

Robert Glaeser

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8117078/publications.pdf>

Version: 2024-02-01

65
papers

3,407
citations

218677

26
h-index

149698

56
g-index

67
all docs

67
docs citations

67
times ranked

3061
citing authors

#	ARTICLE	IF	CITATIONS
1	Perspective: Emerging strategies for determining atomic-resolution structures of macromolecular complexes within cells. <i>Journal of Structural Biology</i> , 2022, 214, 107827.	2.8	3
2	Perspective: Biochemical and Physical Constraints Associated With Preparing Thin Specimens for Single-Particle Cryo-EM. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 864829.	3.5	6
3	Defocus-dependent Thon-ring fading. <i>Ultramicroscopy</i> , 2021, 222, 113213.	1.9	11
4	High-power near-concentric Fabry-Perot cavity for phase contrast electron microscopy. <i>Review of Scientific Instruments</i> , 2021, 92, 053005.	1.3	24
5	Preparing Better Samples for Cryo-Electron Microscopy: Biochemical Challenges Do Not End with Isolation and Purification. <i>Annual Review of Biochemistry</i> , 2021, 90, 451-474.	11.1	33
6	Minimizing Crinkling of Soft Specimens Using Holey Gold Films on Molybdenum Grids for Cryogenic Electron Microscopy. <i>Microscopy and Microanalysis</i> , 2021, 27, 767-775.	0.4	5
7	Simple assay for adsorption of proteins to the air-water interface. <i>Journal of Structural Biology</i> , 2021, 213, 107798.	2.8	3
8	Conquer by cryo-EM without physically dividing. <i>Biochemical Society Transactions</i> , 2021, 49, 2287-2298.	3.4	4
9	Microscale Fluid Behavior during Cryo-EM Sample Blotting. <i>Biophysical Journal</i> , 2020, 118, 708-719.	0.5	43
10	Spectral DQE of the Volta phase plate. <i>Ultramicroscopy</i> , 2020, 218, 113079.	1.9	21
11	Observation of the Relativistic Reversal of the Ponderomotive Potential. <i>Physical Review Letters</i> , 2020, 124, 174801.	7.8	17
12	Current outcomes when optimizing "standard" sample preparation for single-particle cryo-EM. <i>Journal of Microscopy</i> , 2019, 276, 39-45.	1.8	41
13	Laser phase plate for transmission electron microscopy. <i>Nature Methods</i> , 2019, 16, 1016-1020.	19.0	118
14	How Good Can Single-Particle Cryo-EM Become? What Remains Before It Approaches Its Physical Limits?. <i>Annual Review of Biophysics</i> , 2019, 48, 45-61.	10.0	67
15	Proteins, interfaces, and cryo-EM grids. <i>Current Opinion in Colloid and Interface Science</i> , 2018, 34, 1-8.	7.4	117
16	Estimating the effect of finite depth of field in single-particle cryo-EM. <i>Ultramicroscopy</i> , 2018, 184, 94-99.	1.9	19
17	Monolayer-crystal streptavidin support films provide an internal standard of cryo-EM image quality. <i>Journal of Structural Biology</i> , 2017, 200, 307-313.	2.8	34
18	Signalling under the microscope. <i>Nature</i> , 2017, 546, 36-37.	27.8	2

#	ARTICLE	IF	CITATIONS
19	Opinion: hazards faced by macromolecules when confined to thin aqueous films. <i>Biophysics Reports</i> , 2017, 3, 1-7.	0.8	124
20	How Cryo-EM Became so Hot. <i>Cell</i> , 2017, 171, 1229-1231.	28.9	60
21	Multi-pass transmission electron microscopy. <i>Scientific Reports</i> , 2017, 7, 1699.	3.3	44
22	Reducing Electron Beam Damage with Multipass Transmission Electron Microscopy. <i>Microscopy and Microanalysis</i> , 2017, 23, 1794-1795.	0.4	2
23	Development of High-Resolution TEM for Imaging Native, Radiation-Sensitive Biological Macromolecules. <i>Microscopy and Microanalysis</i> , 2017, 23, 2290-2291.	0.4	0
24	Near-concentric Fabry-Pérot cavity for continuous-wave laser control of electron waves. <i>Optics Express</i> , 2017, 25, 14453.	3.4	19
25	Streptavidin Monolayer-Crystal Affinity Grids: A Step Toward Controlling What Happens During Cryo-EM Sample Preparation. <i>Microscopy and Microanalysis</i> , 2017, 23, 820-821.	0.4	0
26	Specimen Behavior in the Electron Beam. <i>Methods in Enzymology</i> , 2016, 579, 19-50.	1.0	67
27	Long shelf-life streptavidin support-films suitable for electron microscopy of biological macromolecules. <i>Journal of Structural Biology</i> , 2016, 195, 238-244.	2.8	58
28	Factors that Influence the Formation and Stability of Thin, Cryo-EM Specimens. <i>Biophysical Journal</i> , 2016, 110, 749-755.	0.5	81
29	Protein complexes in focus. <i>ELife</i> , 2016, 5, .	6.0	3
30	Use of Ultrananocrystalline Diamond as a Phase-contrast Aperture Material. <i>Microscopy and Microanalysis</i> , 2015, 21, 2297-2298.	0.4	0
31	Generalization of the Matsumoto-Tonomura approximation for the phase shift within an open aperture. <i>Ultramicroscopy</i> , 2014, 138, 1-3.	1.9	2
32	Disulfide Linkage and Structure of Highly Stable Yeast-derived Virus-like Particles of Murine Polyomavirus. <i>Journal of Biological Chemistry</i> , 2014, 289, 10411-10418.	3.4	16
33	Automated particle correspondence and accurate tilt-axis detection in tilted-image pairs. <i>Journal of Structural Biology</i> , 2014, 187, 66-75.	2.8	4
34	Investigating the Causes of Electrostatic Charging of Phase-contrast Apertures. <i>Microscopy and Microanalysis</i> , 2014, 20, 210-211.	0.4	0
35	Invited Review Article: Methods for imaging weak-phase objects in electron microscopy. <i>Review of Scientific Instruments</i> , 2013, 84, 111101.	1.3	117
36	Stroboscopic imaging of macromolecular complexes. <i>Nature Methods</i> , 2013, 10, 475-476.	19.0	15

