxybenzaldehyde O-(2-(2,6-dimethylmorpholino)-2-oxoethyl) Oxime (GEBR-7b) related phosphodiesterase 4D (PDE4D) inhibitors. 2014 , 57, 7061-72 | 16 |
| 902 | Biosynthesis of the 5-(Aminomethyl)-3-furanmethanol moiety of methanofuran. 2014 , 53, 4635-47 | 12 |
| 901 | Cytotoxicity of the bisphenolic honokiol from <i>Magnolia officinalis</i> against multiple drug-resistant tumor cells as determined by pharmacogenomics and molecular docking. 2014 , 21, 1525-33 | 19 |
| 900 | Bet v 1 from birch pollen is a lipocalin-like protein acting as allergen only when devoid of iron by promoting Th2 lymphocytes. 2014 , 289, 17416-21 | 47 |
| 899 | Relationship between a point mutation S97C in CK1 β protein and its affect on ATP-binding affinity. 2014 , 32, 394-405 | 19 |
| 898 | Assessing the performance of MM/PBSA and MM/GBSA methods. 5. Improved docking performance using high solute dielectric constant MM/GBSA and MM/PBSA rescoring. 2014 , 16, 22035-45 | 322 |
| 897 | Interference of boswellic acids with the ligand binding domain of the glucocorticoid receptor. 2014 , 54, 978-86 | 13 |
| 896 | Structure-based drug design of diphenyl β -aminoalkylphosphonates as prostate-specific antigen antagonists. 2014 , 54, 2967-79 | 4 |
| 895 | Mutagenesis studies of the 14 β -internal cavity of histone deacetylase 1: insights toward the acetate-escape hypothesis and selective inhibitor design. 2014 , 57, 642-50 | 34 |
| 894 | Synthesis and MRSA PK inhibitory activity of thiazole containing deoxytospentin analogues. 2014 , 70, 7845-7853 | 13 |

| | | |
|-----|---|-----|
| 893 | Synthesis of pyrazolo[4,3-a]phenanthridines, a new scaffold for Pim kinase inhibition. 2014 , 22, 4704-10 | 11 |
| 892 | Rationally designed less toxic SPD-304 analogs and preliminary evaluation of their TNF inhibitory effects. 2014 , 347, 798-805 | 17 |
| 891 | Echinomycin, a potential binder of FKBP12, shows minor effect on calcineurin activity. 2014 , 19, 1275-81 | 5 |
| 890 | Evidence for the involvement of descending pain-inhibitory mechanisms in the attenuation of cancer pain by carvacrol aided through a docking study. 2014 , 116, 8-15 | 24 |
| 889 | Therapeutic potential of targeting the oncogenic SHP2 phosphatase. 2014 , 57, 6594-609 | 96 |
| 888 | Docking ligands into flexible and solvated macromolecules. 7. Impact of protein flexibility and water molecules on docking-based virtual screening accuracy. 2014 , 54, 3198-210 | 23 |
| 887 | Synthesis of imidazothiadiazole-Benzimidazole conjugates as mitochondrial apoptosis inducers. 2014 , 5, 1644-1650 | 11 |
| 886 | Development of quinone analogues as dynamin GTPase inhibitors. 2014 , 85, 191-206 | 15 |
| 885 | Synthesis, biological evaluation and molecular docking of novel chalcone-coumarin hybrids as anticancer and antimalarial agents. 2014 , 85, 65-76 | 139 |
| 884 | Identification of Small Inhibitory Molecules Targeting the Bfl-1 Anti-Apoptotic Protein That Alleviates Resistance to ABT-737. 2014 , 19, 1035-46 | 10 |
| 883 | Structural insights into the substrate-binding mechanism for a novel chitosanase. 2014 , 461, 335-45 | 41 |
| 882 | Relative motions between left flipper and dorsal fin domains favour P2X4 receptor activation. 2014 , 5, 4189 | 29 |
| 881 | Asymmetric phase-transfer catalysis with homo- and heterochiral quaternary ammonium salts: a theoretical study. 2014 , 118, 5154-67 | 19 |
| 880 | The cloud and other new computational methods to improve molecular modelling. 2014 , 9, 1121-31 | 11 |
| 879 | Binding studies of L-3,4-dihydroxyphenylalanine with human serum albumin. 2014 , 10, 3101-10 | 37 |
| 878 | Structure and function of human DnaJ homologue subfamily a member 1 (DNAJA1) and its relationship to pancreatic cancer. 2014 , 53, 1360-72 | 35 |
| 877 | A dynamic niching genetic algorithm strategy for docking highly flexible ligands. 2014 , 289, 206-224 | 83 |
| 876 | Identification of borrelidin binding site on threonyl-tRNA synthetase. 2014 , 451, 485-90 | 5 |

| | | |
|-----|--|----|
| 875 | Targeting the Tcf4 G13ANDE17 binding site to selectively disrupt β -catenin/T-cell factor protein-protein interactions. 2014 , 9, 193-201 | 39 |
| 874 | Modeling the coverage of an AFM tip by enzymes and its application in nanobiosensors. 2014 , 53, 100-104 | 18 |
| 873 | Emergence of pyridoxal phosphorylation through a promiscuous ancestor during the evolution of hydroxymethyl pyrimidine kinases. 2014 , 588, 3068-73 | 7 |
| 872 | Molecular docking and molecular dynamics studies on the structure-activity relationship of fluoroquinolone for the HERG channel. 2014 , 10, 2863-9 | 10 |
| 871 | Benzothiazolyl substituted iminothiazolidinones and benzamido-oxothiazolidines as potent and partly selective aldose reductase inhibitors. 2014 , 5, 1371-1380 | 15 |
| 870 | Computational investigation of cholesterol binding sites on mitochondrial VDAC. 2014 , 118, 9852-60 | 38 |
| 869 | Study on the interaction between bovine serum albumin and 4'-azido-2'-deoxyfluoroarabincytidine or analogs by spectroscopy and molecular modeling. 2014 , 132, 786-94 | 10 |
| 868 | Proposed interaction of some novel antidepressant pyrazolines against monoamine oxidase isoforms. Molecular docking studies and PASS assisted in silico approach. 2014 , 4, 297-301 | 6 |
| 867 | Drug-target interaction prediction via chemogenomic space: learning-based methods. 2014 , 10, 1273-87 | 60 |
| 866 | Isomannide-based peptidomimetics as inhibitors for human tissue kallikreins 5 and 7. 2014 , 5, 128-32 | 26 |
| 865 | Structural and dynamics analysis of matrix metalloproteinases MMP-2 complexed with chemically modified tetracyclines (CMTs). 2014 , 32, 1907-18 | 5 |
| 864 | Endocrine disruptome--an open source prediction tool for assessing endocrine disruption potential through nuclear receptor binding. 2014 , 54, 1254-67 | 73 |
| 863 | Synthesis of novel palladium(II) complexes with oxalic acid diamide derivatives and their interaction with nucleosides and proteins. Structural, solution, and computational study. 2014 , 43, 15126-37 | 22 |
| 862 | Design and synthesis of aminostilbene-arylpropenones as tubulin polymerization inhibitors. 2014 , 9, 2565-79 | 16 |
| 861 | Synthesis, antitumor activity, and structure-activity relationship study of trihydroxylated 2,4,6-triphenyl pyridines as potent and selective topoisomerase II inhibitors. 2014 , 84, 555-65 | 29 |
| 860 | Simulating the catalytic effect of a designed mononuclear zinc metalloenzyme that catalyzes the hydrolysis of phosphate triesters. 2014 , 118, 12146-52 | 6 |
| 859 | Design, synthesis, and preliminary bioactivity studies of substituted purine hydroxamic acid derivatives as novel histone deacetylase (HDAC) inhibitors. 2014 , 5, 1887-1891 | 8 |
| 858 | Morphological effect of gold nanoparticles on the adsorption of bovine serum albumin. 2014 , 16, 20471-82 | 48 |

- 857 Potent and selective inhibitors of class A β -lactamase: 7-prenyloxy coumarins. **2014**, 67, 373-7 10
- 856 New N-1,N-10-bridged pyrrolo[2,3-a]carbazole-3-carbaldehydes: synthesis and biological activities. **2014**, 57, 108-115 12
- 855 New pyrrole derivatives with potent tubulin polymerization inhibiting activity as anticancer agents including hedgehog-dependent cancer. **2014**, 57, 6531-52 57
- 854 Synthesis, characterization, investigation of biological activity and theoretical studies of hydrazone compounds containing chloroacetyl group. **2014**, 1075, 566-578 9
- 853 Ab initio molecular simulations for proposing potent inhibitors to butyrylcholinesterases. **2014**, 54, 54-61 7
- 852 Antipsychotic haloperidol binding to the human dopamine D3 receptor: beyond docking through QM/MM refinement toward the design of improved schizophrenia medicines. **2014**, 5, 1041-54 30
- 851 Computer based screening for novel inhibitors against *Vibrio cholerae* using NCI diversity set-II: an alternative approach by targeting transcriptional activator ToxT. **2014**, 6, 108-17 6
- 850 Molecular docking of bacosides with tryptophan hydroxylase: a model to understand the bacosides mechanism. **2014**, 4, 251-5 7
- 849 Differences in conformational dynamics between *Plasmodium falciparum* and human Hsp90 orthologues enable the structure-based discovery of pathogen-selective inhibitors. **2014**, 57, 2524-35 32
- 848 Construction of a highly stable artificial glutathione peroxidase on a protein nanoring. **2014**, 12, 362-9 18
- 847 Pyridine-3-carboxamide-6-yl-ureas as novel inhibitors of bacterial DNA gyrase: structure based design, synthesis, SAR and antimicrobial activity. **2014**, 86, 31-8 39
- 846 2-Aminothiazole derivatives as antimycobacterial agents: Synthesis, characterization, in vitro and in silico studies. **2014**, 87, 643-56 31
- 845 An in vitro study reveals the nutraceutical potential of punicic acid relevant to diabetes via enhanced GLUT4 expression and adiponectin secretion. **2014**, 5, 2590-601 15
- 844 Identification of novel potential antibiotics against *Staphylococcus* using structure-based drug screening targeting dihydrofolate reductase. **2014**, 54, 1242-53 18
- 843 Further Development of the FFT-based Method for Atomistic Modeling of Protein Folding and Binding under Crowding: Optimization of Accuracy and Speed. **2014**, 10, 2824-2835 21
- 842 Computational evaluation of the dynamic fluctuations of peripheral loops enclosing the catalytic tunnel of a family 7 cellobiohydrolase. **2014**, 118, 5340-9 7
- 841 Engineering the biosynthesis of novel rhamnolipids in *Escherichia coli* for enhanced oil recovery. **2014**, 117, 139-50 25
- 840 Binding of flavonoids to staphylococcal enterotoxin B. **2014**, 74, 1-8 9

| | | |
|-----|--|----|
| 839 | Switching a newly discovered lactonase into an efficient and thermostable phosphotriesterase by simple double mutations His250Ile/Ile263Trp. 2014 , 111, 1920-30 | 24 |
| 838 | Biological evaluation of potent triclosan-derived inhibitors of the enoyl-acyl carrier protein reductase InhA in drug-sensitive and drug-resistant strains of <i>Mycobacterium tuberculosis</i> . 2014 , 9, 2528-37 | 20 |
| 837 | Design of a truncated cardiotoxin-I analogue with potent insulinotropic activity. 2014 , 57, 2623-33 | 8 |
| 836 | Study of the structural and dynamic effects in the FimH adhesin upon α -D-heptyl mannose binding. 2014 , 57, 1416-27 | 38 |
| 835 | Stereoselective inclusion mechanism of ketoprofen into β -cyclodextrin: insights from molecular dynamics simulations and free energy calculations. 2014 , 133, 1 | 10 |
| 834 | Chemo- and bioinformatics resources for in silico drug discovery from medicinal plants beyond their traditional use: a critical review. 2014 , 31, 1585-611 | 85 |
| 833 | Fluorescence quenching studies of β -butyrolactone binding protein (CprB) from <i>Streptomyces coelicolor</i> A3(2). 2014 , 118, 10035-42 | 16 |
| 832 | Cytotoxic activity assessment and c-Src tyrosine kinase docking simulation of thieno[2,3-b]pyridine-based derivatives. 2014 , 23, 1225-1233 | 6 |
| 831 | Novel non-cyclooxygenase inhibitory derivatives of naproxen for colorectal cancer chemoprevention. 2014 , 23, 4177-4188 | 13 |
| 830 | Molecular docking, QSAR and AMD-ET analysis of 6-methyl-1,3,8-trichlorodibenzofuran and its analogs against Estrogen receptor alpha. 2014 , 23, 4724-4748 | 6 |
| 829 | Size does matter! Label-free detection of small molecule-protein interaction. 2014 , 406, 4033-51 | 23 |
| 828 | Identification and functional characterization of sorbitol-6-phosphate dehydrogenase protein from rice and structural elucidation by in silico approach. 2014 , 240, 223-38 | 8 |
| 827 | Insights into the molecular interactions between aminopeptidase and amyloid beta peptide using molecular modeling techniques. 2014 , 46, 1853-66 | 27 |
| 826 | Ab initio molecular simulations for proposing novel peptide inhibitors blocking the ligand-binding pocket of urokinase receptor. 2014 , 20, 2292 | 2 |
| 825 | The ability of molecular docking to unravel the controversy and challenges related to P-glycoprotein--a well-known, yet poorly understood drug transporter. 2014 , 32, 618-25 | 47 |
| 824 | Anti-inflammatory potential of alpha-linolenic acid mediated through selective COX inhibition: computational and experimental data. 2014 , 37, 1297-306 | 33 |
| 823 | Site-Identification by Ligand Competitive Saturation (SILCS) assisted pharmacophore modeling. 2014 , 28, 491-507 | 46 |
| 822 | Simulated Interactions between Endothelin Converting Enzyme and A β Peptide: Insights into Subsite Recognition and Cleavage Mechanism. 2014 , 20, 409-420 | 13 |

| | | |
|-----|---|-----|
| 821 | Autogrid-based clustering of kinases: selection of representative conformations for docking purposes. 2014 , 18, 611-9 | 2 |
| 820 | Adenosine deaminase activity modulation by some street drug: molecular docking simulation and experimental investigation. 2014 , 22, 42 | 5 |
| 819 | Rehabilitating drug-induced long-QT promoters: in-silico design of hERG-neutral cisapride analogues with retained pharmacological activity. 2014 , 15, 14 | 14 |
| 818 | In silico investigation of medicinal spectrum of imidazo-azines from the perspective of multitarget screening against malaria, tuberculosis and Chagas disease. 2014 , 50, 1-9 | 8 |
| 817 | The Complexation of the Anticancer Drug ThioTEPA with Methylated DNA Base Guanine: Combined Ab Initio and QTAIM Investigation. 2014 , 33, 104-14 | 7 |
| 816 | Structural insights into Estrogen Related Receptor- β modulation: 4-methylenesterols from <i>Theonella swinhoei</i> sponge as the first example of marine natural antagonists. 2014 , 80, 51-63 | 14 |
| 815 | Design, synthesis and preliminary bioactivity studies of 1,2-dihydrobenzo[d]isothiazol-3-one-1,1-dioxide hydroxamic acid derivatives as novel histone deacetylase inhibitors. 2014 , 22, 1529-38 | 14 |
| 814 | Direct interaction of garcinol and related polyisoprenylated benzophenones of <i>Garcinia cambogia</i> fruits with the transcription factor STAT-1 as a likely mechanism of their inhibitory effect on cytokine signaling pathways. 2014 , 77, 543-9 | 28 |
| 813 | AutoDock4(Zn): an improved AutoDock force field for small-molecule docking to zinc metalloproteins. 2014 , 54, 2371-9 | 152 |
| 812 | Understanding polyspecificity within the substrate-binding cavity of the human multidrug resistance P-glycoprotein. 2014 , 281, 673-82 | 50 |
| 811 | Zn(II)/pyridyloxime complexes as potential reactivators of OP-inhibited acetylcholinesterase: in vitro and docking simulation studies. 2014 , 134, 12-9 | 2 |
| 810 | Endogenous neurotoxic dopamine derivative covalently binds to Parkinson's disease-associated ubiquitin C-terminal hydrolase L1 and alters its structure and function. 2014 , 130, 826-38 | 20 |
| 809 | Radioiodinated benzyloxybenzene derivatives: a class of flexible ligands target to β -amyloid plaques in Alzheimer's brains. 2014 , 57, 6030-42 | 26 |
| 808 | Alginate-dependent gene expression mechanism in <i>Sphingomonas</i> sp. strain A1. 2014 , 196, 2691-700 | 14 |
| 807 | Synthesis of a D-ring isomer of galanthamine via a radical-based Smiles rearrangement reaction. 2014 , 79, 6759-64 | 17 |
| 806 | Structure-based virtual screening approach for discovery of covalently bound ligands. 2014 , 54, 1941-50 | 85 |
| 805 | DDI-CPI, a server that predicts drug-drug interactions through implementing the chemical-protein interactome. 2014 , 42, W46-52 | 50 |
| 804 | A knowledge-based approach for identification of drugs against vivapain-2 protein of <i>Plasmodium vivax</i> through pharmacophore-based virtual screening with comparative modelling. 2014 , 173, 2174-88 | 4 |

| | | |
|-----|---|-----|
| 803 | A function of SmeDEF, the major quinolone resistance determinant of <i>Stenotrophomonas maltophilia</i> , is the colonization of plant roots. 2014 , 80, 4559-65 | 56 |
| 802 | Exploration of the binding modes of buffalo PGRP1 receptor complexed with meso-diaminopimelic acid and lysine-type peptidoglycans by molecular dynamics simulation and free energy calculation. 2014 , 220, 255-68 | 7 |
| 801 | Structural insight into exosite binding and discovery of novel exosite inhibitors of botulinum neurotoxin serotype A through in silico screening. 2014 , 28, 765-78 | 13 |
| 800 | Synthesis of novel chiral TBBt derivatives with hydroxyl moiety. Studies on inhibition of human protein kinase CK2 and cytotoxicity properties. 2014 , 84, 364-74 | 15 |
| 799 | Fe ²⁺ and Cu ²⁺ increase the production of hyaluronic acid by lactobacilli via affecting different stages of the pentose phosphate pathway. 2014 , 173, 129-42 | 10 |
| 798 | Sterol carrier protein-2: binding protein for endocannabinoids. 2014 , 50, 149-58 | 21 |
| 797 | Synthesis and investigation of new Hesperadin analogues antitumor effects on HeLa cells. 2014 , 7, 85-91 | 4 |
| 796 | Dietary chlorophyllin abrogates TGF β signaling to modulate the hallmark capabilities of cancer in an animal model of forestomach carcinogenesis. 2014 , 35, 6725-37 | 6 |
| 795 | Molecular modeling of the transmembrane domain of mGluR2 metabotropic glutamate receptor and the binding site of its positive allosteric modulators. 2014 , 454, 13-6 | 1 |
| 794 | Role of berberine in anti-bacterial as a high-affinity LPS antagonist binding to TLR4/MD-2 receptor. 2014 , 14, 89 | 58 |
| 793 | Discovery of biaryl aminoquinazolines as novel tubulin polymerization inhibitors. 2014 , 57, 4598-4605 | 21 |
| 792 | Prediction of Potential Kinase Inhibitors in <i>Leishmania</i> spp. through a Machine Learning and Molecular Docking Approach. 2014 , 63-70 | |
| 791 | Insight into the binding mechanism of imipenem to human serum albumin by spectroscopic and computational approaches. 2014 , 11, 1785-97 | 148 |
| 790 | Secret of the major birch pollen allergen Bet v 1: identification of the physiological ligand. 2014 , 457, 379-90 | 61 |
| 789 | Oseltamivir analogues bearing N-substituted guanidines as potent neuraminidase inhibitors. 2014 , 57, 3154-60 | 33 |
| 788 | Computational characterization of ketone-ketal transformations at the active site of matrix metalloproteinases. 2014 , 118, 4345-50 | 5 |
| 787 | Topoisomerase I and II inhibitors: chemical structure, mechanisms of action and role in cancer chemotherapy. 2014 , 83, 82-94 | 22 |
| 786 | Bioactive compounds from culinary herbs inhibit a molecular target for type 2 diabetes management, dipeptidyl peptidase IV. 2014 , 62, 6147-58 | 92 |

