

Herschel Wade

List of Publications by Year in descending order

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Version: 2024-02-01

10
papers

161
citations

1477746

6
h-index

1473754

9
g-index

50
all docs

50
docs citations

50
times ranked

299
citing authors

#	ARTICLE	IF	CITATIONS
1	Naturally Occurring Variants of Muscle Type Creatine Kinase Exhibit Altered Tenofovir Monophosphate Phosphorylation Activity. <i>FASEB Journal</i> , 2020, 34, 1-1.	0.2	0
2	Probing Ligand Structure-Activity Relationships in Pregnane X Receptor (PXR): Efavirenz and 8-Hydroxyefavirenz Exhibit Divergence in Activation. <i>ChemMedChem</i> , 2018, 13, 736-747.	1.6	7
3	Unconventional Coupling between Ligand Recognition and Allosteric Control in the Multidrug Resistance Gene Regulator, BmrR. <i>ChemMedChem</i> , 2017, 12, 426-430.	1.6	2
4	Charge is Major Determinant of Activation of the Ligand-Responsive Multidrug Resistance Gene Regulator, BmrR. <i>ChemMedChem</i> , 2016, 11, 1038-1041.	1.6	4
5	Solution Binding and Structural Analyses Reveal Potential Multidrug Resistance Functions for SAV2435 and CTR107 and Other GyrI-like Proteins. <i>Biochemistry</i> , 2016, 55, 4850-4863.	1.2	11
6	Solution-Binding and Molecular Docking Approaches Combine to Provide an Expanded View of Multidrug Recognition in the MDR Gene Regulator BmrR. <i>Journal of Chemical Information and Modeling</i> , 2016, 56, 377-389.	2.5	4
7	Allosteric Coupling via Distant Disorder-to-Order Transitions. <i>Journal of Molecular Biology</i> , 2015, 427, 1695-1704.	2.0	26
8	A PWWP Domain-Containing Protein Targets the NuA3 Acetyltransferase Complex via Histone H3 Lysine 36 trimethylation to Coordinate Transcriptional Elongation at Coding Regions. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 2883-2895.	2.5	48
9	Structural contributions to multidrug recognition in the multidrug resistance (MDR) gene regulator, BmrR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 11046-11051.	3.3	36
10	MD recognition by MDR gene regulators. <i>Current Opinion in Structural Biology</i> , 2010, 20, 489-496.	2.6	23