

Mikolaj Feliks

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

401
citations

1306789

7
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1281420

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all docs

11
docs citations

11
times ranked

839
citing authors

#	ARTICLE	IF	CITATIONS
1	Chromophore twisting in the excited state of a photoswitchable fluorescent protein captured by time-resolved serial femtosecond crystallography. <i>Nature Chemistry</i> , 2018, 10, 31-37.	6.6	152
2	Structural Basis for a Kolbe-Type Decarboxylation Catalyzed by a Glycyl Radical Enzyme. <i>Journal of the American Chemical Society</i> , 2011, 133, 14666-14674.	6.6	56
3	Photoswitching mechanism of a fluorescent protein revealed by time-resolved crystallography and transient absorption spectroscopy. <i>Nature Communications</i> , 2020, 11, 741.	5.8	56
4	Glycerol Dehydration by the 12-O -Independent Enzyme May Not Involve the Migration of a Hydroxyl Group: A Computational Study. <i>Journal of Physical Chemistry B</i> , 2012, 116, 7076-7087.	1.2	33
5	Catalytic Mechanism of the Glycyl Radical Enzyme 4-Hydroxyphenylacetate Decarboxylase from Continuum Electrostatic and QC/MM Calculations. <i>Journal of the American Chemical Society</i> , 2013, 135, 14574-14585.	6.6	31
6	On the control of the proton current in the voltage-gated proton channel Hv1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 10321-10326.	3.3	28
7	Structural Determinants of Improved Fluorescence in a Family of Bacteriophytochrome-Based Infrared Fluorescent Proteins: Insights from Continuum Electrostatic Calculations and Molecular Dynamics Simulations. <i>Biochemistry</i> , 2016, 55, 4263-4274.	1.2	24
8	Norovirus RNA-dependent RNA polymerase: A computational study of metal-binding preferences. <i>Proteins: Structure, Function and Bioinformatics</i> , 2017, 85, 1435-1445.	1.5	10
9	<i>pCetk</i> : A pDynamo-based Toolkit for Protonation State Calculations in Proteins. <i>Journal of Chemical Information and Modeling</i> , 2015, 55, 2288-2296.	2.5	6
10	Predicting substituent effects on activation energy changes by static catalytic fields. <i>Journal of Molecular Modeling</i> , 2018, 24, 28.	0.8	3
11	Catalytic Mechanism of Peptidoglycan Deacetylase: A Computational Study. <i>Journal of Physical Chemistry B</i> , 2017, 121, 89-99.	1.2	2