| | | |
|-----|--|----|
| 785 | Agonist and antagonist binding in human glycine receptors. 2014 , 53, 6041-51 | 33 |
| 784 | Electrochemistry and molecular modeling of the hemoglobin-Benzene interaction with a nanocrystalline mixed metal oxide. 2014 , 4, 49128-49136 | 10 |
| 783 | Computational evaluations of charge coupling and hydrogen bonding in the active site of a family 7 cellobiohydrolase. 2014 , 118, 434-48 | 8 |
| 782 | Two isorecticular metal-organic frameworks with CdSO ₄ -like topology: selective gas sorption and drug delivery. 2014 , 43, 17265-73 | 48 |
| 781 | Alternative quality assessment strategy to compare performances of GPCR-ligand docking protocols: the human adenosine A _{2A} receptor as a case study. 2014 , 54, 2243-54 | 22 |
| 780 | Synthesis, structure and molecular docking studies of dicopper(II) complexes bridged by N-phenolato-N ² -[2-(dimethylamino)ethyl]oxamide: the influence of terminal ligands on cytotoxicity and reactivity towards DNA and protein BSA. 2014 , 38, 2964-2978 | 95 |
| 779 | Computationally efficient and accurate enantioselectivity modeling by clusters of molecular dynamics simulations. 2014 , 54, 2079-92 | 33 |
| 778 | Interaction between deferiprone and human serum albumin: multi-spectroscopic, electrochemical and molecular docking methods. 2014 , 64, 9-17 | 25 |
| 777 | Binding interaction of a prospective chemotherapeutic antibacterial drug with β -lactoglobulin: results and challenges. 2014 , 30, 5921-9 | 69 |
| 776 | Computer modeling of the complexes of Chlorin e6 with amphiphilic polymers. 2014 , 16, 10903-13 | 8 |
| 775 | (-)-Bisabolol inhibits preferentially electromechanical coupling on rat isolated arteries. 2014 , 63, 37-45 | 11 |
| 774 | Hsp90 inhibitors, part 1: definition of 3-D QSAutogrid/R models as a tool for virtual screening. 2014 , 54, 956-69 | 11 |
| 773 | New in silico insights into the inhibition of RNAP II by Eamanitin and the protective effect mediated by effective antidotes. 2014 , 51, 120-7 | 9 |
| 772 | Substrate specificity of human matriptase-2. 2014 , 97, 121-7 | 21 |
| 771 | Exploring the biophysical aspects and binding mechanism of thionine with bovine hemoglobin by optical spectroscopic and molecular docking methods. 2014 , 131, 43-52 | 45 |
| 770 | In silico design of small molecule inhibitors of CDK9/cyclin T1 interaction. 2014 , 50, 100-12 | 6 |
| 769 | The crystal structure of the amidohydrolase VinJ shows a unique hydrophobic tunnel for its interaction with polyketide substrates. 2014 , 588, 995-1000 | 7 |
| 768 | Discovery of the first potent and selective Mycobacterium tuberculosis Zmp1 inhibitor. 2014 , 24, 2508-11 | 19 |

| | | |
|-----|---|----|
| 767 | Toxicological and pharmacological evaluation, antioxidant, ADMET and molecular modeling of selected racemic chromenotacrine {11-amino-12-aryl-8,9,10,12-tetrahydro-7H-chromeno[2,3-b]quinolin-3-ols} for the potential prevention and treatment of Alzheimer's disease. 2014 , 74, 491-501 | 30 |
| 766 | Molecular insights into the promiscuous interaction of human pregnane X receptor (hPXR) with diverse environmental chemicals and drug compounds. 2014 , 96, 138-45 | 5 |
| 765 | Investigation of the interaction between five alkaloids and human hemoglobin by fluorescence spectroscopy and molecular modeling. 2014 , 123, 176-86 | 24 |
| 764 | Novel homozygous mutations in the WNT10B gene underlying autosomal recessive split hand/foot malformation in three consanguineous families. 2014 , 534, 265-71 | 18 |
| 763 | Pharmacogenomics of cantharidin in tumor cells. 2014 , 87, 399-409 | 41 |
| 762 | Inhibition of methemoglobin formation in aqueous solutions under aerobic conditions by the addition of amino acids. 2014 , 64, 267-75 | |
| 761 | Probing the human estrogen receptor-binding requirements for phenolic mono- and di-hydroxyl compounds: a combined synthesis, binding and docking study. 2014 , 22, 303-10 | 5 |
| 760 | New polyfunctional imidazo[4,5-C]pyridine motifs: synthesis, crystal studies, docking studies and antimicrobial evaluation. 2014 , 77, 288-97 | 24 |
| 759 | Substituent effects on the binding of natural product anthocyanidin inhibitors to influenza neuraminidase with mass spectrometry. 2014 , 828, 61-9 | 12 |
| 758 | Insights on pregnane-X-receptor modulation. Natural and semisynthetic steroids from Theonella marine sponges. 2014 , 73, 126-34 | 10 |
| 757 | Interaction between tachyplesin I, an antimicrobial peptide derived from horseshoe crab, and lipopolysaccharide. 2014 , 1844, 527-34 | 49 |
| 756 | Incisterols, highly degraded marine sterols, are a new chemotype of PXR agonists. 2014 , 83, 80-5 | 10 |
| 755 | Structural basis for the substrate specificity and the absence of dehalogenation activity in 2-chloromuconate cycloisomerase from Rhodococcus opacus 1CP. 2014 , 1844, 1541-9 | 2 |
| 754 | In silico derived small molecules bind the filovirus VP35 protein and inhibit its polymerase cofactor activity. 2014 , 426, 2045-58 | 59 |
| 753 | 1,4-Naphthoquinone, a pro-oxidant, suppresses immune responses via KEAP-1 glutathionylation. 2014 , 88, 95-105 | 23 |
| 752 | Photochemical and photobiological studies on furoquinazolines as new psoralen analogs. 2014 , 138, 43-54 | 7 |
| 751 | Molecular modeling and multispectroscopic studies of the interaction of mesalamine with bovine serum albumin. 2014 , 118, 422-9 | 40 |
| 750 | Electrophysiological and metabolic effects of CHF5074 in the hippocampus: protection against in vitro ischemia. 2014 , 81, 83-90 | 20 |

| | | |
|-----|---|----|
| 749 | The interaction of photoactivators with proteins during microfabrication. 2014 , 275, 81-88 | 3 |
| 748 | Biological activities of new monohydroxylated brassinosteroid analogues with a carboxylic group in the side chain. 2014 , 85, 58-64 | 15 |
| 747 | NMR and computational methods in the structural and dynamic characterization of ligand-receptor interactions. 2014 , 805, 271-304 | 5 |
| 746 | Computational insights into the binding mechanism of antagonists with neuropeptide B/W receptor 1. 2014 , 10, 2236-46 | 1 |
| 745 | Adenine nucleotide translocase is acetylated in vivo in human muscle: Modeling predicts a decreased ADP affinity and altered control of oxidative phosphorylation. 2014 , 53, 3817-29 | 39 |
| 744 | The thiocarbamate disulphide drug, disulfiram induces osteopenia in rats by inhibition of osteoblast function due to suppression of acetaldehyde dehydrogenase activity. 2014 , 139, 257-70 | 14 |
| 743 | Structure-based mutational studies of substrate inhibition of betaine aldehyde dehydrogenase BetB from <i>Staphylococcus aureus</i> . 2014 , 80, 3992-4002 | 31 |
| 742 | Diamidine compounds for selective inhibition of protein arginine methyltransferase 1. 2014 , 57, 2611-22 | 59 |
| 741 | <i>Mycobacterium tuberculosis</i> keto-mycolic acid and macrophage nuclear receptor TR4 modulate foamy biogenesis in granulomas: a case of a heterologous and noncanonical ligand-receptor pair. 2014 , 193, 295-305 | 47 |
| 740 | Inhibition of 17 β -HSD1: SAR of bicyclic substituted hydroxyphenylmethanones and discovery of new potent inhibitors with thioether linker. 2014 , 82, 394-406 | 9 |
| 739 | Supramolecular complex formed by DNA oligonucleotide and thiacalix[4]arene. NMR-spectroscopy and molecular docking. 2014 , 1074, 126-133 | 12 |
| 738 | Alditols and monosaccharides from sorghum vinegar can attenuate platelet aggregation by inhibiting cyclooxygenase-1 and thromboxane-A2 synthase. 2014 , 155, 285-92 | 12 |
| 737 | Progress towards automated Kepler scientific workflows for computer-aided drug discovery and molecular simulations. 2014 , 29, 1745-1755 | 6 |
| 736 | Purification, characterization, and in vitro activity of 2,4-Di-tert-butylphenol from <i>Pseudomonas monteilii</i> PsF84: conformational and molecular docking studies. 2014 , 62, 6138-46 | 57 |
| 735 | SB-RA-2001 inhibits bacterial proliferation by targeting FtsZ assembly. 2014 , 53, 2979-92 | 30 |
| 734 | Preferential binding of an odor within olfactory receptors: a precursor to receptor activation. 2014 , 39, 107-23 | 15 |
| 733 | 7-MEOTA-donepezil like compounds as cholinesterase inhibitors: Synthesis, pharmacological evaluation, molecular modeling and QSAR studies. 2014 , 82, 426-38 | 70 |
| 732 | Traditional Chinese herbs as chemical resource library for drug discovery of anti-infective and anti-inflammatory. 2014 , 155, 589-98 | 20 |

| | | |
|-----|---|----|
| 731 | Reverse docking: a powerful tool for drug repositioning and drug rescue. 2014 , 6, 333-42 | 73 |
| 730 | Structure-based design of small-molecule protein-protein interaction modulators: the story so far. 2014 , 6, 343-57 | 39 |
| 729 | A strategic solution to optimize molecular docking simulations using Fully-Flexible Receptor models. 2014 , 41, 7608-7620 | 10 |
| 728 | An investigation of in vitro cytotoxicity and apoptotic potential of aromatic diselenides. 2014 , 24, 3440-6 | 30 |
| 727 | Applications of structure-based design to antibacterial drug discovery. 2014 , 55, 69-76 | 16 |
| 726 | In vitro and in silico investigations of the binding interactions between chlorophenols and trypsin. 2014 , 278, 55-65 | 30 |
| 725 | Application of ANS fluorescent probes to identify hydrophobic sites on the surface of DREAM. 2014 , 1844, 1472-80 | 19 |
| 724 | Carbohydrate-Small Molecule Hybrids as Lead Compounds Targeting IL-6 Signaling. 2023 , 28, 677 | 0 |
| 723 | Interactions of fentanyl with blood platelets and plasma proteins: platelet sensitivity to prasugrel metabolite is not affected by fentanyl under in vitro conditions. | 0 |
| 722 | DFT study on molecular structure, spectroscopic properties, Hirshfeld surface and molecular docking reveals the potential of flavones based on experimental and theoretical investigations. | 0 |
| 721 | Switching Prenyl Donor Specificities in Squalene Synthase-Like Aromatic Prenyltransferases from Bacterial Carbazole Alkaloid Biosynthesis. | 0 |
| 720 | The molecular classification of cancer-associated fibroblasts on a pan-cancer single-cell transcriptional profiling. | 0 |
| 719 | Structural homology between 11 beta-hydroxysteroid dehydrogenase and Mycobacterium tuberculosis Inh-A enzyme: Dehydroepiandrosterone as a potential co-adjuvant treatment in diabetes-tuberculosis comorbidity. 13, | 0 |
| 718 | In Vitro and In Silico Biological Studies of 4-Phenyl-2-quinolone (4-PQ) Derivatives as Anticancer Agents. 2023 , 28, 555 | 0 |
| 717 | Use of Apatinib as a Bait to Fish Its Unexpected Kinase Targets from the Hepatocellular Carcinoma Druggable Kinome. | 0 |
| 716 | Acarbose May Function as a Competitive Exclusion Agent for the Producing Bacteria. | 0 |
| 715 | Computer-aided drug design for the Pain-Like protease (PLpro) inhibitors against SARS-CoV-2. 2023 , 114247 | 0 |
| 714 | Novel N-benzyl-2-oxo-1,2-dihydrofuro [3,4-d]pyrimidine-3(4H)-carboxamide as anticancer agent: Synthesis, drug-likeness, ADMET profile, DFT and molecular modelling against EGFR target. 2023 , e12948 | 0 |

- 713 Neural Networks in the Design of Molecules with Affinity to Selected Protein Domains. **2023**, 24, 1762 ○
- 712 Activity of Cytosolic Ascorbate Peroxidase (APX) from *Panicum virgatum* against Ascorbate and Phenylpropanoids. **2023**, 24, 1778 ○
- 711 Computational design of prospective molecular targets for *Burkholderia cepacia* complex by molecular docking and dynamic simulation studies. ○
- 710 Native amine dehydrogenases can catalyze the direct reduction of carbonyl compounds to alcohols in the absence of ammonia. 3, ○
- 709 Identification of new dihydrophenanthrene derivatives as promising anti-SARS-CoV-2 drugs through in silico investigations. **2023**, 1-16 ○
- 708 Therapeutic effect of combination vitamin D3 and siponimod on remyelination and modulate microglia activation in cuprizone mouse model of multiple sclerosis. 16, ○
- 707 In Silico Mining of Natural Products Atlas (NPAtlas) Database for Identifying Effective Bcl-2 Inhibitors: Molecular Docking, Molecular Dynamics, and Pharmacokinetics Characteristics. **2023**, 28, 783 ○
- 706 Metal-free visible light mediated direct C-H amination of benzoxazole with secondary amines. 1 ○
- 705 Targeting a transcription factor NF- κ B by green tea catechins using in silico and in vitro studies in pancreatic cancer. 9, ○
- 704 Insights into the mechanism of phospholipid hydrolysis by plant non-specific phospholipase C. **2023**, 14, ○
- 703 Carfentanil structural analogs found in street drugs by paper spray mass spectrometry and their characterization by high-resolution mass spectrometry. ○
- 702 6-Shogaol Exhibits Anti-viral and Anti-inflammatory Activity in COVID-19-Associated Inflammation by Regulating NLRP3 Inflammasomes. **2023**, 8, 2618-2628 1
- 701 Nano-Zirconium Dioxide Catalyzed Multicomponent Synthesis of Bioactive Pyranopyrazoles That Target Cyclin Dependent Kinase 1 in Human Breast Cancer Cells. **2023**, 11, 172 1
- 700 Improving Lurasidone Hydrochloride's Solubility and Stability by Higher-Order Complex Formation with Hydroxypropyl- β -cyclodextrin. **2023**, 15, 232 ○
- 699 Xanthatin and 8-epi-xanthatin as new potential colchicine binding site inhibitors: a computational study. **2023**, 29, ○
- 698 Designing Tailored Thiosemicarbazones with Bespoke Properties: The Styrene Moiety Imparts Potent Activity, Inhibits Heme Center Oxidation, and Results in a Novel Stealth Zinc(II) Complex 1 ○
- 697 Structural characterization on a β -galactosidase Aga86A_Wa from *Wenyngzhuangia aestuarii* reveals the prevalent methyl-galactose accommodation capacity of GH86 enzymes at subsite II. **2023**, 120594 ○
- 696 Recent applications of computational methods to allosteric drug discovery. 9, ○

| | | |
|-----|--|---|
| 695 | Structural and functional analysis of a tandem repeat galacturonic acid-binding lectin from the sea hare <i>Aplysia californica</i> . 2023 , 132, 108513 | 0 |
| 694 | Determination of nucleoside DOT1L inhibitors' residence times by RAMD simulations. 2, | 0 |
| 693 | Exploration on Ononin and Corylin molecule Against Anti-Influenza H1N1 A Virus via Molecular Docking, Molecular dynamics simulation and Binding free energy calculations. | 0 |
| 692 | Identifying Dopamine D3 Receptor Ligands through Virtual Screening and Exploring the Binding Modes of Hit Compounds. 2023 , 28, 527 | 1 |
| 691 | Targeting RNA:protein interactions with an integrative approach leads to the identification of potent YBX1 inhibitors. 12, | 0 |
| 690 | In silico Screening of Potential SARS-CoV-2 Main Protease Inhibitors from <i>Thymus schimperii</i> . Volume 16, 1-13 | 0 |
| 689 | Silibinin chronic treatment in a rat model of Parkinson disease: A comprehensive in-vivo evaluation and in silico molecular modeling. 2023 , 175517 | 1 |
| 688 | Peonidin-3-O-Glucoside from Purple Corncob Ameliorates Nonalcoholic Fatty Liver Disease by Regulating Mitochondrial and Lysosome Functions to Reduce Oxidative Stress and Inflammation. 2023 , 15, 372 | 0 |
| 687 | Design, synthesis, spectroscopic characterizations, single crystal X-ray analysis, in vitro xanthine oxidase and acetylcholinesterase inhibitory evaluation as well as in silico evaluation of selenium-based N-heterocyclic carbene compounds. 1-20 | 0 |
| 686 | Investigation of pharmacodynamic material basis of <i>Anemarrhenae Rhizoma</i> and its processed products based on plant metabolomics and molecular docking technology. | 1 |
| 685 | Proteogenomic Approaches to Understand Gene Mutations and Protein Structural Alterations in Colon Cancer. 2023 , 3, 11-29 | 0 |
| 684 | The mitigative effect of lotus root (<i>Nelumbo nucifera Gaertn</i>) extract on acute alcoholism through activation of alcohol catabolic enzyme, reduction of oxidative stress, and protection of liver function. 9, | 0 |
| 683 | GABAA and serotonergic receptors participation in anxiolytic effect of chalcones in adult zebrafish. 1-19 | 0 |
| 682 | Antileishmanial Activity of <i>Clinanthus milagroanthus</i> S. Leiva & Meerow (<i>Amaryllidaceae</i>) Collected in Peru. 2023 , 12, 322 | 0 |
| 681 | Drug Repurposing to Inhibit Histamine N-Methyl Transferase. 2023 , 28, 576 | 0 |
| 680 | Exploring pradimicin-IRD antineoplastic mechanisms and related DNA repair pathways. 2023 , 371, 110342 | 0 |
| 679 | Bionics design of affinity peptide inhibitors for SARS-CoV-2 RBD to block SARS-CoV-2 RBD-ACE2 interactions. 2023 , e12890 | 0 |
| 678 | New amidine-benzenesulfonamides as iNOS inhibitors for the therapy of the triple negative breast cancer. 2023 , 248, 115112 | 1 |

