Willy Wriggers

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/156063/publications.pdf

Version: 2024-02-01

1039880 940416 23 272 9 16 citations g-index h-index papers 24 24 24 350 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Interpreting Cytoskeletal Filaments in Cryo-Electron Tomograms with Shape-Constrained Deconvolution. Microscopy and Microanalysis, 2021, 27, 72-73.	0.2	O
2	Tracing Filaments in Simulated and Experimental 3D Cryo-Electron Tomography Maps Using a Fast Dynamic Programming Algorithm. Microscopy and Microanalysis, 2021, 27, 3236-3237.	0.2	1
3	Mechanism for the Unfolding of the TOP7 Protein in Steered Molecular Dynamics Simulations as Revealed by Mutual Information Analysis. Frontiers in Molecular Biosciences, 2021, 8, 696609.	1.6	1
4	TomoSim: Simulation of Filamentous Cryo-Electron Tomograms., 2021,,.		5
5	Tracing Filaments in Simulated 3D Cryo-Electron Tomography Maps Using a Fast Dynamic Programming Algorithm. , 2021, , .		4
6	An Experimental Approach to Mapping of Magnetic Fields of CubeSat Attitude Actuator Representations. , 2020, , .		1
7	Faces of Contemporary CryoEM Information and Modeling. Journal of Chemical Information and Modeling, 2020, 60, 2407-2409.	2.5	1
8	Correction of Missing-Wedge Artifacts in Filamentous Tomograms by Template-Based Constrained Deconvolution. Journal of Chemical Information and Modeling, 2020, 60, 2626-2633.	2.5	9
9	A cryo-tomography-based volumetric model of the actin core of mouse vestibular hair cell stereocilia lacking plastin 1. Journal of Structural Biology, 2020, 210, 107461.	1.3	14
10	Cylindrical Similarity Measurement for Helices in Medium-Resolution Cryo-Electron Microscopy Density Maps. Journal of Chemical Information and Modeling, 2020, 60, 2644-2650.	2.5	12
11	Frontiers in CryoEM Modeling. Journal of Chemical Information and Modeling, 2019, 59, 3091-3093.	2.5	2
12	Accurate flexible refinement of atomic models against medium-resolution cryo-EM maps using damped dynamics. BMC Structural Biology, 2018, 18, 12.	2.3	29
13	Tracing Actin Filament Bundles in Three-Dimensional Electron Tomography Density Maps of Hair Cell Stereocilia. Molecules, 2018, 23, 882.	1.7	15
14	Comparing an Atomic Model or Structure to a Corresponding Cryo-electron Microscopy Image at the Central Axis of a Helix. Journal of Computational Biology, 2017, 24, 52-67.	0.8	8
15	Flexible Fitting and Refinement of Atomic Structures Using the Coarse-Grained DDFF Force Field Tailored to 5-10A Resolution Cryo-TEM Maps. Microscopy and Microanalysis, 2017, 23, 1226-1227.	0.2	O
16	An Information Theoretic Approach for Creating 3D Spatial Images from 4D Time Series Data. Microscopy and Microanalysis, 2017, 23, 100-101.	0.2	0
17	Detection of Protein Secondary Structure Patterns from 3D Cryo-TEM Maps at Medium Resolution - Combining the Best of SSETracer and VolTrac. Microscopy and Microanalysis, 2017, 23, 242-243.	0.2	1
18	Spatial Heat Maps from Fast Information Matching of Fast and Slow Degrees of Freedom: Application to Molecular Dynamics Simulations. Journal of Physical Chemistry B, 2016, 120, 8473-8484.	1.2	11

#	Article	IF	CITATIONS
19	Conserved Intramolecular Interactions Maintain Myosin Interacting-Heads Motifs Explaining Tarantula Muscle Super-Relaxed State Structural Basis. Journal of Molecular Biology, 2016, 428, 1142-1164.	2.0	82
20	Numerical geometry of map and model assessment. Journal of Structural Biology, 2015, 192, 255-261.	1.3	18
21	Comparison of an atomic model and its cryo-EM image at the central axis of a helix. , 2015, 2015, 1253-1259.		1
22	Multi-scale Visualization of Molecular Architecture Using Real-Time Ambient Occlusion in Sculptor. PLoS Computational Biology, 2015, 11, e1004516.	1.5	10
23	Automated tracing of filaments in 3D electron tomography reconstructions using Sculptor and Situs. Journal of Structural Biology, 2012, 178, 121-128.	1.3	47