

List of Publications by Year in descending order

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citing authors

#	ARTICLE	IF	CITATIONS
1	Ligand pathways in estrogen-related receptors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2023, 41, 1639-1648.	2.0	1
2	Evaluation of Xa inhibitors as potential inhibitors of the SARS-CoV-2 Mpro protease. <i>PLoS ONE</i> , 2022, 17, e0262482.	1.1	7
3	Activation of retinoic acid-related orphan receptor $\hat{I}^3(t)$ by parabens and benzophenone UV-filters. <i>Toxicology</i> , 2022, 471, 153159.	2.0	5
4	Conformational Landscape of Cytochrome P450 Reductase Interactions. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1023.	1.8	8
5	Conformational Changes of Thyroid Receptors in Response to Antagonists. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 1010-1019.	2.5	4
6	Computational Assessment of Combination Therapy of Androgen Receptor-Targeting Compounds. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 1001-1009.	2.5	4
7	Decision Making in Structure-Based Drug Discovery: Visual Inspection of Docking Results. <i>Journal of Medicinal Chemistry</i> , 2021, 64, 2489-2500.	2.9	113
8	Computational Selectivity Assessment of Protease Inhibitors against SARS-CoV-2. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2065.	1.8	3
9	A Conserved Allosteric Site on Drug-Metabolizing CYPs: A Systematic Computational Assessment. <i>International Journal of Molecular Sciences</i> , 2021, 22, 13215.	1.8	9
10	Allosteric Binding Sites On Nuclear Receptors: Focus On Drug Efficacy and Selectivity. <i>International Journal of Molecular Sciences</i> , 2020, 21, 534.	1.8	11
11	Potential Inhibitors for Novel Coronavirus Protease Identified by Virtual Screening of 606 Million Compounds. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3626.	1.8	131
12	Ligand Pathways in Nuclear Receptors. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 3100-3109.	2.5	23
13	Spontaneous Ligand Access Events to Membrane-Bound Cytochrome P450 2D6 Sampled at Atomic Resolution. <i>Scientific Reports</i> , 2019, 9, 16411.	1.6	13
14	Molecular Dynamics Simulations Reveal Structural Differences among Allelic Variants of Membrane-Anchored Cytochrome P450 2D6. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 1962-1975.	2.5	21