- 677 Unravelling structural, functional, evolutionary and genetic basis of SWEET transporters regulating abiotic stress tolerance in maize. **2023**, 229, 539-560 1
- 676 Albumin-binding properties of an aromatic N-acylhydrazone. **2023**, 372, 121180 0
- 675 Synthesis of piperine analogues as AChE and BChE inhibitors for the treatment of Alzheimer's disease. **2023**, 53, 216-221 0
- 674 Rationally designed donepezil-based hydroxamates modulate Sig-1R and HDAC isoforms to exert anti-glioblastoma effects. **2023**, 248, 115054 0
- 673 Machine intelligence-guided selection of optimized inhibitor for human immunodeficiency virus (HIV) from natural products. **2023**, 153, 106525 0
- 672 Synthesis, structural and theoretical investigations on 3-diethyl 2-((4-[3-ethoxy-2-(ethoxycarbonyl)-3-oxo-2-phenylpropyl]-2,5-dimethylphenyl)methyl)-2-phenylpropanedioate. **2023**, 100, 100869 0
- 671 Identification of a type II LacNAc specific binding lectin CMRBL from *Cordyceps militaris*. **2023**, 230, 123207 0
- 670 Biological activity and inhibition potential against α -glucosidase and α -amylase of 2,4-di-tert-butylphenol from bamboo shoot extract by in vitro and in silico studies. **2023**, 126, 15-22 0
- 669 Affinity screening of potential anti-obesity and anti-diabetic component from pomegranate peel by co-immobilization of lipase and α -amylase using carbon nanotube and hydrogel. **2023**, 126, 51-60 0
- 668 Synthesis, crystal structure, Hirshfeld surface, energy framework, NCI-RDG, theoretical calculations and molecular docking of (Z)-4,4'-bis[3-N-ethyl-2-N'-(phenylimino)thiazolidin-4-one]methane. **2023**, 1277, 134781 0
- 667 DFT-aided infrared and electronic circular dichroism spectroscopic study of cyclopeptide S-PK6 and the exploration of its antitumor potential by molecular docking. **2023**, 1278, 134903 1
- 666 Investigation of crystal structures, spectral (FT-IR and NMR) analysis, DFT, and molecular docking studies of novel piperazine derivatives as antineurotic drugs. **2023**, 1278, 134937 0
- 665 Enhancing the asymmetric reduction activity of ene-reductases for the synthesis of a brivaracetam precursor. **2023**, 126, 108-116 0
- 664 Ligand-observed in-tube NMR in natural products research: A review on enzymatic biotransformations, protein-ligand interactions, and in-cell NMR spectroscopy. **2023**, 16, 104536 0
- 663 Design, synthesis and antifungal activity of novel pyrazole amides derivatives. **2023**, 1277, 134881 0
- 662 An easy-to-use phosphate triggered zinc-azophenine complex assisted metal extrusion assay: A diagnostic approach for chronic kidney disease and in silico docking studies. **2023**, 548, 121364 0
- 661 Molecular interactions of resveratrol with A β 42 peptide and fibril during in-vitro A β 42 aggregation. **2023**, 7, 100060 0
- 660 Chemical profile, antiproliferative and antibacterial activities and docking studies of essential oil and hexane fraction of hydrosol from fresh leaf of *Plectranthus amboinicus* (Lour.) Spreng.. **2023**, 107, 104595 1

- 659 Extraction of protein with protease inhibitor activity from Brazilwood (*Caesalpinia echinata* LAM.) seeds using choline-based ionic liquids. **2023**, 31, 100956 ○
- 658 Mitigating the inhibition of antibacterial agent chloroxylenol on nitrification system-The role of *Rhodococcus ruber* in a bioaugmentation system. **2023**, 447, 130758 1
- 657 Identification of Key Genes as Potential Drug Targets for Gastric Cancer. **2023**, 28, 649-664 ○
- 656 Multi spectroscopic and molecular simulation studies of propyl acridone binding to calf thymus DNA in the presence of electromagnetic force. **2023**, 13, 5-16 ○
- 655 Comparative In-Silico Molecular Docking of Silymarin for SARS-CoV-2 Receptor. **2022**, 2, 58-67 ○
- 654 Synthesis and Antiproliferative Activity against Cancer Cells of Indole-Aryl-Amide Derivatives. **2023**, 28, 265 ○
- 653 Synthesis, Photochemistry, Computational Study and Potential Application of New Styryl-Thiophene and Naphtho-Thiophene Benzylamines. **2023**, 24, 610 ○
- 652 Ethanolic Extracts of *Datura innoxia* Have Promising Acaricidal Activity against *Rhipicephalus microplus* as It Blocks the Glutathione S-Transferase Activity of the Target Tick. **2023**, 14, 118 1
- 651 In silico Screening, Docking, and Redesigning of Traditional Chinese Medicinal Compounds Against *Streptococcus pneumoniae* Glycosyl Hydrolase GHIP and Peptidoglycan Hydrolase LytB. **2022**, ○
- 650 Investigating Key Targets of Dajianzhong Decoction for Treating Crohn's Disease Using Weighted Gene Co-Expression Network. **2023**, 11, 112 ○
- 649 Chemical and Biological Investigations of *Allium scorodoprasum* L. Flower Extracts. **2023**, 16, 21 1
- 648 *Rosa gallica* and its active compound, cyanidin-3, 5- O -diglucoside, improve skin hydration via the GLK signaling pathway. ○
- 647 A sorghum ascorbate peroxidase with four binding sites has activity against ascorbate and phenylpropanoids. ○
- 646 Crystal structure, in silico molecular docking, DFT analysis and ADMET studies of N-(2-methoxy-benzyl)-acetamide. **2022**, 13, 440-450 ○
- 645 Design and Synthesis of Conformationally Flexible Scaffold as Bitopic Ligands for Potent D3-Selective Antagonists. **2023**, 24, 432 1
- 644 Deaza-modification of MR1 ligands modulates recognition by MR1-restricted T cells. **2022**, 12, ○
- 643 Effects of Trimethylamine and Trimethylamine Oxide on Human Serum Albumin Observed by Tryptophan Fluorescence and Absorbance Spectroscopies. **2023**, 13, 421 ○
- 642 A Guide to In Silico Drug Design. **2023**, 15, 49 ○

- 641 Computational Drug Repurposing Approach to Identify Novel Inhibitors of ILK Protein for Treatment of Esophageal Squamous Cell Carcinoma. **2022**, 2022, 1-10 ○
- 640 In-Silico analysis of potential antidiabetic phytoconstituents from Manilkara hexandra against mitoNEET. ○
- 639 Structure Prediction and Binding Site Analysis of Human Sperm Hyaluronidases. **2022**, 13, 96-100 ○
- 638 ViTRMSE: a three-dimensional RMSE scoring method for protein-ligand docking models based on Vision Transformer. **2022**, ○
- 637 IDENTIFICATION OF FLAVONOIDS FROM ACALYPHA INDICA L. (EUPHORBIACEAE) AS CASPASE-3 ACTIVATORS USING MOLECULAR DOCKING AND MOLECULAR DYNAMICS. 162-166 ○
- 636 Synthesis, Crystallographic Structure, Theoretical Analysis, Molecular Docking Studies, and Biological Activity Evaluation of Binuclear Ru(II)-1-Naphthylhydrazine Complex. **2023**, 24, 689 1
- 635 In silico docking based screening of constituents from Persian shallot as modulators of human glucokinase. ○
- 634 Techniques of Bioremediation using bacteria for the treatment of polycyclic aromatic hydrocarbons: A Review. **2022**, 10, 1318-1330 ○
- 633 Curcumin Decreases Viability and Inhibits Proliferation of Imatinib-Sensitive and Imatinib-Resistant Chronic Myeloid Leukemia Cell Lines. **2023**, 13, 58 1
- 632 Binding to Iron Quercetin Complexes Increases the Antioxidant Capacity of the Major Birch Pollen Allergen Bet v 1 and Reduces Its Allergenicity. **2023**, 12, 42 ○
- 631 Design, Synthesis, Characterization, and Analysis of Antimicrobial Property of Novel Benzophenone Fused Azetidinone Derivatives through In Vitro and In Silico Approach. **2023**, 45, 92-109 ○
- 630 Identification and structural analysis of a carbohydrate binding module specific to alginate, a representative of a new family, CBMxx#. **2022**, 102854 ○
- 629 Corosolic Acid Inhibits Secretory Phospholipase A2IIa as an Anti-Inflammatory Function and Exhibits Anti-Tumor Activity in Ehrlich Ascites Carcinoma Bearing Mice. Volume 15, 6905-6921 ○
- 628 Disruption of DYRK1A-induced hyperphosphorylation of amyloid-beta and tau protein in Alzheimer's disease: An integrative molecular modeling approach. 9, ○
- 627 Synthesis, Quantum Computational Analysis and Molecular Docking of 3-(2-Hydroxyphenyl)-1-Phenyl Propanone: A Combined Experimental and Theoretical Analysis. 1-30 ○
- 626 Elucidating the toxicity mechanism of AFM2 and the protective role of quercetin in albino mice. **2023**, 13, ○
- 625 DFT calculation and molecular docking of lawsone and its derivatives as antibacterial. **2023**, ○
- 624 Small Molecules Targeting the RNA-Binding Protein HuR Inhibit Tumor Growth in Xenografts. **2023**, 66, 2032-2053 ○

- 623 Combination of DNA Damage, Autophagy, and ERK Inhibition: Novel Evodiamine-Inspired Multi-Action Pt(IV) Prodrugs with High-Efficiency and Low-Toxicity Antitumor Activity. **2023**, 66, 1852-1872 ○
- 622 Graphene oxide and flavonoids as potential inhibitors of the spike protein of SARS-CoV-2 variants and interaction between ligands: a parallel study of molecular docking and DFT. ○
- 621 Conjugates of Tacrine and Salicylic Acid Derivatives as New Promising Multitarget Agents for Alzheimer's Disease. **2023**, 24, 2285 ○
- 620 Potential of hydroethanolic leaf extract of *Ocimum sanctum* in ameliorating redox status and lung injury in COPD: an in vivo and in silico study. **2023**, 13, ○
- 619 Nur77 Serves as a Potential Prognostic Biomarker That Correlates with Immune Infiltration and May Act as a Good Target for Prostate adenocarcinoma. **2023**, 28, 1238 ○
- 618 In silico Structure Prediction, Molecular Docking, and Dynamic Simulation of Plasmodium falciparum AP2-I Transcription Factor. **2023**, 17, 117793222211496 ○
- 617 Improve the Biodistribution with Bulky and Lipophilic Modification Strategies on Lys-Urea-Glu-Based PSMA-Targeting Radiotracers. **2023**, 20, 1435-1446 ○
- 616 Robust anti-inflammatory activity of genistein against neutrophil elastase: a microsecond molecular dynamics simulation study. 1-17 1
- 615 Molecular dynamics simulations depict structural motions of the whole human aryl hydrocarbon receptor influencing its binding of ligands and HSP90. 1-16 ○
- 614 Withanolides of *Athenaea velutina* with potential inhibitory properties against SARS coronavirus main protease (mpro): molecular modeling studies. 1-9 ○
- 613 Identification of potential inhibitor molecule against MabA protein of *Mycobacterium leprae* by integrated in silico approach. 1-16 ○
- 612 HuR modulation with tanshinone mimics impairs LPS response in murine macrophages. ○
- 611 In silico Identification of Triclosan Derivatives as Potential Inhibitors of Mutant *Mycobacterium tuberculosis* InhA. ○
- 610 Enzyme PTP-1B Inhibition Studies by Vanadium Metal Complexes: a Kinetic Approach. ○
- 609 Investigation on Phyto-active Constituent of *Clerodendrum paniculatum* as Therapeutic Agent against Viral Diseases. **2023**, 8, ○
- 608 Synthesis, Molecular Dynamics Simulation, and In-vitro Antitumor Activity of Quinazoline-2,4,6-triamine Derivatives as Novel EGFR Tyrosine Kinase Inhibitors. **2023**, 21, ○
- 607 Biologically Potent Benzimidazole-Based-Substituted Benzaldehyde Derivatives as Potent Inhibitors for Alzheimer's Disease along with Molecular Docking Study. **2023**, 16, 208 ○
- 606 New Quinoline Analogues: As Potential Diabetics Inhibitors and Molecular Docking Study. 1-23 ○

- 605 Contribution of circulating host and microbial tryptophan metabolites towards Ah receptor activation. ○
- 604 Discovery of a Novel Covalent EZH2 Inhibitor Based on Tazemetostat Scaffold for the Treatment of Ovarian Cancer. **2023**, 66, 1725-1741 ○
- 603 Antimicrobial, antiproliferative effects and docking studies of methoxy group enriched coumarin-chalcone hybrids. ○
- 602 Molecular dynamic, Hirshfeld surface, molecular docking and drug likeness studies of a potent anti-oxidant, anti-malaria and anti-inflammatory medicine: Pyrogallol. **2023**, 5, 100763 ○
- 601 PTCHD1 Binds Cholesterol but Not Sonic Hedgehog, Suggesting a Distinct Cellular Function. **2023**, 24, 2682 ○
- 600 Polyphenolic Profile of Herniaria hemistemon Aerial Parts Extract and Assessment of Its Anti-Cryptosporidiosis in a Murine Model: In Silico Supported In Vivo Study. **2023**, 15, 415 ○
- 599 Interaction of copper potential metallodrugs with TMPRSS2: A comparative study of docking tools and its implications on COVID-19. 11, ○
- 598 Water-SDS-Ionic Liquid Catalytic System for the Synthesis of Pyrano-chromenes and in-silico Approach to Predict Inhibitory Activity Against Mpro of SARS-CoV-2**. **2023**, 8, ○
- 597 DeepMPF: deep learning framework for predicting drug-target interactions based on multi-modal representation with meta-path semantic analysis. **2023**, 21, 1
- 596 Diclofenac and Meloxicam Exhibited Anti-Virulence Activities Targeting Staphyloxanthin Production in Methicillin-Resistant Staphylococcus aureus. **2023**, 12, 277 ○
- 595 Mining and Characterization of Thermophilic Glucose Isomerase Based on Virtual Probe Technology. ○
- 594 Phylogenetic analysis, computer modeling and catalytic prediction of an Amazonian soil β -glucosidase against a soybean saponin. ○
- 593 Regulation of Intersubunit Interactions in Homotetramer of Glyceraldehyde-3-Phosphate Dehydrogenases upon Its Immobilization in Protein κ -Carrageenan Gels. **2023**, 15, 676 ○
- 592 Design, Synthesis and Pharmacological Evaluation of New Quinoline-Based Panx-1 Channel Blockers. **2023**, 24, 2022 ○
- 591 Antiproliferative Activity of Antibiotics through DNA Binding Mechanism: Evaluation and Molecular Docking Studies. **2023**, 24, 2563 ○
- 590 Ecofriendly aminochalcogenation of alkenes: a green alternative to obtain compounds with potential anti SARS-CoV-2 activity. ○
- 589 The Effect of Alkali Iodide Salts in the Inclusion Process of Phenolphthalein in β -Cyclodextrin: A Spectroscopic and Theoretical Study. **2023**, 28, 1147 ○
- 588 Recent advances in the area of plant-based anti-cancer drug discovery using computational approaches. ○

- 587 Using Defect Control To Break the Stability-Activity Trade-Off in Enzyme Immobilization via Competitive Coordination. ○
- 586 Bioinformatics-Structural Approach to the Search for New D-Amino Acid Oxidases. **2022**, 14, 57-68 ○
- 585 Cyclohexane-1,3-dione Derivatives as Future Therapeutic Agents for NSCLC: QSAR Modeling, In Silico ADME-Tox Properties, and Structure-Based Drug Designing Approach. **2023**, 8, 4294-4319 ○
- 584 Molecular Docking in the Study of Ligand-Protein Recognition: An Overview. ○
- 583 MetalProGNet: a structure-based deep graph model for metalloprotein-ligand interaction predictions. ○
- 582 Identification of a new class of proteasome inhibitors based on a naphthyl-azotricyclic-urea-phenyl scaffold. ○
- 581 In vitro contraceptive activities, molecular docking, molecular dynamics, MM-PBSA, non-covalent interaction and DFT studies of bioactive compounds from *Aegle marmelos*. Linn., leaves. 11, ○
- 580 In silico discovery of antioxidant peptides from the sea grass *Posidonia australis*. **2023**, 197-213 ○
- 579 Essential Oils Composition and Biological Activity of *Chamaecyparis obtusa*, *Chrysopogon nigrifolius* and *Lavandula coronopifolia* Grown Wild in Sudan. **2023**, 28, 1005 1
- 578 Valeramide and Halo-phenol in a Non-polar Liquid: DFT Based Characterization and Reactivity, Non-covalent Interaction, and Dielectric Relaxation Studies. 1-26 ○
- 577 From big data to complex network: a navigation through the maze of drug-target interaction. **2023**, 407-436 ○
- 576 Targeting Y220C mutated p53 by *Foeniculum vulgare*-derived phytochemicals as cancer therapeutics. **2023**, 29, ○
- 575 Molecular interaction modeling of carbon nanotubes and fullerene toward prioritized targets of SARS-CoV-2 by computer-aided screening and docking studies. **2023**, 157-179 ○
- 574 The In Silico Characterization of Monocotyledonous β -Arabinofuranosidases on the Example of Maize. **2023**, 13, 266 ○
- 573 Novel Inhibitors of Acetyl- and Butyrylcholinesterase Derived from Benzohydrazides: Synthesis, Evaluation and Docking Study. **2023**, 16, 172 ○
- 572 Long-Term Supplementation of *Syzygium cumini* (L.) Skeels Concentrate Alleviates Age-Related Cognitive Deficit and Oxidative Damage: A Comparative Study of Young vs. Old Mice. **2023**, 15, 666 ○
- 571 UHPLC-MS Phytochemical Profiling and Insight into Bioactivity of *Rabelera holostea* (Greater Stitchwort) Extract. **2023**, 28, 1274 ○
- 570 Click-designed vanilloid-triazole conjugates as dual inhibitors of AChE and A β aggregation. **2023**, 13, 2871-2883 ○