#	ARTICLE	IF	CITATIONS
37	Replication and validation of cryo-EM structures. <i>Journal of Structural Biology</i> , 2013, 184, 379-380.	2.8	3
38	Ranking TEM cameras by their response to electron shot noise. <i>Ultramicroscopy</i> , 2013, 133, 1-7.	1.9	24
39	Minimizing electrostatic charging of an aperture used to produce in-focus phase contrast in the TEM. <i>Ultramicroscopy</i> , 2013, 135, 6-15.	1.9	18
40	Electron microscopy of biotinylated protein complexes bound to streptavidin monolayer crystals. <i>Journal of Structural Biology</i> , 2012, 180, 249-253.	2.8	43
41	Human Tripeptidyl Peptidase II: A Gentle Giant. <i>Structure</i> , 2012, 20, 565-566.	3.3	1
42	Reaching the Information Limit in Cryo-EM of Biological Macromolecules: Experimental Aspects. <i>Biophysical Journal</i> , 2011, 100, 2331-2337.	0.5	60
43	Precise beam-tilt alignment and collimation are required to minimize the phase error associated with coma in high-resolution cryo-EM. <i>Journal of Structural Biology</i> , 2011, 174, 1-10.	2.8	80
44	The surface of evaporated carbon films is an insulating, high-bandgap material. <i>Journal of Structural Biology</i> , 2011, 174, 420-423.	2.8	22
45	Experimental evaluation of support vector machine-based and correlation-based approaches to automatic particle selection. <i>Journal of Structural Biology</i> , 2011, 175, 319-328.	2.8	24
46	Design of a hybrid double-sideband/single-sideband (schlieren) objective aperture suitable for electron microscopy. <i>Ultramicroscopy</i> , 2011, 111, 1688-1695.	1.9	24
47	Practical factors affecting the performance of a thin-film phase plate for transmission electron microscopy. <i>Ultramicroscopy</i> , 2009, 109, 312-325.	1.9	116
48	Restoration of weak phase-contrast images recorded with a high degree of defocus: The "twin image" problem associated with CTF correction. <i>Ultramicroscopy</i> , 2008, 108, 921-928.	1.9	34
49	Retrospective: Radiation damage and its associated "Information Limitations". <i>Journal of Structural Biology</i> , 2008, 163, 271-276.	2.8	81
50	Retrospective on the early development of cryoelectron microscopy of macromolecules and a prospective on opportunities for the future. <i>Journal of Structural Biology</i> , 2008, 163, 214-223.	2.8	143
51	Macromolecular structures without crystals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1779-1780.	7.1	5
52	Aspects of Using a Boersch Type Phase Shifting Device for Contrast Enhancement in Macromolecular Electron Microscopy. <i>Microscopy and Microanalysis</i> , 2008, 14, 74-75.	0.4	1
53	Historical background: why is it important to improve automated particle selection methods?. <i>Journal of Structural Biology</i> , 2004, 145, 15-18.	2.8	23
54	A binary segmentation approach for boxing ribosome particles in cryo EM micrographs. <i>Journal of Structural Biology</i> , 2004, 145, 142-151.	2.8	27

#	ARTICLE	IF	CITATIONS
55	Crystal structures of bR(D85S) favor a model of bacteriorhodopsin as a hydroxyl-ion pump. FEBS Letters, 2004, 564, 301-306.	2.8	24
56	Specimen Charging on Thin Films with One Conducting Layer: Discussion of Physical Principles. Microscopy and Microanalysis, 2004, 10, 790-796.	0.4	58
57	Water structure as a function of temperature from X-ray scattering experiments and ab initio molecular dynamics. Physical Chemistry Chemical Physics, 2003, 5, 1981.	2.8	189
58	Crystallization of membrane proteins from media composed of connected-bilayer gels. Biopolymers, 2002, 66, 300-316.	2.4	9
59	Characterization of Conditions Required for X-Ray Diffraction Experiments with Protein Microcrystals. Biophysical Journal, 2000, 78, 3178-3185.	0.5	32
60	A high-quality x-ray scattering experiment on liquid water at ambient conditions. Journal of Chemical Physics, 2000, 113, 9140-9148.	3.0	280
61	What can x-ray scattering tell us about the radial distribution functions of water?. Journal of Chemical Physics, 2000, 113, 9149-9161.	3.0	381
62	Solution X-ray scattering as a probe of hydration-dependent structuring of aqueous solutions. Journal of Computer - Aided Molecular Design, 1999, 17, 97-118.	1.0	30
63	Review: Electron Crystallography: Present Excitement, a Nod to the Past, Anticipating the Future. Journal of Structural Biology, 1999, 128, 3-14.	2.8	137
64	Combining noisy images of small crystalline domains in high resolution electron microscopy. Journal of Applied Statistics, 1989, 16, 165-175.	1.3	10
65	Limitations to significant information in biological electron microscopy as a result of radiation damage. Journal of Ultrastructure Research, 1971, 36, 466-482.	1.1	348