

Laercio Pol-Fachin

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

619
citations

759233

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docs citations

22
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973
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#	ARTICLE	IF	CITATIONS
1	GROMOS 53A6 _{GLYC} , an Improved GROMOS Force Field for Hexopyranose-Based Carbohydrates. <i>Journal of Chemical Theory and Computation</i> , 2012, 8, 4681-4690.	5.3	132
2	GROMOS96 43a1 performance on the characterization of glycoprotein conformational ensembles through molecular dynamics simulations. <i>Carbohydrate Research</i> , 2009, 344, 491-500.	2.3	93
3	Polymyxin Binding to the Bacterial Outer Membrane Reveals Cation Displacement and Increasing Membrane Curvature in Susceptible but Not in Resistant Lipopolysaccharide Chemotypes. <i>Journal of Chemical Information and Modeling</i> , 2017, 57, 2181-2193.	5.4	54
4	Depiction of the forces participating in the 2-O-sulfo- α -L-iduronic acid conformational preference in heparin sequences in aqueous solutions. <i>Carbohydrate Research</i> , 2008, 343, 1435-1445.	2.3	49
5	Anticoagulant Activity of a Unique Sulfated Pyranosic (1 \rightarrow 3)- β -L-Arabinan through Direct Interaction with Thrombin. <i>Journal of Biological Chemistry</i> , 2013, 288, 223-233.	3.4	46
6	A Unique 2-Sulfated β -Galactan from the Egg Jelly of the Sea Urchin <i>Glyptocidaris crenularis</i> . <i>Journal of Biological Chemistry</i> , 2009, 284, 18790-18800.	3.4	44
7	Extension and validation of the GROMOS 53A6 _{scp>glyc</sub> parameter set for glycoproteins. <i>Journal of Computational Chemistry</i>, 2014, 35, 2087-2095.}	3.3	42
8	Effects of glycosylation on heparin binding and antithrombin activation by heparin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2011, 79, 2735-2745.	2.6	22
9	Atomic Model and Micelle Dynamics of QS-21 Saponin. <i>Molecules</i> , 2014, 19, 3744-3760.	3.8	21
10	Inhibition of the hemolytic activity caused by <i>Staphylococcus aureus</i> α -hemolysin through isatin-Schiff copper(II) complexes. <i>FEMS Microbiology Letters</i> , 2016, 363, fmv207.	1.8	20
11	Conformational stability of the epidermal growth factor (EGF) receptor as influenced by glycosylation, dimerization and EGF hormone binding. <i>Proteins: Structure, Function and Bioinformatics</i> , 2017, 85, 561-570.	2.6	18
12	Solution conformation and dynamics of exopolysaccharides from <i>Burkholderia</i> species. <i>Carbohydrate Research</i> , 2010, 345, 1922-1931.	2.3	13
13	Glycosylation is crucial for a proper catalytic site organization in human glucocerebrosidase. <i>Glycoconjugate Journal</i> , 2016, 33, 237-244.	2.7	13
14	Characterization of the conformational ensemble from bioactive N-acylhydrazone derivatives. <i>Journal of Molecular Graphics and Modelling</i> , 2010, 28, 446-454.	2.4	12
15	Assessment of Glycoproteins Dynamics from Computer Simulations. <i>Mini-Reviews in Organic Chemistry</i> , 2011, 8, 229-238.	1.3	12
16	Unrestrained Conformational Characterization of <i>Stenocereus eruca</i> Saponins in Aqueous and Nonaqueous Solvents. <i>Journal of Natural Products</i> , 2012, 75, 1196-1200.	3.0	11
17	Structural glycobiology of the major allergen of <i>Artemisia vulgaris</i> pollen, Art v 1: O-glycosylation influence on the protein dynamics and allergenicity. <i>Glycobiology</i> , 2012, 22, 817-825.	2.5	7
18	<i>Staphylococcus aureus</i> α -toxin in aqueous solution: Behavior in monomeric and multimeric states. <i>Biophysical Chemistry</i> , 2017, 227, 21-28.	2.8	4

#	ARTICLE	IF	CITATIONS
19	The Lazy Life of Lipid-Linked Oligosaccharides in All Life Domains. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 631-643.	5.4	4
20	Insights into the effects of glycosylation and the monosaccharide-binding activity of the plant lectin CrataBL. <i>Glycoconjugate Journal</i> , 2017, 34, 515-522.	2.7	2
21	Cover Image, Volume 85, Issue 4. <i>Proteins: Structure, Function and Bioinformatics</i> , 2017, 85, C4.	2.6	0
22	Duffy binding-like 1 adhesin from <i>Plasmodium falciparum</i> recognizes ABH histo-blood group saccharide in a type specific manner. <i>Carbohydrate Polymers</i> , 2019, 207, 266-275.	10.2	0