- 569 Employing Molecular Docking Calculations for the Design of Alkyl (2-Alcoxy-2-Hydroxypropanoyl)-L-Tryptophanate Derivatives as Potential Inhibitors of 11 β -Hydroxysteroid Dehydrogenase Type 1 (11 β -HSD1). **2023**, 4, 108-116 ○
- 568 In Vitro, In Vivo, and In Silico Analyses of Molecular Anti-Pigmentation Mechanisms of Selected Thai Rejuvenating Remedy and Bioactive Metabolites. **2023**, 28, 958 ○
- 567 Nano-ZrO₂-Catalyzed Biginelli Reaction and the Synthesis of Bioactive Dihydropyrimidinones That Targets PPAR- γ in Human Breast Cancer Cells. **2023**, 13, 228 ○
- 566 PI(3,5)P₂ Controls the Signaling Activity of Class I PI3K. ○
- 565 MetaDOCK: A Combinatorial Molecular Docking Approach. ○
- 564 Chitinase is a Potent Insecticidal Molecular Target of Camptothecin and Its Derivatives. **2023**, 71, 1845-1851 ○
- 563 Long-Lived States Provide Insights from NMR into the β -Cyclodextrin Drug Assemblies. **2023**, 127, 1158-1167 ○
- 562 Computational design and structure dynamics analysis of bifunctional chimera of endoxylanase from *Clostridium thermocellum* and xylosidase from *Bacteroides ovatus*. **2023**, 13, ○
- 561 Novel Signposts on the Road from Natural Sources to Pharmaceutical Applications: A Combinative Approach between LC-DAD-MS and Offline LC-NMR for the Biochemical Characterization of Two Hypericum Species (*H. montbretii* and *H. origanifolium*). **2023**, 12, 648 ○
- 560 MANF regulates neuronal survival and UPR through its ER-located receptor IRE1 β . **2023**, 42, 112066 ○
- 559 Spectral-fluorescent and photochemical study of 6,6'-di(benzoylamino)trimethine cyanine dyes in solutions as possible probes for DNA. **2023**, 292, 122416 ○
- 558 Curcumin and whey protein concentrate binding: Thermodynamic and structural approach. **2023**, 139, 108547 ○
- 557 Computational investigation into Nirematrelvir/Ritonavir synergetic efficiency compared with some approved antiviral drugs targeting main protease (Mpro) SARS-CoV-2 Omicron variant. **2023**, 100, 100891 ○
- 556 The chemical reactivity and antimalarial investigation of crystal structure (2E)-3-(biphenyl-4-yl)-1-(4-chlorophenyl)prop-2-en-1-one and hydroxyphenyl, nitrophenyl substituted chalcone derivative molecules. **2023**, 1280, 135001 ○
- 555 Comparative biochemical characterization of mammalian-derived CYP11A1s with cholesterol side-chain cleavage activities. **2023**, 229, 106268 ○
- 554 Pathway network-based quantitative modeling of the time-dependent and dose-response anti-inflammatory effect of Reduning Injection. **2023**, 307, 116216 ○
- 553 A rhodanine derivative as a potential antibacterial and anticancer agent: Crystal structure, spectral characterization, DFT calculations, Hirshfeld surface analysis, in silico molecular docking and ADMET studies. **2023**, 1280, 135025 ○
- 552 Molecular dynamics simulations suggest Thiosemicarbazones can bind p53 cancer mutant R175H. **2023**, 1871, 140903 ○

- 551 A novel Cu(II)-based DNA-intercalating agent: Structural and biological insights using biophysical and in silico techniques. **2023**, 293, 122438 ○
- 550 Unveiling the tartrazine binding mode with dsDNA by UV-visible spectroscopy, electrochemical, and QM/MM methods. **2023**, 292, 122400 ○
- 549 Experimental and theoretical studies of a thiourea derivative: 1-(4-chloro-benzoyl)-3-(2-trifluoromethyl-phenyl)thiourea. **2023**, 1279, 134996 ○
- 548 In silico Investigation of Immulina Glycosidic Residues as Immunostimulants by Targeting Penaeus monodon Shrimp Toll-Receptor 9. **2022**, ○
- 547 Screening of Secondary Metabolite Compounds of Gorontalo Traditional Medicinal Plants Using the In Silico Method as a Candidate for SARS-CoV-2 Antiviral. **2023**, 25, 382-393 ○
- 546 Organotin compound as an inhibitor of nitric oxide formation. **2022**, 71, 2605-2611 ○
- 545 The substrate specificity in the O-demethylation of 4-alkylguaiacols by cytochrome P450 AgcAP450. **2023**, 13, 2070-2079 ○
- 544 A tris(2-aminoethyl)amine-based zinc complex as a highly water-soluble drug carrier for the anti-COVID-19 drug favipiravir: a joint experimental and theoretical study. ○
- 543 Coumarin-based derivatives targeting Trypanosoma cruzi cruzain and Trypanosoma brucei cathepsin L-like proteases. ○
- 542 Structural and functional characterization of 1-deoxy-D-xylulose-5-phosphate synthase (DXS) from Acinetobacter baumannii: identification of promising lead molecules from virtual screening, molecular docking and molecular dynamics simulations. 1-14 ○
- 541 GC/MS Analysis, Cytotoxicity, and Antiviral Activities of Annona glabra Hexane Extract Supported by In Silico Study. **2023**, 28, 1628 ○
- 540 Inhibitory effect and underlying mechanism of cinnamon and clove essential oils on Botryosphaeria dothidea and Colletotrichum gloeosporioides causing rots in postharvest bagging-free apple fruits. 14, ○
- 539 Scalable Inhibitors of the Nsp3-Nsp4 Coupling in SARS-CoV-2. **2023**, 8, 5349-5360 ○
- 538 Phosphomevalonate Kinase Controls β -Catenin Signaling via the Metabolite 5-Diphosphomevalonate. 2204909 ○
- 537 RNA as a component of fibrils from Alzheimer's disease and other neurodegenerations. ○
- 536 Enzymatic Glyco-Modification of Synthetic Membrane Systems. **2023**, 13, 335 ○
- 535 Template Entrance Channel as Possible Allosteric Inhibition and Resistance Site for Quinolines Tricyclic Derivatives in RNA Dependent RNA Polymerase of Bovine Viral Diarrhea Virus. **2023**, 16, 376 ○
- 534 New, Low-Molecular Weight Chemical Compounds Inhibiting Biological Activity of Interleukin 15. **2023**, 28, 2287 ○

- 533 Exploring the structural basis to develop efficient multi-epitope vaccines displaying interaction with HLA and TAP and TLR3 molecules to prevent NIPAH infection, a global threat to human health. **2023**, 18, e0282580 ○
- 532 Evidence That Binding of Cyclic GMP to the Extracellular Domain of Na⁺/K⁺-ATPase Mediates Natriuresis. ○
- 531 Coumarin-Based Compounds as Inhibitors of Tyrosinase/Tyrosine Hydroxylase: Synthesis, Kinetic Studies, and In Silico Approaches. **2023**, 24, 5216 ○
- 530 Synthesis and Spectroscopic Characterization of Novel Pyridine-based N-acyl Hydrazone Derivatives and Molecular Docking Studies on Glucosamine-6-Phosphate. **2023**, 13, 135-152 ○
- 529 Aging-Accelerated Mouse Prone 8 (SAMP8) Mice Experiment and Network Pharmacological Analysis of Aged Liupao Tea Aqueous Extract in Delaying the Decline Changes of the Body. **2023**, 12, 685 ○
- 528 In silico and in vitro analyses of a novel FoxO1 agonist reducing A β levels via downregulation of BACE1. ○
- 527 Biophysical insights on the interaction of anticoagulant drug dicoumarol with calf thymus-DNA: deciphering the binding mode and binding force with thermodynamics. 1-12 ○
- 526 Anxiolytic effects of N-(4,5-dihydro-5-oxo-1,2-dithiolo-[4,3,b]-pyrrole-6-yl)-N-methylformamide, a pyrroloformamide isolated from a marine Streptomyces sp., in adult zebrafish by the 5-HT system. 1-16 ○
- 525 A new series of IDO1 inhibitors derived from microbial metabolites. **2023**, 54, 76-80 ○
- 524 A novel molecular docking program based on a multi-swarm competitive algorithm. **2023**, 78, 101292 ○
- 523 Genetically engineered PD-1 displaying nanovesicles for synergistic checkpoint blockades and chemo-metabolic therapy against non-small cell lung cancer. **2023**, 161, 184-200 ○
- 522 Bioinformatics and computational chemistry approaches to explore the mechanism of the anti-depressive effect of ligustilide. **2023**, 13, ○
- 521 Atropostatin: Design and Total Synthesis of an Atropisomeric Lactone-Atorvastatin Prodrug. **2023**, 28, 3176 ○
- 520 A consensual machine-learning-assisted QSAR model for effective bioactivity prediction of xanthine oxidase inhibitors using molecular fingerprints. ○
- 519 Towards a better understanding of commonly used medicinal plants from Turkiye: Detailed phytochemical screening and biological activity studies of two Teucrium L. species with in vitro and in silico approach. **2023**, 116482 ○
- 518 A self-reporting photosensitizer for inducing and in-situ monitoring lysosomal damage and cell apoptosis. **2023**, 382, 133482 ○
- 517 Cymbaria daurica L.: A Mongolian herbal medicine for treating eczema via natural killer cell-mediated cytotoxicity pathway. **2023**, 308, 116246 ○
- 516 Synthesis and preliminary in vitro cytotoxic activity of Pd(II) complexes including Salen- or Salphen-Ligands. **2023**, 550, 121450 ○

- 515 Protective effect of *Nardostachys jatamansi* extract against lithium-pilocarpine-induced spontaneous recurrent seizures and associated cardiac irregularities in a rat model. **2023**, 308, 116280 ○
- 514 Computational perspectives on Chlorpyrifos and its degradants as human glutathione S-transferases inhibitors: DFT calculations, molecular docking study and MD simulations. **2023**, 26, 100264 ○
- 513 Proposal of novel ApoE4 inhibitors from the natural spice Cinnamon for the treatment of Alzheimer's disease: Ab initio molecular simulations. **2023**, 296, 106990 ○
- 512 In silico repurposing of CNS drugs for multiple sclerosis. **2023**, 73, 104622 ○
- 511 Korupensamine A, but not its atropisomer, korupensamine B, inhibits SARS-CoV-2 in vitro by targeting its main protease (Mpro). **2023**, 251, 115226 ○
- 510 A microRNA, PC-5p-30_205949, regulates triflumezopyrim susceptibility in *Laodelphax striatellus* (Fallén) by targeting CYP419A1 and ABCG23. **2023**, 192, 105413 ○
- 509 Carboxymuconolactone decarboxylase is a prospective molecular target for multi-drug resistant *Acinetobacter baumannii*-computational modeling, molecular docking and dynamic simulation studies. **2023**, 157, 106793 ○
- 508 Probing the interactions of dansyl appended sequence-defined oligomers with serum albumins: Effect of functionality, hydrophobicity, and architecture. **2023**, 439, 114640 ○
- 507 Vasicinone, a pyrroloquinazoline alkaloid from *Adhatoda vasica* Nees enhances memory and cognition by inhibiting cholinesterases in Alzheimer's disease. **2023**, 3, 100439 ○
- 506 Multi-model in silico characterization of 3-benzamidobenzoic acid derivatives as partial agonists of Farnesoid X receptor in the management of NAFLD. **2023**, 157, 106789 ○
- 505 Astragaloside IV attenuates indoxyl sulfate-induced injury of renal tubular epithelial cells by inhibiting the aryl hydrocarbon receptor pathway. **2023**, 308, 116244 ○
- 504 Synthesis, biochemical characterization and molecular modeling studies of 5-(substituted benzylidene) pyrimidine-2,4,6-trione: Potential inhibitors of alkaline phosphatase. **2023**, 1282, 135225 ○
- 503 DNA-binding affinity and molecular docking studies of the PEGylated binuclear palladacycle, BTC2, an efficient metallodrug against triple-negative breast cancer. **2023**, 243, 112191 ○
- 502 New thieno[2,3-b]pyridine-fused pyrimidin-4(3H)-ones as potential thymidylate synthase inhibitors: Synthesis, SAR, in vitro and in silico study. **2023**, 1282, 135236 ○
- 501 A combined DFT and molecular docking study on novel tricarbonylrhenium(I) complexes bearing mono- and bivalent benzenesulfonamide scaffolds as human carbonic anhydrase IX and XII inhibitors. **2023**, 1282, 135211 ○
- 500 Pharmacological targeting of MTHFD2 suppresses NSCLC via the regulation of ILK signaling pathway. **2023**, 161, 114412 ○
- 499 New anticancer potential Pt complex with tertamyl dithiocarbamate ligand: Synthesis, DNA targeting behavior, molecular dynamic, and biological activity. **2023**, 379, 121651 ○
- 498 Synthesis of benzylidene-indandione derivatives as quantification of amyloid fibrils. **2023**, 296, 106982 ○

- 497 Insights into the inhibitory activity and mechanism of action of flavonoids from the stems and branches of *Acer mono Maxim.* against α -glucosidase via kinetic analysis, molecular docking, and molecular dynamics simulations. **2023**, 1282, 135188 1
- 496 Molecular level solvent interaction (microscopic), electronic, covalent assembly (RDG, AIM & ELF), ADMET prediction and anti-cancer activity of 1-(4-Fluorophenyl)-1-propanone): Cytotoxic agent. **2023**, 380, 121714 0
- 495 N-sulfonyl peptide-hybrids as a new class of dengue virus protease inhibitors. **2023**, 251, 115227 0
- 494 A flaw in applying the FRET technique to evaluate the distance between ligands and tryptophan residues in human serum albumin: Proposal of correction. **2023**, 242, 112693 0
- 493 Biogenic silver NPs alleviate LPS-induced neuroinflammation in a human fetal brain-derived cell line: Molecular switch to the M2 phenotype, modulation of TLR4/MyD88 and Nrf2/HO-1 signaling pathways, and molecular docking analysis. **2023**, 148, 213363 0
- 492 Spectroscopic, electronic properties analysis for 2, 6-Bis (phenylamino)-4-(iminophenyl) benzoquinone molecule and molecular docking clarification for its anticancer activity detected by strong inhibition of NQO1 enzyme. **2023**, 1282, 135231 0
- 491 Synthesis, solvent polarity(polar and nonpolar), structural and electronic properties with diverse solvents and biological studies of (E)-3-((3-chloro-4-fluorophenyl) imino) indolin-2-one. **2023**, 380, 121709 0
- 490 Physicochemical insights into the micellar delivery of doxycycline and minocycline to the carrier protein in aqueous environment. **2023**, 379, 121675 0
- 489 A novel thermostable and salt-tolerant carboxylesterase involved in the initial aerobic degradation pathway for pyrethroids in *Glycomyces salinus*. **2023**, 451, 131128 0
- 488 Bioassay guided isolation of caffeoylquinic acids from the leaves of *Ilex pubescens* Hook. et Arn. and investigation of their anti-influenza mechanism. **2023**, 309, 116322 0
- 487 Multimodal ionic liquid-based chromatographic supports for an effective RNA purification. **2023**, 315, 123676 0
- 486 4-(4-methoxyphenyl)-6-methyl-3-phenyl-4H-1,2,4-oxadiazin-5(6H)-one: Synthesis, crystal structure, Hirshfeld surface analysis, noncovalent, ADMET studies and biological evaluation. **2023**, 1282, 135197 0
- 485 In silico prediction of Antifungal compounds from Natural sources towards Lanosterol 14-alpha demethylase (CYP51) using Molecular docking and Molecular dynamic simulation. **2023**, 121, 108435 0
- 484 Antiproliferative evaluation and supramolecular properties of a Pd(II) complex harvested from benzil bis(pyridyl hydrazone) ligand: Combined experimental and theoretical studies. **2023**, 152, 110646 0
- 483 Voltage dependent anion channel and its interaction with N-acetyl-L-Cysteine (NAC) under oxidative stress on planar lipid bilayer. **2023**, 209, 150-160 0
- 482 Biochemical interaction of human hemoglobin with ionic liquids of noscapinoids: Spectroscopic and computational approach. **2023**, 239, 124227 0
- 481 Ethyl palmitate, an anti-chikungunya virus principle from *Sauropus androgynus*, a medicinal plant used to alleviate fever in ethnomedicine. **2023**, 309, 116366 0
- 480 Design, synthesis and biological evaluation of novel thiosemicarbazones as cruzipain inhibitors. **2023**, 254, 115345 0

- 479 Computational identification of drug-like marine natural products as potential RNA polymerase inhibitors against Nipah virus. **2023**, 104, 107850 ○
- 478 Synthesis, in vitro evaluation and molecular docking studies of hybrid 4-quinolinyl bearing 1,3,4-thiadiazole-2-amine as a new inhibitor of α -amylase and α -glucosidase. **2023**, 1282, 135173 ○
- 477 In silico characterization of the psilocybin biosynthesis pathway. **2023**, 104, 107854 ○
- 476 In silico toxicity studies of traditional Chinese herbal medicine: A mini review. **2023**, 80, 102588 ○
- 475 Design and synthesis of piano-stool ruthenium(II) complexes and their studies on the inhibition of amyloid β (1-42) peptide aggregation. **2023**, 239, 124197 ○
- 474 Assessing protein homology models with docking reproducibility. **2023**, 121, 108430 ○
- 473 Synthesis, characterization, biological activity and molecular docking study of transition metal complexes from heterocyclic ligand system. **2023**, 1282, 135162 ○
- 472 Targeting HER2-receptors with ^{177}Lu -labeled triazole stapled cyclic peptidomimetic. **2023**, 135, 106503 ○
- 471 Exploring the structural, photophysical and optoelectronic properties of a diaryl heptanoid curcumin derivative and identification as a SARS-CoV-2 inhibitor. **2023**, 1281, 135110 ○
- 470 Synthesis of Novel Hybrid Lonidamine-Coumarin Derivatives and Their Anticancer Activities. **2023**, 1281, 135114 ○
- 469 Key essential oil components delocalize *Candida albicans* Kar3p and impact microtubule structure. **2023**, 272, 127373 ○
- 468 Hydrazones tethered disubstituted 1,2,3-triazoles: Design, synthesis, antitubercular and antimicrobial evaluation. **2023**, 1283, 135163 ○
- 467 A novel benzothiophene incorporated Schiff base acting as a turn-on sensor for the selective detection of Serine in organic medium. **2023**, 136, 106525 ○
- 466 Synthesis, characterization and antimicrobial properties of silver complexes derived from 5,6-Dimethylbenzimidazol-2-ylidene. **2023**, 237, 116383 ○
- 465 Experimental and theoretical insights about the effect of some newly designed azomethine group-contained macroheterocycles on oxidative stress and DNA repair gene profiles in neuroblastoma cell lines. **2023**, 1285, 135432 ○
- 464 Synthesis, spectral studies, biological evaluation and molecular docking studies of metal complexes from coumarin derivative. **2023**, 1285, 135443 ○
- 463 5-Oxohexahydroquinolines bearing 4-pyridyl methyl carboxylate as P-glycoprotein inhibitors and multidrug resistance reversal agents in cancer cells. **2023**, 1285, 135427 ○
- 462 The *Haemonchus contortus* LGC-39 subunit is a novel subtype of an acetylcholine-gated chloride channel. **2023**, 22, 20-26 ○

