

Vinicius G Maltarollo

List of Publications by Year in descending order

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26
papers

466
citations

687363

13
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713466

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26
all docs

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docs citations

26
times ranked

597
citing authors

#	ARTICLE	IF	CITATIONS
1	Transfer and Multi-task Learning in QSAR Modeling: Advances and Challenges. <i>Frontiers in Pharmacology</i> , 2018, 9, 74.	3.5	87
2	Advances and Perspectives in Applying Deep Learning for Drug Design and Discovery. <i>Frontiers in Robotics and AI</i> , 2019, 6, 108.	3.2	60
3	Recent Advances in the Prediction of Pharmacokinetics Properties in Drug Design Studies: A Review. <i>ChemMedChem</i> , 2022, 17, .	3.2	44
4	Advanced QSAR studies on PPAR γ ligands related to metabolic diseases. <i>Journal of the Brazilian Chemical Society</i> , 2012, 23, 78-84.	0.6	23
5	Structure-Based Virtual Screening and Discovery of New PPAR γ / δ Dual Agonist and PPAR γ and δ Agonists. <i>PLoS ONE</i> , 2015, 10, e0118790.	2.5	21
6	Theoretical study of tautomers and photoisomers of avobenzone by DFT methods. <i>Journal of Molecular Modeling</i> , 2015, 21, 319.	1.8	20
7	Computational studies of TGF- β RI (ALK-5) inhibitors: Analysis of the binding interactions between ligand α -receptor using 2D and 3D techniques. <i>European Journal of Pharmaceutical Sciences</i> , 2013, 49, 542-549.	4.0	18
8	Quantitative structure α -activity relationships (HQSAR, CoMFA, and CoMSIA) studies for COX-2 selective inhibitors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 1436-1445.	3.5	17
9	Antibacterial activity of synthetic 1,3 α -bis(aryloxy)propan α -2 α -amines against Gram α -positive bacteria. <i>MicrobiologyOpen</i> , 2019, 8, e814.	3.0	16
10	Knowing and combating the enemy: a brief review on SARS-CoV-2 and computational approaches applied to the discovery of drug candidates. <i>Bioscience Reports</i> , 2021, 41, .	2.4	16
11	Theoretical study on the molecular and electronic properties of some substances used for diabetes mellitus treatment. <i>Journal of Molecular Modeling</i> , 2010, 16, 799-804.	1.8	15
12	Hologram QSAR Studies of Antiprotozoal Activities of Sesquiterpene Lactones. <i>Molecules</i> , 2014, 19, 10546-10562.	3.8	15
13	GQ-11: A new PPAR agonist improves obesity-induced metabolic alterations in LDLr α / β mice. <i>International Journal of Obesity</i> , 2018, 42, 1062-1072.	3.4	15
14	Application of bioisosterism in design of the semicarbazone derivatives as cruzain inhibitors: a theoretical and experimental study. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 1244-1259.	3.5	14
15	Role of physicochemical properties in the activation of peroxisome proliferator-activated receptor γ . <i>Journal of Molecular Modeling</i> , 2011, 17, 2549-2558.	1.8	12
16	Hologram quantitative structure α -activity relationship and comparative molecular interaction field analysis of aminothiazole and thiazolesulfonamide as reversible LSD1 inhibitors. <i>Future Medicinal Chemistry</i> , 2015, 7, 1381-1394.	2.3	12
17	Synthesis of arylfuran derivatives as potential antibacterial agents. <i>Medicinal Chemistry Research</i> , 2021, 30, 1074-1086.	2.4	9
18	Ligand α -and Structure α -Based Drug Design Strategies and PPAR γ / δ Selectivity. <i>Chemical Biology and Drug Design</i> , 2012, 80, 533-544.	3.2	8

#	ARTICLE	IF	CITATIONS
19	Current trends in quantitative structure–activity relationship validation and applications on drug discovery. <i>Future Science OA</i> , 2017, 3, FSO214.	1.9	8
20	In silico studies on the interaction between bioactive ligands and ALK5, a biological target related to the cancer treatment. <i>Journal of Biomolecular Structure and Dynamics</i> , 2016, 34, 2045-2053.	3.5	7
21	Structure-Based Virtual Screening, Molecular Dynamics and Binding Free Energy Calculations of Hit Candidates as ALK-5 Inhibitors. <i>Molecules</i> , 2020, 25, 264.	3.8	7
22	The Role of QSAR and Virtual Screening Studies in Type 2 Diabetes Drug Discovery. <i>Medicinal Chemistry</i> , 2017, 13, 706-720.	1.5	7
23	Molecular Features Related to HIV Integrase Inhibition Obtained from Structure- and Ligand-Based Approaches. <i>PLoS ONE</i> , 2014, 9, e81301.	2.5	6
24	On the relationship of anthranilic derivatives structure and the FXR (Farnesoid X receptor) agonist activity. <i>Journal of Biomolecular Structure and Dynamics</i> , 2018, 36, 4378-4391.	3.5	5
25	Understanding electrostatic and steric requirements related to hypertensive action of AT1 antagonists using molecular modeling techniques. <i>Journal of Molecular Modeling</i> , 2014, 20, 2231.	1.8	2
26	Understanding PPAR- γ affinity and selectivity using hologram quantitative structure–activity modeling, molecular docking and GRID calculations. <i>Future Medicinal Chemistry</i> , 2016, 8, 1913-1926.	2.3	2