- 461 Design, synthesis, and computational studies of novel imidazo[1,2-a]pyrimidine derivatives as potential dual inhibitors of hACE2 and spike protein for blocking SARS-CoV-2 cell entry. **2023**, 1285, 135525 ○
- 460 Synthesis, in silico studies and investigations on antimicrobial, antimalarial activities of p-toluenesulphonamoyl β -Leu-Gly dipeptide carboxamide derivatives. **2023**, 1285, 135529 ○
- 459 Design, synthesis and molecular docking studies of 5,6-difluoro-1H-benzo[d]imidazole derivatives as effective binders to GABAA receptor with potent anticonvulsant activity. **2023**, 1285, 135502 ○
- 458 The molecular interplay of known phytochemicals as *Culex pipiens* and Rift Valley fever virus inhibitors through molecular docking. **2023**, 30, 103611 ○
- 457 Designing of 2,3-dihydrobenzofuran derivatives as inhibitors of PDE1B using pharmacophore screening, ensemble docking and molecular dynamics approach. **2023**, 159, 106869 ○
- 456 Photophysical study on DNA & BSA binding and cytotoxic behaviour of piperidine-Pt(II) complexes: Their kinetics & mechanism and molecular docking. **2023**, 441, 114740 ○
- 455 Synthesis, crystal structure, Hirshfeld surface analysis, DFT and antihyperglycemic activity of 9-allyl-2,3,9,10a-tetrahydrobenzo[b]cyclopenta[e][1,4]diazepin-10(1H)-one. **2023**, 1283, 135283 ○
- 454 Cellular uptake of metal oxide-based nanocomposites and targeting of chikungunya virus replication protein nsP3. **2023**, 78, 127176 ○
- 453 Synthesis, crystal structure, DFT calculations, and Hirshfeld surface analysis of an NNN pincer type compound. **2023**, 1283, 135252 ○
- 452 Local QSAR modeling of cytotoxic activity of newly designed androstane 3-oximes towards malignant melanoma cells. **2023**, 1283, 135272 ○
- 451 Synthesis, structure and α -glucosidase inhibitor activity evaluation of some acetamide derivatives starting from 2-(naphthalen-1-yl) acetic acid, containing a 1,2,4-triazole. **2023**, 1284, 135321 ○
- 450 Synthesis and evaluation of the antioxidant and anti-tyrosinase activities of thiazolyl hydrazone derivatives and their application in the anti-browning of fresh-cut potato. **2023**, 414, 135745 ○
- 449 Polymerized stimuli-responsive microgels for the removal of organic dye from water. **2023**, 375, 121267 ○
- 448 Design, synthesis, biological evaluation, and docking study of chromone-based phenylhydrazone and benzoylhydrazone derivatives as antidiabetic agents targeting α -glucosidase. **2023**, 132, 106384 ○
- 447 Fluorescence labeling of anchor-modified Mart-1 peptide for increasing its affinity for HLA-A*0201: Hit two targets with one arrow. ○
- 446 Galantamine Based Novel Acetylcholinesterase Enzyme Inhibitors: A Molecular Modeling Design Approach. **2023**, 28, 1035 1
- 445 Propitious Indazole Compounds as β -ketoacyl-ACP Synthase Inhibitors and Mechanisms Unfolded for TB Cure: Integrated Rational Design and MD Simulations. **2023**, 8, ○
- 444 The heptapeptide somatostatin analogue TT-232 exerts analgesic and anti-inflammatory actions via SST4 receptor activation: In silico, in vitro and in vivo evidence in mice. **2023**, 209, 115419 ○

- 443 Design, synthesis and biological evaluation of purine-based derivatives as novel JAK2/BRD4(BD2) dual target inhibitors. **2023**, 132, 106386 ○
- 442 Mechanisms Contributing to Acquired Activated Protein C Resistance in Patients Treated with Thalidomide: A Molecular Dynamics Study. **2022**, 22, 237-244 ○
- 441 Exploration of the anti-inflammatory mechanism of Lanqin oral solution based on the network pharmacology analysis optimized by Q-markers selection. **2023**, 154, 106607 ○
- 440 Discovery of multi-target directed 3-OH pyrrolidine derivatives through a semisynthetic approach from alkaloid vasicine for the treatment of Alzheimer's disease. **2023**, 249, 115145 1
- 439 Molecular targeting of prodigiosin against anti-inflammatory genes cyclooxygenase-1 and -2. **2023**, 126, 260-271 ○
- 438 A facile synthesis of 1,3,4-oxadiazole-based carbamothioate molecules: Antiseizure potential, EEG evaluation and in-silico docking studies. **2023**, 16, 104610 ○
- 437 Onopordum acanthium L. extract attenuates pancreatic β -Cells and cardiac inflammation in streptozocin-induced diabetic rats. **2023**, 18, e0280464 ○
- 436 Anxiolytic and anticonvulsant effect of Ibuprofen derivative through GABAergic neuromodulation in adult Zebrafish. 1-8 ○
- 435 Design, synthesis, docking and mechanistic studies of new thiazolyl/thiazolidinylpyrimidine-2,4-dione antiproliferative agents. **2023**, 16, 104612 1
- 434 Design, synthesis and in vitro biological evaluation of 2-aminopyridine derivatives as novel PI3K α inhibitors for hematological cancer. **2023**, 82, 129152 ○
- 433 Integrated computational and experimental approach for novel anti-leishmanial molecules by targeting Dephospho-coenzyme A kinase. **2023**, 232, 123441 ○
- 432 A computational strategy for therapeutic development against superoxide dismutase (SOD1) amyloid formation: effect of polyphenols on the various events in the aggregation pathway. **2023**, 25, 6232-6246 ○
- 431 Chemical composition and pesticide activity of essential oils from Artemisia annua L. harvested in the rainy and winter seasons. **2023**, 107, 104601 1
- 430 Cyclic 1H-Phospolane Oxides as a Potential Candidate for Cancer Therapy. ○
- 429 Advances in the discovery of new chemotypes through ultra-large library docking. **2023**, 18, 303-313 ○
- 428 A high-resolution β glucosidase inhibition profiling for targeted identification of natural antidiabetic products from Lycopodiella cernua (L.) Pic. Serm and their inhibitory mechanism study. 1-13 ○
- 427 Exploring AlphaFold2's Performance on Predicting Amino Acid Side-Chain Conformations and Its Utility in Crystal Structure Determination of B318L Protein. **2023**, 24, 2740 ○
- 426 Identification of potential drug candidates as TGR5 agonist to combat type II diabetes using in silico docking and molecular dynamics simulation studies. 1-18 ○

- 425 Efficient synthesis of chromeno[2,3-b]pyridine derivatives using Zn(OTf)₂ as a catalyst: DFT computations, molecular docking and ADME studies. **2023**, 375, 121364 ○
- 424 Antimicrobial Activity of Catechol-Containing Biopolymer Poly[3-(3,4-dihydroxyphenyl)glyceric Acid] from Different Medicinal Plants of Boraginaceae Family. **2023**, 12, 285 ○
- 423 Design and synthesis of Diphenyl-1H-imidazole analogs towards SARS CoV-2 3CLpro inhibition for the treatment of COVID-19. ○
- 422 Discovery of novel inhibition site centered on 114-bit tryptophan of Thioredoxin reductase 1 through computer-aided drug design. **2023**, 16, 104642 ○
- 421 Structural, Spectral, Molecular Docking, and Molecular Dynamics Simulations of Phenylthiophene-2-Carboxylate Compounds as Potential Anticancer Agents. 1-23 1
- 420 Synthesis, enzyme inhibition, and molecular docking studies of a novel chalcone series bearing benzothiazole scaffold. ○
- 419 Genome-wide analysis of Catalase gene family reveal insights into abiotic stress response mechanism in Brassica juncea and B. rapa.. **2023**, 330, 111620 ○
- 418 Synthesis, Biological and In Silico Studies of Griseofulvin and Usnic Acid Sulfonamide Derivatives as Fungal, Bacterial and Human Carbonic Anhydrase Inhibitors. **2023**, 24, 2802 1
- 417 Comparative Genomics of Histoplasma capsulatum and Prediction of New Vaccines and Drug Targets. **2023**, 9, 193 1
- 416 Interaction of Positively Charged Oligopeptides with Blood Plasma Proteins. **2023**, 24, 2836 ○
- 415 Molecular surveillance of Kelch-13 gene in Plasmodium falciparum field isolates from Mayurbhanj District, Odisha, India, and in silico artemisinin-Kelch-13 protein interaction study. **2023**, 122, 717-727 ○
- 414 4-Hydroxy Enigmol, a 1-Deoxyphytosphingolipid that Exhibit Good Activity against Prostate and Colon Cancer. **2023**, 8, ○
- 413 New Piperazine Derivatives of 6-Acetyl-7-hydroxy-4-methylcoumarin as 5-HT_{1A} Receptor Agents. **2023**, 24, 2779 ○
- 412 Thioxanthenone-based derivatives as multitarget therapeutic leads for Alzheimer's disease. **2023**, 250, 115169 ○
- 411 Structural Basis for Agonistic Activity and Selectivity toward Melatonin Receptors hMT1 and hMT2. **2023**, 24, 2863 1
- 410 Novel polyvinyl-alcohol microsphere for everolimus delivery for subependymal giant cell astrocytoma. **2023**, 81, 104204 ○
- 409 Structure-function analysis of CYP719As involved in methylenedioxy bridge-formation in the biosynthesis of benzyloquinoline alkaloids and its de novo production. **2023**, 22, ○
- 408 New Glucosamine-Based TLR4 Agonists: Design, Synthesis, Mechanism of Action, and In Vivo Activity as Vaccine Adjuvants. **2023**, 66, 3010-3029 1

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| 407 | Anticancer cytotoxicity of indole-thiazolidinone hybrids, in silico study of mechanisms of their action. 2022 , 38, 257-277 | 0 |
| 406 | Chemical Constituents from <i>Streblus taxoides</i> Wood with Their Antibacterial and Antityrosinase Activities Plus in Silico Study. 2023 , 12, 319 | 0 |
| 405 | The Structural Basis of African Swine Fever Virus pS273R Protease Binding to E64 through Molecular Dynamics Simulations. 2023 , 28, 1435 | 0 |
| 404 | Docking and Molecular Dynamics Simulation Revealed the Potential Inhibitory Activity of Amygdalin in Triple-Negative Breast Cancer Therapeutics Targeting the BRCT Domain of BARD1 Receptor. | 0 |
| 403 | Environmentally-Friendly Pesticidal Activities of <i>Callicarpa</i> and <i>Karomia</i> Essential Oils from Vietnam and Their Microemulsions. 2023 , 20, | 0 |
| 402 | Identification of corticosteroids as potential inhibitor against glycolytic enzyme hexokinase II role in cancer glycolysis pathway: a molecular docking study. 2023 , 36, 173-180 | 0 |
| 401 | Identification of potential novel inhibitors against glutamine synthetase enzyme of <i>Leishmania major</i> by using computational tools. 1-9 | 0 |
| 400 | In silico analysis of low molecular weight sulfated chitosan from <i>Sepia brevimana</i> as potential inhibitors of white spot syndrome envelope proteins. | 0 |
| 399 | Universal Approach to De Novo Drug Design for Target Proteins Using Deep Reinforcement Learning. 2023 , 8, 5464-5474 | 1 |
| 398 | Multiple-P450 Gene Co-Up-Regulation in the Development of Permethrin Resistance in the House Fly, <i>Musca domestica</i> . 2023 , 24, 3170 | 0 |
| 397 | Synthesis, structure elucidation, Hirshfeld surface analysis, DFT, and molecular docking of new 6-bromo-imidazo[4,5-b]pyridine derivatives as potential tyrosyl-tRNA synthetase inhibitors. 1-16 | 0 |
| 396 | Fluorinated benzimidazolium salts: Synthesis, characterization, molecular docking studies and inhibitory properties against some metabolic enzymes. 2023 , 267, 110094 | 1 |
| 395 | Sphingoid Bases Regulate the Sigma-1 Receptor Sphingosine and N,N-Dimethylsphingosine Are Endogenous Agonists. 2023 , 24, 3103 | 0 |
| 394 | RGD and Scutellarin Conjugate (WK001) Targeting Platelet Glycoprotein IIb/IIIa Receptor Protects from Myocardial Ischemia/Reperfusion Injury: Synthesis, Characterization, and Bioactivity Evaluation. 2023 , 34, 477-488 | 0 |
| 393 | AI-accelerated protein-ligand docking for SARS-CoV-2 is 100-fold faster with no significant change in detection. 2023 , 13, | 0 |
| 392 | Integrated use of ligand and structure-based virtual screening, molecular dynamics, free energy calculation and ADME prediction for the identification of potential PTP1B inhibitors. | 0 |
| 391 | Computational modeling of cyanobacterial phytoconstituents against toll-like receptors of skin cancer. 1-13 | 0 |
| 390 | QSAR study of tetrahydropteridin derivatives as polo-like kinase 1 (PLK1) Inhibitors with molecular docking and dynamics study. 2023 , 34, 91-116 | 0 |

- 389 Structural Insight into the Working Mechanism of the FAD Synthetase from the Human Pathogen *Streptococcus pneumoniae*: A Molecular Docking Simulation Study. **2023**, 24, 3121 ○
- 388 High-affinity binding of celastrol to monomeric β -synuclein mitigates in vitro aggregation. 1-11 ○
- 387 Exploring anticancer potential of nintedanib conjugated magnetic nanoparticles: In-vitro and in-silico studies. **2023**, 81, 104213 ○
- 386 Dinuclear Phenoxo-Bridged Nickel(II) and Copper(II) Complexes of Phenolate-Based Tripodal Ligand: Theoretical and Experimental Insights. 1-20 ○
- 385 Targeting *Acanthamoeba* proteins interaction with flavonoids of Propolis extract by in vitro and in silico studies for promising therapeutic effects. 11, 1274 ○
- 384 Phlorizin isolated from seagrass *Syringodium isoetifolium* inhibits diethylnitrosamine and carbon tetrachloride-induced hepatocellular carcinoma in BALB/c mice. **2023**, 155, 1-15 ○
- 383 N-Amidation of Nitrogen-Containing Heterocyclic Compounds: Can We Apply Enzymatic Tools?. **2023**, 10, 222 ○
- 382 Identification of pyrrolizidine alkaloids and flavonoid glycosides through HR-LCMS/MS analysis, biological screening, DFT and molecular docking studies on *Heliotropium dasycarpum* Ledeb.. **2023**, 16, 104655 ○
- 381 Can Resveratrol Influence the Activity of 11 β -Hydroxysteroid Dehydrogenase Type 1? A Combined In Silico and In Vivo Study. **2023**, 16, 251 ○
- 380 Discovery of Highly Potent CRBN Ligands and Insight into Their Binding Mode through Molecular Docking and Molecular Dynamics Simulations. **2023**, 18, ○
- 379 Structure-based virtual screening of novel natural products as chalcone derivatives against SARS-CoV-2 Mpro. 1-15 ○
- 378 Systematic investigation of the multi-scale mechanisms of herbal medicine on treating ventricular remodeling: Theoretical and experimental studies. **2023**, 112, 154706 ○
- 377 In Silico study and design of some new potent threonine tyrosine kinase inhibitors using molecular docking simulation. **2023**, 49, 517-524 ○
- 376 Histidinal-Based Potent Antimalarial Agents. ○
- 375 Mitochondrial Peroxiredoxin 3 Is Rapidly Oxidized and Hyperoxidized by Fatty Acid Hydroperoxides. **2023**, 12, 408 ○
- 374 Computational, Equilibrium, Structural, and Biological Study of the Novel 1-Formyl-4-phenyl-3-semicarbazide and Its Complexes. **2022**, 92, 2708-2722 ○
- 373 Design, Synthesis, and Molecular Docking Study of 6-Aryl-3-(quinolin-3-yl)-7H-[1,2,4]triazolo[3,4-b][1,3,4]thiazidiazines as Novel Antimicrobial Agents. **2022**, 58, 1851-1860 ○
- 372 Alizarin as a potential protector of proteins against damage caused by hydroperoxyl radical. **2023**, 373, 110395 ○

- 371 SS148 and WZ16 inhibit the activities of nsp10-nsp16 complexes from all seven human pathogenic coronaviruses. **2023**, 1867, 130319 ○
- 370 Investigation of the Underlying Mechanism of Huangqi-Dangshen for Myasthenia Gravis Treatment via Molecular Docking and Network Pharmacology. **2023**, 2023, 1-18 ○
- 369 Molecular docking and molecular simulation studies for N-degron selectivity of chloroplastic ClpS from *Chlamydomonas reinhardtii*. **2023**, 103, 107825 ○
- 368 An investigation into solubility and dissolution improvement of alectinib hydrochloride as a third-generation amorphous solid dispersion. **2023**, 81, 104259 ○
- 367 Resveratrol analog, triacetylresveratrol, a potential immunomodulator of lung adenocarcinoma immunotherapy combination therapies. 12, ○
- 366 Cyano-2-oxopyridines: Green synthesis, cytotoxicity evaluation and molecular docking study. **2023**, 299, 127451 ○
- 365 Synergistic Antimicrobial Action of Lactoferrin-Derived Peptides and Quorum Quenching Enzymes. **2023**, 24, 3566 ○
- 364 Challenging breast cancer through novel sulfonamide-pyridine hybrids: design, synthesis, carbonic anhydrase IX inhibition and induction of apoptosis. **2023**, 15, 147-166 ○
- 363 Search for immunomodulatory compounds with antiproliferative activity against melanoma. **2023**, 160, 114374 ○
- 362 Interaction study with DNA/HSA, anti-topoisomerase II β cytotoxicity and in vitro antiproliferative evaluations and molecular docking of indole-thiosemicarbazone compounds. **2023**, 234, 123606 ○
- 361 In vitro efficacy of green synthesized silver nanoconjugates of *Valeriana wallichii* methanolic root extract against MDR strain of *Acinetobacter baumannii*. ○
- 360 Exploring the potential and identifying *Withania somnifera* alkaloids as novel dihydrofolate reductase (DHFR) inhibitors by the AlteQ method. 1-14 ○
- 359 Synthesis of hydrocortisone esters targeting androgen and glucocorticoid receptors in prostate cancer in vitro. **2023**, 229, 106269 ○
- 358 Hope toward potent drug target using tertiary topological instances of Mycobacterial transmembrane protein. **2022**, 10, 69-75 ○
- 357 JFD, a Novel Natural Inhibitor of Keap1 Alkylation, Suppresses Intracellular Mycobacterium Tuberculosis Growth through Keap1/Nrf2/SOD2-Mediated ROS Accumulation. **2023**, 2023, 1-21 ○
- 356 Dockey: a modern integrated tool for large-scale molecular docking and virtual screening. **2023**, 24, 1
- 355 Faster and Protective Wound Healing Mechanistic of Para-Coumaric Acid Loaded Liver ECM Scaffold Cross-linked with Acellular Marine Kelp. 2212325 ○
- 354 In Vitro and In Silico Study on the Impact of Chlorogenic Acid in Colorectal Cancer Cells: Proliferation, Apoptosis, and Interaction with β -Catenin and LRP6. **2023**, 16, 276 ○

- 353 Design, Synthesis, and Metabolism Studies of N-1,4-Diketophenyltriazinones as Protoporphyrinogen IX Oxidase Inhibitors. ○
- 352 Molecular Thermodynamic Origin of Substrate Promiscuity in the Enzyme Laccase: Toward a Broad-Spectrum Degrader of Dye Effluents. **2023**, 14, 1892-1898 ○
- 351 Phenolic compounds as histone deacetylase inhibitors: binding propensity and interaction insights from molecular docking and dynamics simulations. ○
- 350 Illumination of a progressive allosteric mechanism mediating the glycine receptor activation. **2023**, 14, ○
- 349 Identification of a novel adjuvant loperamide that enhances the antibacterial activity of colistin against MCR-1-positive pathogens in vitro/vivo. **2023**, 76, ○
- 348 Identification of Potential Anti-COVID-19 Drug Leads from Medicinal Plants through Virtual High-Throughput Screening. **2023**, 8, ○
- 347 Exploration of the anti-liver injury active components of Shaoyao Gancao decoction by network pharmacology and experiments in vivo. **2023**, 112, 154717 ○
- 346 Effect of sulfonamide derivatives of phenylglycine on scopolamine-induced amnesia in rats. **2023**, 9, 13-31 ○
- 345 The Strategies of Development of New Non-Toxic Inhibitors of Amyloid Formation. **2023**, 24, 3781 ○
- 344 Characterization of the interaction between boscalid and tannic acid and its effect on the antioxidant properties of tannic acid. **2023**, 88, 1325-1335 ○
- 343 Novel Inhibitory Role of Fenofibric Acid by Targeting Cryptic Site on the RBD of SARS-CoV-2. **2023**, 13, 359 ○
- 342 Dectin-1 Acts as a Non-Classical Receptor of Ang II to Induce Cardiac Remodeling. **2023**, 132, 707-722 ○
- 341 Insights into the Degradation of Polymer-Drug Conjugates by an Overexpressed Enzyme in Cancer Cells. **2023**, 66, 2761-2772 ○
- 340 Cytochrome P450 Surface Domains Prevent the β -Carotene Monohydroxylase CYP97H1 of *Euglena gracilis* from Acting as a Dihydroxylase. **2023**, 13, 366 ○
- 339 Exposure to bisphenol A alternatives bisphenol AF and fluorene-9-bisphenol induces gonadal injuries in male zebrafish. **2023**, 253, 114634 ○
- 338 Engineered Glycosidase for Significantly Improved Production of Naturally Rare Vina-Ginsenoside R7. **2023**, 71, 3852-3861 ○
- 337 Pharmacophore Modeling and Molecular Docking of Flavonoid Derivatives in *Abelmoschus manihot* Against Human Estrogen Receptor Alpha of Breast Cancer. **2022**, 1, 1-9 ○
- 336 *Clostridium perfringens* Sialidase Interaction with Neu5Ac β Gal Sialic Acid Receptors by In-Silico Observation and Its Impact on Monolayers Cellular Behaviour Structure. ○

- 335 Rheum rhaponticum and Rheum rhabarbarum Extracts as Modulators of Endothelial Cell Inflammatory Response. **2023**, 15, 949 0
- 334 In Silico Investigation of the Molecular Mechanism of PARP1 Inhibition for the Treatment of BRCA-Deficient Cancers. **2023**, 28, 1829 1
- 333 Binding profile of a mixed-ligand silver(I) complex with DNA and Topoisomerase I. **2023**, 103, 107831 0
- 332 Multicomponent synthesis of new 5-thiourea-4-aza-2,3 didehydropodophyllotoxins as potent cytotoxic agents. 0
- 331 Artificial Intelligence uncovers Evolutionarily Conserved Intracellular Allosteric Modulators of GPCR-G β Interface. 0
- 330 An Insight into Wheat Germ Oil Nutrition, Identification of Its Bioactive Constituents and Computer-Aided Multidimensional Data Analysis of Its Potential Anti-Inflammatory Effect via Molecular Connections. **2023**, 13, 526 2
- 329 How ractopamine binds to bovine serum albumin at the drug site 1. **2023**, 49, 599-607 0
- 328 Chemical Characterization and Leishmanicidal Activity In Vitro and In Silico of Natural Products Obtained from Leaves of Vernonanthur a brasili ana (L.) H. Rob (Asteraceae). **2023**, 13, 285 0
- 327 Toxoplasma gondii aspartic protease 5 (TgASP5): Understanding structural details and inhibition mechanism. 0
- 326 Functionalized Sulfur-Containing Heterocyclic Analogs Induce Sub-G1 Arrest and Apoptotic Cell Death of Laryngeal Carcinoma In Vitro. **2023**, 28, 1856 0
- 325 R97 at Handlebar Binding Mode in Active Pocket Plays an Important Role in Fe(II)/Ketoglutaric Acid-Dependent Dioxygenase cis-P3H-Mediated Selective Synthesis of (2S,3R)-3-Hydroxypip ecolic Acid. **2023**, 28, 1854 0
- 324 Complexation between rice starch and cellulose nanocrystal from black tea residues: Gelatinization properties and digestibility in vitro. **2023**, 234, 123695 0
- 323 Molecular Modes of Action of an Aqueous Nerium oleander Extract in Cancer Cells In Vitro and In Vivo. **2023**, 28, 1871 0
- 322 Study of the competition between Pi and Cr (VI) for the use of Pi-transporter at Vicia faba L. using molecular modeling. **2023**, 196, 695-702 0
- 321 Metallo drugs against Breast Cancer: Combining the Tamoxifen Vector with Platinum(II) and Palladium(II) Complexes. **2023**, 15, 682 0
- 320 Systematic analysis on the mechanism of Zhizi-Bopi decoction against hepatitis B via network pharmacology and molecular docking. **2023**, 45, 463-478 0
- 319 Computational Chemistry for the Identification of Lead Compounds for Radiotracer Development. **2023**, 16, 317 1
- 318 Molecular Screening of Bioactive Compounds of Garlic for Therapeutic Effects against COVID-19. **2023**, 11, 643 0

- 317 Therapeutic Effect of Costunolide in Autoimmune Hepatitis: Network Pharmacology and Experimental Validation. **2023**, 16, 316 ○
- 316 Selective Targeting of Cancer-Related G-Quadruplex Structures by the Natural Compound Dicentrine. **2023**, 24, 4070 ○
- 315 Investigation of autism-related transcription factors underlying sex differences in the effects of bisphenol A on transcriptome profiles and synaptogenesis in the offspring hippocampus. **2023**, 14, 1 ○
- 314 In silico analysis of dietary polyphenols and their gut microbial metabolites suggest inhibition of SARS-CoV-2 infection, replication, and host inflammatory mediators. 1-19 ○
- 313 Pyrazolones Potentiate Colistin Activity against MCR-1-Producing Resistant Bacteria: Computational and Microbiological Study. **2023**, 8, 8366-8376 ○
- 312 Antifungal Activity, Structure-Activity Relationship and Molecular Docking Studies of 1,2,4-Triazole Schiff Base Derivatives. **2023**, 20, ○
- 311 Exploring the Binding Mechanism of Novel T. Brucei Leucine tRNA Synthase Inhibitors Based on QSAR Modeling, Molecular Docking, ADMET Prediction and Molecular Dynamics Simulations . ○
- 310 All-Atom Molecular Dynamics Simulations Indicated the Involvement of a Conserved Polar Signaling Channel in the Activation Mechanism of the Type I Cannabinoid Receptor. **2023**, 24, 4232 ○
- 309 Identification of and Mechanistic Insights into SARS-CoV-2 Main Protease Non-Covalent Inhibitors: An In-Silico Study. **2023**, 24, 4237 ○
- 308 A Simple Way to Incorporate Target Structural Information in Molecular Generative Models. ○
- 307 GalaxyDock2-HEME : P rotein ligand docking for heme proteins. ○
- 306 Exploring quantum computational, molecular docking, and molecular dynamics simulation with MMGBSA studies of ethyl-2-amino-4-methyl thiophene-3-carboxylate. 1-19 ○
- 305 Coumarin linked thiazole derivatives as potential αGlucosidase inhibitors to treat Diabetes Mellitus. ○
- 304 Molecular Docking and In Vitro Studies of Ochratoxin A (OTA) Biodetoxification Testing Three Endopeptidases. **2023**, 28, 2019 ○
- 303 Liquid Crystal Monomer: A Potential PPAR α Antagonist. **2023**, 57, 3758-3771 ○
- 302 Identification and Functional Characterization of General Odorant Binding Proteins in *Orthaga achatina*. **2023**, 14, 216 ○
- 301 Complex chemical signals dictate Ah receptor activation through the gut-lung axis. ○
- 300 Venetoclax analogs as promising anticancer therapeutics via targeting Bcl-2 protein: in-silico drug discovery study. 1-17 ○

- 299 Molecular and Biochemical Characterization of Field Resistant Isolates of *Glomerella cingulata* to Pyraclostrobin in China. **2023**, 71, 3960-3966 ○
- 298 Anastrozole-mediated modulation of mitochondrial activity by inhibition of mitochondrial permeability transition pore opening: an initial perspective. 1-17 ○
- 297 Isolation, characterization and in silico study of propenamide alkaloids from *Hymenopmecis bicolor* poison against active μ opioid receptor. 1-17 ○
- 296 Sterol Structural Features Impact on the Spontaneous Membrane Insertion of CLIC1 into Artificial Lipid Membranes. **2023**, 39, 3286-3300 ○
- 295 Potential diagnostic markers and therapeutic targets for rheumatoid arthritis with comorbid depression based on bioinformatics analysis. 14, ○
- 294 Lessons Learnt from COVID-19: Computational Strategies for Facing Present and Future Pandemics. **2023**, 24, 4401 ○
- 293 A Comprehensive Molecular and Clinical Investigation of Approved Anti-HCV Drugs Repurposing against SARS-CoV-2 Infection: A Glaring Gap between Benchside and Bedside Medicine. **2023**, 11, 515 ○
- 292 Eaton's reagent is an alternative of PPA: Solvent free synthesis, molecular docking and ADME studies of new angular and linear carbazole based naphtho naphthyridines. **2023**, 135, 133320 ○
- 291 Identification of novel C-15 fluoro isosteviol derivatives for GABA-AT inhibition by in silico investigations. **2023**, 29, ○
- 290 Machine learning combines atomistic simulations to predict SARS-CoV-2 Mpro inhibitors from natural compounds. ○
- 289 Investigating a Library of Flavonoids as Potential Inhibitors of a Cancer Therapeutic Target MEK2 Using in Silico Methods. **2023**, 24, 4446 ○
- 288 Dihydrokaempferol attenuates CCl₄-induced hepatic fibrosis by inhibiting PARP-1 to affect multiple downstream pathways and cytokines. **2023**, 464, 116438 ○
- 287 In silico target specific design of potential quinazoline-based anti-NSCLC agents. 1-12 ○
- 286 Drug repurposing and molecular mechanisms of the antihypertensive drug candesartan as a TMEM16A channel inhibitor. **2023**, 235, 123839 ○
- 285 A comparative study on DNA and protein binding properties of thymol and thymoquinone. 1-13 ○
- 284 Discovery of potential quality markers of *Fritillariae thunbergii* bulbus in pneumonia by combining UPLC-QTOF-MS, network pharmacology, and molecular docking. ○
- 283 Insights into the binding mechanism of ascorbic acid and violaxanthin with violaxanthin de-epoxidase (VDE) and chlorophycean violaxanthin de-epoxidase (CVDE) enzymes. ○
- 282 5,8-Dimethyl-9H-carbazole Derivatives Blocking hTopo I Activity and Actin Dynamics. **2023**, 16, 353 ○

- 281 Synthesis, Reactions, and Antioxidant Properties of Bis(3-amino-1-hydroxybenzyl)diselenide. **2023**, 88, 3509-3522 ○
- 280 Docking strategies. **2023**, 243-258 ○
- 279 Comparative Assessment of Docking Programs for Docking and Virtual Screening of Ribosomal Oxazolidinone Antibacterial Agents. **2023**, 12, 463 ○
- 278 New Insight into the Pharmacological Importance of Atropine as the Potential Inhibitor of AKR1B1 via Detailed Computational Investigations: DFTs, ADMET, Molecular Docking, and Molecular Dynamics Studies. ○
- 277 Molecular docking and dynamic approach to screen the drug candidate against the Imipenem-resistant CarO porin in *Acinetobacter baumannii*. **2023**, 177, 106049 ○
- 276 The molecular mechanism of Y473 phosphorylation of UGDH relieves the inhibition effect of UDP-glucose on HuR. **2023**, 25, 8714-8724 ○
- 275 Facile, Single-Step Synthesis of a Series of D-Ring Ethisterones Substituted with 1,4-1,2,3-Triazoles: Preliminary Evaluation of Cytotoxic Activities. **2023**, 18, ○
- 274 β -Pinene: Docking Study, Cytotoxicity, Mechanism of Action, and Anti-Biofilm Effect against *Candida albicans*. **2023**, 12, 480 ○
- 273 In-vitro antigout potential of *Alstonia scholaris* flower, characterization and prospective ligand-receptor interaction of bioactive lead compound. **2023**, 9, e14093 ○
- 272 In-silico natural product database mining for novel neuropilin-1 inhibitors: molecular docking, molecular dynamics and binding energy computations. **2023**, 17, ○
- 271 The antiphage defense system CBASS controls resistance and enables killing by antifolate antibiotics in *Vibrio cholerae*. ○
- 270 Synthesis, cytotoxicity, and docking based analysis of acridone-N-acetamides as AKT kinase inhibitors. ○
- 269 Exploring the Interaction Between the Newly Designed Antitumor Zn(II) Complex and CT-DNA/BSA: Spectroscopic Methods, DFT Computational Analysis, and Docking Simulation. ○
- 268 The Importance of Epigallocatechin as a Scaffold for Drug Development against Flaviviruses. **2023**, 15, 803 ○
- 267 IN SILICO LIBRARY SCREENING TO FIND NOVEL ANTICANCER AGENT WITH CHEMOSENSITIZING PROPERTIES: FOCUS ON TARGETING ATAXIA TELANGIECTASIA AND Rad3 RELATED KINASE. ○
- 266 Computational Docking Study of the Phytochemical Constituent, Silybin (*Silybum marianum*) against SARS-CoV-2 Omicron Variant Spike Glycoprotein: An In-silico Approach. **2023**, 17, 385-394 ○
- 265 In silico investigation of binding propensity of hematoxylin derivative and damnacanthal for their potential inhibitory effect on HIV-1 Vpr from different subtypes. 1-12 ○
- 264 Design, synthesis, and biological evaluation of new 2-(4-(methylsulfonyl)phenyl)-N-phenylimidazo[1,2-a]pyridin-3-amine as selective COX-2 inhibitors. **2023**, 32, 856-868 ○

- 263 PAMAM-G4 protect the N-(2-hydroxyphenyl)-2-propylpentanamide (HO-AAVPA) and maintain its antiproliferative effects on MCF-7. **2023**, 13, ○
- 262 Evaluation of antioxidant potentials and acetylcholinesterase inhibitory effects of some new salicylic acid-salicylamide hybrids. ○
- 261 Analyzing the Impact of the Substituent on the Quinazolinone Schiff Base and the Interaction of the Fe (III) and Cr (III) with Different Quinazolinone Schiff Base for Antioxidant and Anti-inflammatory Activity. **2023**, 13, 39-59 ○
- 260 Computational and experimental studies of salvianolic acid A targets 3C protease to inhibit enterovirus 71 infection. 14, ○
- 259 Experimental spectroscopic and quantum computational investigation and molecular docking analysis of N-benzyloxycarbonyl-L-serine as an anticancer agent. **2023**, 56, 147-165 ○
- 258 Therapeutic capability of selected medicinal plants' bioactive constituents against the mutant ovarian TP53 gene; a computational approach. **2023**, 5, 8-32 ○
- 257 LC/MS/MS Phenolic Profile, Antioxidant Capacity, Angiotensin-Converting Enzyme (ACE) and Glutathione S Transferase (GST) Inhibitory Properties of Ultrasound-Assisted Propolis Extracts. ○
- 256 Structural Characterization of the Chlorophyllide a Oxygenase (CAO) Enzyme Through an In Silico Approach. **2023**, 91, 225-235 ○
- 255 Effects of synonymous mutations on kinetic properties and structure of firefly luciferase: Molecular dynamics simulation, molecular docking, RNA folding, and experimental study. **2023**, 235, 123835 ○
- 254 Insights into the effect of protein glycosylation on carbohydrate substrate binding. **2023**, 235, 123833 ○
- 253 Adding New Scientific Evidences on the Pharmaceutical Properties of Pelargonium quercetorum Agnew Extracts by Using In Vitro and In Silico Approaches. **2023**, 12, 1132 1
- 252 SCARDock: A Web Server and Manually Curated Resource for Discovering Covalent Ligands. **2023**, 8, 10397-10402
- 251 Structure modelling of odorant receptor from Aedes aegypti and identification of potential repellent molecules. **2023**, 21, 2204-2214 ○
- 250 In Silico Exploration of Microtubule Agent Griseofulvin and Its Derivatives Interactions with Different Human β -Tubulin Isoforms. **2023**, 28, 2384 ○
- 249 9-Butyl-Harmol Exerts Antiviral Activity against Newcastle Disease Virus through Targeting GSK-3 β and HSP90 α **2023**, 97, ○
- 248 Targeting Sterylglucosidase A to Treat Aspergillus fumigatus Infections. ○
- 247 An Impedimetric Biosensor for Detection of Volatile Organic Compounds in Food. **2023**, 13, 341 ○
- 246 Inducible Gut-Specific Carboxylesterase SICOE030 in Polyphagous Pests of Spodoptera litura Conferring Tolerance between Nicotine and Cyantraniliprole. **2023**, 71, 4281-4291 ○

- 245 The unique three-dimensional arrangement of Macrophage Galactose Lectin enables E. coli LipoPolySaccharides recognition through two distinct interfaces. ○
- 244 Interaction of the nanobio-based reagent with sodium fluorescein and lipids via bioinformatics for forensic fingerprint visualisations. 1-8 ○
- 243 Discovery of 2H-Indazole-3-carboxamide Derivatives as Novel Potent Prostanoid EP4 Receptor Antagonists for Colorectal Cancer Immunotherapy. ○
- 242 Synthesis, In Vitro Biological Evaluation of Antiproliferative and Neuroprotective Effects and In Silico Studies of Novel 16E-Arylidene-5 β -Epoxyepiandrosterone Derivatives. **2023**, 11, 812 ○
- 241 Functional identification of two novel carbohydrate-binding modules of glucuronoxylanase CrXyl30 and their contribution to the lignocellulose saccharification. **2023**, 16, ○
- 240 Potential drug candidates as P-glycoprotein inhibitors to reverse multidrug resistance in cancer: an in silico drug discovery study. 1-16 ○
- 239 In silico Rivaroxaban binding affinity to β -ketoacyl[ACP]synthase I: search for new pharmacophore. **2023**, 59, 42-48 ○
- 238 Terpenic Constituents of Essential Oils with Larvicidal Activity against Aedes Aegypti: A QSAR and Docking Molecular Study. **2023**, 28, 2454 ○
- 237 Assessment of Biological Activities and Genomic Changes on Microorganisms, Wheat, and Wilding Arise from Poly(pyrazole). ○
- 236 Genetic algorithm-de novo, molecular dynamics and MMGBSA based modelling of a novel Benz-pyrazole based anticancer ligand to functionally revert mutant P53 into wild type P53. **2023**, 49, 678-689 ○
- 235 Interaction of tacrine-coumarin derivatives with double stranded DNA: spectroscopic and molecular docking study. ○
- 234 DFT calculations, spectroscopic investigations, and molecular docking study of Methylprednisolone with some selective cancer proteins. **2023**, ○
- 233 Chemical Characterization and Multidirectional Biological Effects of Different Solvent Extracts of *Arum elongatum* : in Vitro and in Silico Approaches. ○
- 232 Insight into the conserved structural dynamics of the C-terminus of mammal PrPC identifies structural core and possible structural role of pharmacological chaperones. **2023**, 17, 55-66 ○
- 231 Purification, Identification, and Mechanistic Investigation of Novel Selenium-Enriched Antioxidant Peptides from *Moringa oleifera* Seeds. **2023**, 71, 4625-4637 ○
- 230 C-N bond formation by a polyketide synthase. **2023**, 14, ○
- 229 Resistance Risk Assessment for the New OSBP Inhibitor Y18501 in *Pseudoperonospora cubensis* and Point Mutations (G705V, L798W, and I812F) in PscORP1 that Confer Resistance. **2023**, 71, 4510-4520 ○
- 228 Investigation of Antimicrobial, Anti-Quorum Sensing, and Cytotoxic Activities of Flavonoids Isolated from *Pulicaria armena* Boiss. & Kotschy ex Boiss. (Asteraceae). ○

- 227 Piano-stool ruthenium(ii) complexes with maleimide and phosphine or phosphite ligands: synthesis and activity against normal and cancer cells. **2023**, 52, 4237-4250 ○
- 226 Metformin May Alter the Metabolic Reprogramming in Cancer Cells by Disrupting the L-Arginine Metabolism: A Preliminary Computational Study. **2023**, 24, 5316 ○
- 225 Antimicrobial Peptides Designed against the β -Loop of Class A β -Lactamases to Potentiate the Efficacy of β -Lactam Antibiotics. **2023**, 12, 553 ○
- 224 Structural and functional characterization of the cytotoxic protein ledodin, an atypical ribosome-inactivating protein from shiitake mushroom (*Lentinula edodes*). **2023**, 32, ○
- 223 Advances and critical assessment of machine learning techniques for prediction of docking scores. ○
- 222 Computational insight into structural basis of human ELOVL1 inhibition. **2023**, 157, 106786 ○
- 221 Probing mitochondrial damage using a fluorescent probe with mitochondria-to-nucleolus translocation. **2023**, 108323 ○
- 220 A bivalent β -carboline derivative inhibits macropinocytosis-dependent entry of pseudorabies virus by targeting the kinase DYRK1A. **2023**, 299, 104605 ○
- 219 Cytotoxic Activity of Amaryllidaceae Plants against Cancer Cells: Biotechnological, In Vitro, and In Silico Approaches. **2023**, 28, 2601 ○
- 218 In Silico and In Vitro Approach for Validating the Inhibition of Matrix Metalloproteinase-9 by Quercetin. **2023**, Publish Ahead of Print, ○
- 217 HuR modulation counteracts lipopolysaccharide response in murine macrophages. **2023**, 16, ○
- 216 Identification of novel potential inhibitor of thymidylate kinase from Variola virus. 1-11 ○
- 215 A Study on the Interactions of Proteinase K with Myricetin and Myricitrin by Multi-Spectroscopy and Molecular Modeling. **2023**, 24, 5317 ○
- 214 Exploring Probenecid Derived 1,3,4-Oxadiazole-Phthalimide Hybrid as α -Amylase Inhibitor: Synthesis, Structural Investigation, and Molecular Modeling. **2023**, 16, 424 ○
- 213 Spectroscopic and Theoretical Analysis of the Interaction between Plasma Proteins and Phthalimide Analogs with Potential Medical Application. **2023**, 13, 760 ○
- 212 Performance of Eversa Transform 2.0 Lipase in Ester Production Using Babassu Oil (*Orbignya sp.*) and Tucuman Oil (*Astrocaryum vulgare*): A Comparative Study between Liquid and Immobilized Forms in Fe₃O₄ Nanoparticles. **2023**, 13, 571 1
- 211 A comparative analysis of daunorubicin and its metabolite daunorubicinol interaction with apoptotic and drug resistance proteins using in silico approach. 1-13 ○
- 210 Computational Analysis and Experimental Testing of the Molecular Mode of Action of Gatastatin and Its Derivatives. **2023**, 15, 1714 ○

- 209 In silico study of interaction of (ZnO)₁₂ nanocluster to glucose oxidase-FAD in absence and presence of glucose. 1-9 ○
- 208 Bioinformatics Approaches Applied to the Discovery of Antifungal Peptides. **2023**, 12, 566 ○
- 207 In Silico Methodologies to Improve Antioxidants Characterization from Marine Organisms. **2023**, 12, 710 ○
- 206 Effect of in vitro simulated digestion on the anti-Helicobacter Pylori activity of different Propolis extracts. **2023**, 38, ○
- 205 Revisiting the inhibitory potential of protein kinase inhibitors against NEK7 protein via comprehensive computational investigations. **2023**, 13, ○
- 204 Quinoline-Malonitrile-Based Aggregation-Induced Emission Probe for Monoamine Oxidase Detection in Living Cells. **2023**, 28, 2655 ○
- 203 Disaggregation-induced resurgence of quenched emission of a self-assembled bis-indole system: photophysics, energetics, and dynamics. **2023**, 25, 10166-10174 ○
- 202 Virtual Screening of Novel 24-Dehydroxysterol Reductase (DHCR24) Inhibitors and the Biological Evaluation of Irbesartan in Cholesterol-Lowering Effect. **2023**, 28, 2643 ○
- 201 High throughput screening of a new fluorescent G-quadruplex ligand having telomerase inhibitory activity in human A549 cells. 1-22 ○
- 200 Identification of small molecule antivirals against HTLV-1 by targeting the hDLG1-Tax-1 protein-protein interaction. ○
- 199 Development of machine learning models based on molecular fingerprints for selection of small molecule inhibitors against JAK2 protein. ○
- 198 The role of RNA methyltransferase METTL3 in gynecologic cancers: Results and mechanisms. 14, 1
- 197 Identification of novel inhibitors of tetranectin-plasminogen interaction to suppress breast cancer invasion: an integrated computational and cell-based investigation. 1-10 ○
- 196 Molecular docking of 5-o-benzoylpinostrobin derivatives from boesenbergia pandurata roxb. as anti-inflammatory. ○
- 195 Synthesis of vanillin derivatives with 1,2,3-triazole fragments and evaluation of their fungicide and fungistatic activities. ○
- 194 New Fluoroquinolone-1,2,4-triazoles as Potent Antibacterial Agents: Design, Synthesis, Docking Studies and in Silico ADME Profiles. ○
- 193 Molecular Structure, Spectral Analysis, Molecular Docking and Physicochemical Studies of 3-Bromo-2-hydroxypyridine Monomer and Dimer as Bromodomain Inhibitors. **2023**, 28, 2669 ○
- 192 Hypoglycemic effect of Coffea arabica leaf extracts and major bioactive constituents. 1-16 ○

- 191 Dirigent gene editing of gossypol enantiomers for toxicity-depleted cotton seeds. **2023**, 9, 605-615 ○
- 190 Novel Steroidal[17,16-d]pyrimidines Derived from Epiandrosterone and Androsterone: Synthesis, Characterization and Configuration-Activity Relationships. **2023**, 28, 2691 ○
- 189 Exploring the Potential of Black Soldier Fly Larval Proteins as Bioactive Peptide Sources through in Silico Gastrointestinal Proteolysis: A Cheminformatic Investigation. **2023**, 13, 605 ○
- 188 Eco-friendly Synthesis of Schiff base Mn(II) and Cu(II) Complexes: Antimicrobial, Antifungal, Molecular Docking and Anticancer Studies. ○
- 187 MM/GBSA prediction of relative binding affinities of carbonic anhydrase inhibitors: effect of atomic charges and comparison with Autodock4Zn. **2023**, 37, 167-182 ○
- 186 Green products in the management of stored food grains: Challenges, recent advances and future prospects. **2023**, 1-27 ○
- 185 Bio-efficacy of insecticidal molecule emodin against dengue, filariasis, and malaria vectors. ○
- 184 Human Gut Microbial Sulfonolipids are linked to Inflammatory Bowel Diseases through Toll-Like Receptor 4 Signaling. ○
- 183 Interactions of N-Mannich Bases of Pyrrolo[3,4-c]pyrrole with Artificial Models of Cell Membranes and Plasma Proteins, Evaluation of Anti-Inflammatory and Antioxidant Activity. **2023**, 13, 349 ○
- 182 Phytomolecules as potential candidates to intervene the function of E. coli sodium-proton antiporters; Ec-NhaA. 1-12 ○
- 181 In Silico Exploration of Metabolically Active Peptides as Potential Therapeutic Agents against Amyotrophic Lateral Sclerosis. **2023**, 24, 5828 ○
- 180 Ze-Qi decoction inhibits non-small cell lung cancer growth and metastasis by modulating the PI3K/Akt/p53 signaling pathway. **2023**, ○
- 179 Comprehensive Evaluation of End-Point Free Energy Techniques in Carboxylated-Pillar[6]arene Host-Guest Binding: III. Force-Field Comparison, Three-Trajectory Realization and Further Dielectric Augmentation. **2023**, 28, 2767 ○
- 178 Pharmacological Chaperones and Protein Conformational Diseases: Approaches of Computational Structural Biology. **2023**, 24, 5819 ○
- 177 New Thienobenzonaphthotriazoles as Butyrylcholinesterase Inhibitors: Design, Synthesis and Computational Study. **2023**, 24, 5879 ○
- 176 Identification of peptide epitopes of the gp120 protein of HIV-1 capable of inducing cellular and humoral immunity. **2023**, 13, 9078-9090 ○
- 175 Immunoinformatics aided approach for predicting potent cytotoxic T cell epitopes of respiratory syncytial virus. 1-13 ○
- 174 G2/M-Phase-Inhibitory Mitochondrial-Depolarizing Re(I)/Ru(II)/Ir(III)-2,2'-Bipyrimidine-Based Heterobimetallic Luminescent Complexes: An Assessment of In Vitro Antiproliferative Activity and Bioimaging for Targeted Therapy toward Human TNBC Cells. **2023**, 8, 12283-12297 ○

- 173 Inhibition of α -glucosidase enzyme by click'-inspired pharmacophore framework 1,3,4-thiadiazole π ,2,3-triazole hybrids. **2023**, 15, 345-363 ○
- 172 Elucidating the molecular mechanisms of essential oils' insecticidal action using a novel cheminformatics protocol. **2023**, 13, ○
- 171 Synthesis, DFT calculations, α -glucosidase inhibitor activity, and docking studies on Schiff base metal complexes containing isothiocyanate. **2023**, 37, ○
- 170 Piperazine- and Pyrazole-Based Heterocyclic Scaffold Derivatives Connected with Urea and Thiourea for Anti-Inflammatory Activity. **2023**, 8, ○
- 169 Synthesis, Characterization and Structural Studies of Novel Pyrazoline Derivatives as Potential Inhibitors of NAD⁺ Synthetase in Bacteria and Cytochrome P450 51 in Fungi. **2023**, 8, ○
- 168 Batch and continuous flow asymmetric synthesis of anabolic-androgenic steroids via a single-cell biocatalytic α -dehydrogenation and C17 π -carbonyl reduction cascade. ○
- 167 Rigid Scaffolds are Promising for Designing Macrocyclic Kinase Inhibitors. ○
- 166 Synthesis, characterization, docking and antimicrobial studies of binol based amide linked symmetrical bistriazoles. **2023**, 100, 100973 ○
- 165 Combination of furosemide, gold, and dopamine as a potential therapy for breast cancer. **2023**, 23, ○
- 164 Efficient screening of protein-ligand complexes in lipid bilayers using LoCoMock score. **2023**, 37, 217-225 ○
- 163 Bioinformatics-based investigation on the genetic influence between SARS-CoV-2 infections and idiopathic pulmonary fibrosis (IPF) diseases, and drug repurposing. **2023**, 13, ○
- 162 Computer-Aided Drug Design and Synthesis of Rhenium Clotrimazole Antimicrobial Agents. **2023**, 12, 619 ○
- 161 Purification and activity evaluation of novel anti-inflammatory peptides from pearl oyster (*Pinctada martensii*) hydrolysates. ○
- 160 Drug repurposing for Mpox: Discovery of small molecules as potential inhibitors against DNA-dependent RNA polymerase using molecular modeling approach. ○
- 159 Structural rationale to understand the effect of disease-associated mutations on Myotubularin. **2023**, 5, 100100 ○
- 158 Conformationally Restricted β Receptor Antagonists from β -Isopulegol. **2023**, 66, 4999-5020 ○
- 157 Impaired brain equanimity and neurogenesis in the diet-induced overweight mouse: a preventive role by syringic acid treatment. 1-18 ○
- 156 Restriction of access to the central cavity is a major contributor to substrate selectivity in plant ABCG transporters. **2023**, 80, ○

- 155 Unraveling the Aquaporin-3 Inhibitory Effect of Rottlerin by Experimental and Computational Approaches. **2023**, 24, 6004 ○
- 154 Intermolecular And Dynamic Investigation of The Mechanism of Action of Reldesemtiv on Fast Skeletal Muscle Troponin Complex Toward the Treatment of Impaired Muscle Function. ○
- 153 Some Nanocarrier's Properties and Chemical Interaction Mechanisms with Flavones. **2023**, 28, 2864 ○
- 152 Non-covalent interaction, adsorption characteristics and solvent effect of procainamide anti-arrhythmias drug on silver and gold loaded silica surfaces: SERS spectroscopy, density functional theory and molecular docking investigations. **2023**, 13, 9539-9554 ○
- 151 Pyrrole-Based Schiff-Bases: Synthesis, Fluorescent Properties, Molecular Docking and in silico ADME/Tox Profiling Studies. **2023**, 8, ○
- 150 Docking experiments suggest that gloriosine has microtubule-targeting properties similar to colchicine. **2023**, 13, ○
- 149 Identification of Natural Compounds of the Apple as Inhibitors against Cholinesterase for the Treatment of Alzheimer's Disease: An In Silico Molecular Docking Simulation and ADMET Study. **2023**, 15, 1579 ○
- 148 Flipping the Substrate Creates a Highly Selective Halohydrin Dehalogenase for the Synthesis of Chiral 4-Aryl-2-oxazolidinones from Readily Available Epoxides. **2023**, 13, 4768-4777 ○
- 147 The 8-bromobaicalein inhibited the replication of dengue, and Zika viruses and targeted the dengue polymerase. **2023**, 13, ○
- 146 Surface-up click access to allylimidazolium bridged cyclodextrin dimer phase for efficient enantioseparation. ○
- 145 Antibacterial activity of a novel compound isolated from *Bacillus licheniformis* for treating bacterial infections in fishes: An in-silico approach. ○
- 144 Visual dynamics: a WEB application for molecular dynamics simulation using GROMACS. **2023**, 24, ○
- 143 *Candida glycerinogenes*-Promoted β -Pinene and Squalene Co-production Strategy Based on β -Pinene Stress. **2023**, 71, 5250-5260 ○
- 142 Targeting *Acanthamoeba* proteins interaction with flavonoids of Propolis extract by in vitro and in silico studies for promising therapeutic effects. 11, 1274 ○
- 141 Docking cholesterol to integral membrane proteins with Rosetta. **2023**, 19, e1010947 ○
- 140 Detailed Experimental and In Silico Investigation of Indomethacin Binding with Human Serum Albumin Considering Primary and Secondary Binding Sites. **2023**, 28, 2979 ○
- 139 Site-directed mutagenesis identified the key active site residues of 2,3-oxidosqualene cyclase HcOSC6 responsible for cucurbitacins biosynthesis in *Hemsleya chinensis*. 14, ○
- 138 Pharmacological Mechanism of *Aucklandia Radix* against Gastric Ulcer Based on Network Pharmacology and In Vivo Experiment. **2023**, 59, 666 ○

- 137 Potent and selective covalent inhibition of the papain-like protease from SARS-CoV-2. **2023**, 14, ○
- 136 Docking and Molecular Dynamics Identify Leads against 5 Alpha Reductase 2 for Benign Prostate Hyperplasia Treatment. **2023**, 2023, 1-20 ○
- 135 SARS-CoV2 billion-compound docking. **2023**, 10, ○
- 134 Investigation of the Entry Pathway and Molecular Nature of β Receptor Ligands. **2023**, 24, 6367 ○
- 133 New insight of chemical constituents in *Persea americana* fruit against obesity: A data-driven study. ○
- 132 Binding mechanism of andrographolide with intramolecular antiparallel G-quadruplexes of therapeutic importance: an in-silico analysis. **2023**, 49, 816-828 ○
- 131 Identification of a Putative β -Synuclein Radioligand Using an in silico Similarity Search. ○
- 130 Targeting Shikimate Kinase Pathway of *Acinetobacter baumannii*: A Structure-Based Computational Approach to Identify Antibacterial Compounds. **2023**, 2023, 1-14 ○
- 129 Screening of Prospective Antiallergic Compound as Fc ϵ R1 Inhibitors and Its Antiallergic Efficacy Through Immunoinformatics Approaches. ○
- 128 Inhibition Analysis and High-Resolution Crystal Structure of *Mus musculus* Glutathione Transferase P1-1. **2023**, 13, 613 ○
- 127 Phytochemical Profiling, Biological Activities, and In Silico Molecular Docking Studies of *Causonis trifolia* (L.) Mabb. & J.Wen Shoot. **2023**, 12, 1495 ○
- 126 Identification of potential inhibitor against CTX-M-3 and CTX-M-15 proteins: an in silico and in vitro study. 1-17 ○
- 125 Synthesis, DFT Calculation, DNA Binding, and Biological Evaluation of Some Mononuclear Ru(III) Complexes with 2,6-Bis(2-benzimidazolyl)pyridine Bearing Different p-Substituted Heterochalcones. **2023**, 93, 375-388 ○
- 124 Effect of temperature on hepatitis a virus and exploration of binding mode mechanism of phytochemicals from *tinospora cordifolia*: an insight into molecular docking, MM/GBSA, and molecular dynamics simulation study. 1-17 ○
- 123 Spectroscopy, docking and molecular dynamics studies on the interaction between cis and trans palladium-alanine complexes with calf-thymus DNA and antitumor activities. 1-24 ○
- 122 4,5-Dihydro-5-Oxo-Pyrazolo[1,5-a]Thieno[2,3-c]Pyrimidine: A Novel Scaffold Containing Thiophene Ring. Chemical Reactivity and In Silico Studies to Predict the Profile to GABAA Receptor Subtype. **2023**, 28, 3054 1
- 121 Molecular docking and dynamics simulation of *Orthosiphon stamineus* against SGLT1 and SGLT2. 1-16 ○
- 120 Discovery of *Mycobacterium tuberculosis* CYP121 New Inhibitor via Structure-based Drug Repurposing. **2023**, 31, 1503-1521 ○

- 119 Curcumin-Based Heterocycles: Synthesis, Antimicrobial Genotoxicity and Molecular Docking. **2023**, 35, 951-960 ○
- 118 Synthesis of Novel 2,9-Disubstituted-6-morpholino Purine Derivatives Assisted by Virtual Screening and Modelling of Class I PI3K Isoforms. **2023**, 15, 1703 ○
- 117 Synthesis, crystal structure, DFT/HF, Hirshfeld surface, and molecular docking analysis of 4-(tert-butyl)-4-nitro-1,1-biphenyl. **2023**, 14, 90-98 ○
- 116 Orphan receptor GPR158 serves as a metabotropic glycine receptor: mGlyR. **2023**, 379, 1352-1358 ○
- 115 A network pharmacology and molecular docking approach in the exploratory investigation of the biological mechanisms of lagundi (*Vitex negundo* L.) compounds against COVID-19. **2023**, 21, e4 ○
- 114 Large-scale Annotation of Biochemically Relevant Pockets and Tunnels in Cognate Enzyme-Ligand Complexes. ○
- 113 Green biosynthesis of rare DHA-phospholipids by lipase-catalyzed transesterification with edible algal oil in solvent-free system and catalytic mechanism study. 11, ○
- 112 Development of New Targeted Nanotherapy Combined with Magneto-Fluorescent Nanoparticles against Colorectal Cancer. **2023**, 24, 6612 ○
- 111 Quantitative structure activity relationship studies of androgen receptor binding affinity of endocrine disruptor chemicals with index of ideality of correlation, their molecular docking, molecular dynamics and ADME studies. 1-16 ○
- 110 Therapeutic Role of DGJ (1-deoxygalactonojirimycin) in Fabry Disease: Theoretical Insights. **2023**, 8, ○
- 109 Synthesis, X-ray Crystal Structure, Anticancer, Hirshfeld Surface Analysis, DFT, TD-DFT, ADMET, and Molecular Docking of 3-Phenyl-1,2,4-triazolo[3,4-h]-13,4-thiaza-11-crown-4. **2023**, 28, 3166 ○
- 108 Tailoring Hydrophobicity of Thioflavin T to Optimize A β Fibril Bioimaging. ○
- 107 Curcumin and Plumbagin Synergistically Target the PI3K/Akt/mTOR Pathway: A Prospective Role in Cancer Treatment. **2023**, 24, 6651 ○
- 106 Prediction of dynamic allostery for the transmembrane domain of the sweet taste receptor subunit, TAS1R3. **2023**, 6, ○
- 105 Synthesis of Pyrazoline-Embedded 1,2,3-Triazole Derivatives via 1,3-Dipolar Cycloaddition Reactions with in vitro and in silico Studies. **2023**, 8, ○
- 104 Computational Investigations into Two-Photon Fibril Imaging Using the DANIR-2c Probe. **2023**, 127, 3119-3125○
- 103 Anti-inflammatory activities of black raspberry seed ellagitannins and their structural effects on the stimulation of glucagon-like peptide-1 secretion and intestinal bitter taste receptors. ○
- 102 In silico identification and molecular dynamic simulations of derivatives of 6,6-dimethyl-3-azabicyclo[3.1.0]hexane-2-carboxamide against main protease 3CLpro of SARS-CoV-2 viral infection. **2023**, 29, ○

- 101 Biologically potent organotin(IV) complexes of N-acetylated α -amino acids with spectroscopic, X-ray powder diffraction and molecular docking studies. **2023**, 13, 10768-10789 ○
- 100 Crystal Structure and Functional Characterization of an S-Formylglutathione Hydrolase (BuSFGH) from Burkholderiaceae sp.. **2023**, 13, 621 ○
- 99 Identifying novel selective PPO inhibitors through structure-based virtual screening and bio-evaluation. **2023**, 13, 10873-10883 ○
- 98 First In Vivo Insights on the Effects of Tempol-Methoxycinnamate, a New UV Filter, as Alternative to Octyl Methoxycinnamate, on Zebrafish Early Development. **2023**, 24, 6767 ○
- 97 Combinatorial Effect of Ligand Aromaticity, Torsion Number and Hydrophobicity on the Predictability of AutoDock for Protein-Ligand Binding Conformation. **2023**, 38, 70-75 ○
- 96 Synthesis, spectral, DFT calculation, antimicrobial, antioxidant, DNA/BSA binding and molecular docking studies of bio-pharmacologically active pyrimidine appended Cu(II) and Zn(II) complexes. 1-15 ○
- 95 In Silico Structural and Functional Analysis of the Mitochondrial Malate Transporters in Oleaginous Fungus *Mucor circinelloides* WJ11. **2023**, 13, 705 ○
- 94 Synthesis, Characterization and Crystal Structures of Thiosemicarbazones with Urease Inhibitory Activity. **2023**, 64, 462-473 ○
- 93 Metagenome mining and functional analysis reveal oxidized guanine DNA repair at the Lost City Hydrothermal Field. ○
- 92 SARS-CoV-2 evolved variants optimize binding to cellular glycocalyx. **2023**, 4, 101346 ○
- 91 Structure-based virtual screening and molecular dynamics simulations for detecting novel candidates for allosteric inhibition of EGFR T790M. 1-27 ○
- 90 Synthesis and anxiolytic effect of europium metallic complex containing lapachol [Eu(DBM)₃. LAP] in adult zebrafish through serotonergic neurotransmission: in vivo and in silico approach. 1-13 ○
- 89 Riparin-B as a Potential Inhibitor of AdeABC Efflux System from *Acinetobacter baumannii*. **2023**, 2023, 1-10 ○
- 88 Natural Andrographolide Isolated from *Andrographis paniculata* as Potent Epileptic Agent: Spectroscopy, Molecular Structure, and Molecular Docking Investigation. ○
- 87 Structure-based molecular networking, molecular docking, dynamics simulation and pharmacokinetic studies of *Olax subscorpioidea* for identification of potential inhibitors against selected cancer targets. 1-16 ○
- 86 A combined spectroscopic and quantum chemical approach to study the molecular interaction between anti-inflammatory drug Hydrocortisone and amino acid L-Phenylalanine. **2023**, 135546 ○
- 85 The Yeast Permease Agp2 Senses Cycloheximide and Undergoes Degradation That Requires the Small Protein Brp1-Cellular Fate of Agp2 in Response to Cycloheximide. **2023**, 24, 6975 ○
- 84 Tadalafil Rescues the p.M325T Mutant of Best1 Chloride Channel. **2023**, 28, 3317 ○

- 83 Exploring polyamine interactions and binding pockets in SARS-CoV-2 ORF3a. **2023**, 122, 108487 ○
- 82 Green Synthesis, Antimycobacterial Evaluation and Molecular Docking Studies of Novel 2,3-Dihydro-1H-pyrazol-4-yl-naphthalene-1,4-diones. **2023**, 135556 ○
- 81 Comparative modeling and enzymatic affinity of novel haloacid dehalogenase from *Bacillus megaterium* strain BHS1 isolated from alkaline Blue Lake in Turkey. 1-14 ○
- 80 Tool and Techniques on Computer-Aided Drug Design for Targeted Cancer Therapy. **2023**, 781-829 ○
- 79 Molecular interactions, binding stability, and synergistic inhibition on Acetylcholinesterase activity of Safranin O in combination with Quercetin and Gallic acid: In vitro and in silico study. **2023**, 1286, 135562 ○
- 78 Clinical side-effects based drug repositioning for anti-epileptic activity. 1-12 ○
- 77 Structure-Activity Relationship of Methyl 4-Aminobenzoate Derivatives as Being Drug Candidate Targeting Glutathione Related Enzymes: in Vitro and in Silico Approaches. ○
- 76 The comparison between zein-anthocyanins complex and nanoparticle systems: Stability enhancement, interaction mechanism, and in silico approaches. **2023**, 420, 136136 ○
- 75 Computational Insights into the Dynamic Structural Features and Binding Characteristics of Recombinase UvsX Compared with RecA. **2023**, 28, 3363 ○
- 74 Enhancing Enzyme Activity and Thermostability of *Bacillus amyloliquefaciens* Chitosanase BaCsn46A Through Saturation Mutagenesis at Ser196. **2023**, 80, ○
- 73 Investigation on the growth, structure and physical properties of pyridin-1-ium-2-carboxylate benzimidazole (1:1) hydrate single crystal. **2023**, 34, ○
- 72 Binding kinetics study of SARS-CoV-2 main protease and potential inhibitors via molecular dynamics simulations. ○
- 71 Interaction between escitalopram and ibuprofen or paracetamol: DFT and molecular docking on the drug-drug interactions. 1-15 ○
- 70 Interactions between curcumin and human salt-induced kinase 3 elucidated from computational tools and experimental methods. 14, ○
- 69 In vitro, in vivo, and in silico analysis of synbiotics as preventive interventions for lipid metabolism in ethanol-induced adipose tissue injury. **2023**, 22, ○
- 68 Identification of Novel Cyclooxygenase-1 Selective Inhibitors of Thiadiazole-Based Scaffold as Potent Anti-Inflammatory Agents with Safety Gastric and Cytotoxic Profile. **2023**, 28, 3416 ○
- 67 Pomiferin targets SERCA, mTOR, and P-gp to induce autophagic cell death in apoptosis-resistant cancer cells, and reverses the MDR phenotype in cisplatin-resistant tumors in vivo. **2023**, 106769 ○
- 66 Amorphous Pterostilbene Delivery Systems Preparation Innovative Approach to Preparation Optimization. **2023**, 15, 1231 ○

- 65 Discovery of the Universal tRNA Binding Mode for the TsaD-like Components of the t6A tRNA Modification Pathway. **2023**, 3, 288-306 ○
- 64 Identification of novel drug targets in *Porphyromonas gingivalis* and proposing inhibitors against Acetate kinase using structure-based virtual screening. **2023**, ○
- 63 Carotenoids from female *Grapsus albolineatus* as potential anti-ageing compounds. 11, 1457 ○
- 62 Gibberellin 2-Oxidases in Potato (*Solanum tuberosum* L.): Cloning, Characterization, In Silico Analysis and Molecular Docking. ○
- 61 Identification of Potential p38 α Inhibitors via In Silico Screening, In Vitro Bioassay and Molecular Dynamics Simulation Studies. **2023**, 24, 7360 ○
- 60 Identification of merozoite secreted repertoire and immuno-pharmacological inhibition of a novel host-parasite interaction to block malarial infection. ○
- 59 Cytochrome P450 mediates the formation of four new citrinin metabolites. **2023**, 102663 ○
- 58 Phytochemical profiling, antimicrobial, antibiofilm, insecticidal, and anti-leishmanial properties of aqueous extract from *Juglans regia* L. root bark: In vitro and in silico approaches. **2023**, 26, 1079-1097 ○
- 57 Potential mechanisms of osthole against bladder cancer cells based on network pharmacology, molecular docking, and experimental validation. **2023**, 23, ○
- 56 Drug-target interaction prediction based on spatial consistency constraint and graph convolutional autoencoder. **2023**, 24, ○
- 55 Plumbagin Suppresses Growth, Induces Apoptosis, and Inhibits Migration in Cholangiocarcinoma via Reactive Oxygen Species Generation and Mitochondrial Function. 097312962311582 ○
- 54 Repositioning Cannabinoids and Terpenes as Novel EGFR-TKIs Candidates for Targeted Therapy Against Cancer: A virtual screening model using CADD and biophysical simulations. **2023**, 9, e15545 ○
- 53 Exploring the hub genes and mechanisms of *Daphne altaica*. treating esophageal squamous cell carcinoma based on Network Pharmacology and Bioinformatics Analysis. ○
- 52 Identification and molecular interactions of novel ACE inhibitory peptides from rapeseed protein. **2023**, 136085 ○
- 51 Discovery of Highly Potent HDAC8 PROTACs with Anti-tumor Activity. **2023**, 106546 ○
- 50 In Silico Analysis of Anti-Inflammatory and Antioxidant Properties of Bioactive Compounds from *Crescentia cujete* L.. **2023**, 28, 3547 ○
- 49 Study on molecular mechanisms of destabilizing A β (142) protofibrils by licochalcone A and licochalcone B using molecular dynamics simulations. **2023**, 108500 ○
- 48 Zeolite@-Ag-S-CH₂-COOH catalyzed synthesis of bis-cyclohexenones and their antibacterial evaluation and molecular docking study. **2023**, 122710 ○

- 47 Visfatin-induced upregulation of lipogenesis via EGFR/AKT/GSK3 β pathway promotes breast cancer cell growth. **2023**, 110686
- 46 Small Molecule Attenuates Bacterial Virulence by Targeting Conserved Response Regulator.
- 45 The Cytotoxicity Profile, Apoptosis Mechanism, and Molecular Docking Studies of a Series of Benzimidazolium Derivative Morpholine-Substituted Ag(I) Heterocyclic Carbene Complexes.
- 44 Essential Oil of *Origanum vulgare* var. *aureum* L. from Western Romania: Chemical Analysis, In Vitro and In Silico Screening of Its Antioxidant Activity. **2023**, 13, 5076
- 43 Identification of Novel Compounds Inhibiting the Kinase Activity of the CDK5/p25 Complex via Direct Binding to p25.
- 42 Characterization of the bioactive compounds with efficacy against epilepsy from the herb pairs *Polygala tenuifolia* - *Zizyphus jujuba* by modulating CHRNA4/CaMKII signaling pathway: LC-MS/MS combined with network pharmacology analysis and experimental evidence.
- 41 Solvent effect on molecular, electronic parameters, topological analysis and Fukui function evaluation with biological studies of Imidazo [1, 2-a] pyridine-8-carboxylic acid. **2023**, 121863
- 40 Potential profound fluctuation in tacrolimus concentration on consumption of pomegranate rind extract: A Pharmacokinetic Experiment. 14,
- 39 Catalytic resilience of multicomponent aromatic ring-hydroxylating dioxygenases in *Pseudomonas* for degradation of polycyclic aromatic hydrocarbons. **2023**, 39,
- 38 Structure based virtual screening, molecular dynamic simulation to identify the oxadiazole derivatives as inhibitors of Enterococcus D-Ala-D-Ser ligase for combating vancomycin resistance. **2023**, 106965
- 37 Study of the binding interaction of salmon sperm DNA with nintedanib, a tyrosine kinase inhibitor using multi-spectroscopic, thermodynamic, and in silico approaches. 1-11
- 36 Allosteric modulation of GluN1/GluN3 NMDA receptors by GluN1-selective competitive antagonists. **2023**, 155,
- 35 Concussion Diagnostics: A New Diagnostic Approach Using PET Technology. **2023**, 4,
- 34 Molecular Structure, Hydrogen Bonding Interactions and Docking Simulations of Nicotinamide (Monomeric and Trimeric Models) by Using Spectroscopy and Theoretical Approach. 1-19
- 33 Insight into the phytochemical profile and antimicrobial activities of *Amomum subulatum* and *Amomum xanthioides*: an in vitro and in silico study. 14,
- 32 Biocatalytic Access to Chiral Benzazepines Using Imine Reductases. 6185-6194
- 31 Repurposing of phytocompounds-derived novel bioactive compounds possessing promising anticancer and cancer therapeutic efficacy through molecular docking, MD simulation, and drug-likeness/ADMET studies. **2023**, 201-222
- 30 Synthesis, Spectroscopic, Crystallographic, Quantum and Molecular Docking Investigations of *cis*-4,5-diphenylimidazolidine-2-thione. **2023**, 135633

- 29 Characterizing Soft Matter Self-Assembly and Material Properties with Advanced Molecular Dynamics and Data-Driven Methods. **2023**, 1197-1220 ○
- 28 Dihydrotestosterone-based A-ring-fused pyridines: microwave-assisted synthesis and biological evaluation in prostate cancer cells compared to structurally related quinolines. **2023**, 106315 ○
- 27 The mechanism of peach kernel and safflower herb-pair for the treatment of liver fibrosis based on network pharmacology and molecular docking technology: A review. **2023**, 102, e33593 ○
- 26 Design, Synthesis and Molecular Docking of 1,2,4-Triazole Schiff Base Hybrids as Tubulin, EGFR Inhibitors and Apoptosis-Inducers. **2023**, 135621 ○
- 25 Pharmacological mechanism of indigenous herb *Exacum lawii* on cisplatin instigated toxicity in human embryonic kidney cells (HEK-293). **2023**, 100454 ○
- 24 Thioether-substituted Benzimidazolium Salts: Synthesis, Characterization, Crystal Structure, and Their Inhibitory Properties Against Carbonic Anhydrase and Xanthine Oxidase. **2023**, 135640 ○
- 23 Identification of novel Zika virus NS3 protease inhibitors with different inhibition modes by integrative experimental and computational approaches. **2023**, ○
- 22 Investigation of the inhibitory behavior of XFE and mitoxantrone molecules in interaction with AKT1 protein: a molecular dynamics simulation study. **2023**, 29, ○
- 21 Homology Modeling, Screening, and Identification of Potential FOXO6 Inhibitors Curtail Gastric Cancer Progression: an In Silico Drug Repurposing Approach. ○
- 20 Molecular dynamics simulation of the interaction of a raspberry polygalacturonase (RiPG) with a PG inhibiting protein (RiPGIP) isolated from ripening raspberry (*Rubus idaeus* cv. Heritage) fruit as a model to understand proteins interaction during fruit softening. **2023**, 108502 ○
- 19 In Silico Interactions of Natural and Synthetic Compounds with Key Proteins Involved in Alzheimer's Disease: Prospects for Designing New Therapeutics Compound. ○
- 18 The signal conversion strategy using the lytic transglycosylase Cag4-double nanoporous gold co-catalysis for the rapid screening of drugs against *Helicobacter pylori* infection. **2023**, 115345 ○
- 17 Targeting the Human Influenza A Virus: The Methods, Limitations, and Pitfalls of Virtual Screening for Drug-like Candidates Including Scaffold Hopping and Compound Profiling. **2023**, 15, 1056 ○
- 16 A pyrazolone-based dinuclear Cu(II) Schiff-base complex: DFT studies on the rate-determining steps of the tautomerism in the ligand and molecular docking modelling with COVID-19 main protease (6LU7). 1-17 ○
- 15 Development of quinazolinone and vanillin acrylamide hybrids as multi-target directed ligands against Alzheimer's disease and mechanistic insights into their binding with acetylcholinesterase. 1-18 ○
- 14 VirtualFlow 2.0 - The Next Generation Drug Discovery Platform Enabling Adaptive Screens of 69 Billion Molecules. ○
- 13 Uni-Dock: GPU-Accelerated Docking Enables Ultralarge Virtual Screening. ○
- 12 In-silico method in predicting potential targets from marine natural products in the waters of South Sulawesi, Indonesia. **2023**, ○

- 11 Proximity-Induced Nucleic Acid Degradation (PINAD) Approach to Targeted RNA Degradation Using Small Molecules. ○
- 10 Experimental and Theoretical Investigation of Inclusion Complexes of β -Cyclodextrin with Fingolimod. **2023**, 97, 469-476 ○
- 9 Targeting the vital non-structural proteins (NSP12, NSP7, NSP8 and NSP3) from SARS-CoV-2 and inhibition of RNA polymerase by natural bioactive compound naringenin as a promising drug candidate against COVID-19. **2023**, 1287, 135642 ○
- 8 Exploration of natural compounds against the human mpox virus DNA-dependent RNA polymerase in silico. **2023**, 16, 996-1003 ○
- 7 The difference in the cellular uptake of tocopherol and tocotrienol is influenced by their affinities to albumin. **2023**, 13, ○
- 6 Discovery of pyrimidine-5-carboxamide derivatives as novel salt-inducible kinases (SIKs) inhibitors for inflammatory bowel disease (IBD) treatment. **2023**, 256, 115469 ○
- 5 Exploring the mechanism by which quercetin re-sensitizes breast cancer to paclitaxel: network pharmacology, molecular docking, and experimental verification. ○
- 4 Genomic analyses provide insights into the polyploidization-driven herbicide adaptation in *Leptochloa* weeds. ○
- 3 Estimation of Antioxidant Potential, Phytochemical Profiling, and In Silico Characterization of Hepatoprotective Biomarkers of *Phyllanthus fraternus* G.L. Webster Leaves. **2023**, 8, ○
- 2 Exploring the Antioxidant and Anti-Inflammatory Potential of *Wilckia maritima*: In Vitro and In Silico Investigations. **2023**, 11, 1497 ○
- 1 Spectroscopic analysis of 2-amino-1-naphthalenesulfonic acid, molecular docking, and evaluation of the electronic properties of several solvents. 1-20 ○