

Ming Sun

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4508556/publications.pdf>

Version: 2024-02-01

19
papers

1,631
citations

687363

13
h-index

940533

16
g-index

28
all docs

28
docs citations

28
times ranked

3807
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative host-coronavirus protein interaction networks reveal pan-viral disease mechanisms. <i>Science</i> , 2020, 370, .	12.6	508
2	An ultrapotent synthetic nanobody neutralizes SARS-CoV-2 by stabilizing inactive Spike. <i>Science</i> , 2020, 370, 1473-1479.	12.6	336
3	A Fast and Effective Microfluidic Spraying-Plunging Method for High-Resolution Single-Particle Cryo-EM. <i>Structure</i> , 2017, 25, 663-670.e3.	3.3	112
4	Late steps in bacterial translation initiation visualized using time-resolved cryo-EM. <i>Nature</i> , 2019, 570, 400-404.	27.8	103
5	Structural Dynamics of Ribosome Subunit Association Studied by Mixing-Spraying Time-Resolved Cryogenic Electron Microscopy. <i>Structure</i> , 2015, 23, 1097-1105.	3.3	78
6	Key Intermediates in Ribosome Recycling Visualized by Time-Resolved Cryoelectron Microscopy. <i>Structure</i> , 2016, 24, 2092-2101.	3.3	68
7	Structure and assembly model for the <i>Trypanosoma cruzi</i> 60S ribosomal subunit. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 12174-12179.	7.1	63
8	Structural basis for the function of a small GTPase RsgA on the 30S ribosomal subunit maturation revealed by cryoelectron microscopy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 13100-13105.	7.1	57
9	Dynamical features of the <i>Plasmodium falciparum</i> ribosome during translation. <i>Nucleic Acids Research</i> , 2015, 43, gkv991.	14.5	48
10	The structural basis for release-factor activation during translation termination revealed by time-resolved cryogenic electron microscopy. <i>Nature Communications</i> , 2019, 10, 2579.	12.8	43
11	<i>Escherichia coli</i> NusG Links the Lead Ribosome with the Transcription Elongation Complex. <i>Science</i> , 2020, 23, 101352.	4.1	43
12	Identification of Changing Ribosome Protein Compositions using Mass Spectrometry. <i>Proteomics</i> , 2018, 18, e1800217.	2.2	29
13	Determination of the ribosome structure to a resolution of 2.5 Å... by single-particle cryo-EM. <i>Protein Science</i> , 2017, 26, 82-92.	7.6	26
14	A Time-Resolved Cryo-EM Study of <i>Saccharomyces cerevisiae</i> 80S Ribosome Protein Composition in Response to a Change in Carbon Source. <i>Proteomics</i> , 2021, 21, 2000125.	2.2	7
15	Critical Role for <i>Saccharomyces cerevisiae</i> Asc1p in Translational Initiation at Elevated Temperatures. <i>Proteomics</i> , 2018, 18, e1800208.	2.2	4
16	The Structural Basis for Release Factor Activation during Translation Termination Revealed by Time-Resolved Cryogenic Electron Microscopy. <i>Biophysical Journal</i> , 2019, 116, 574a-575a.	0.5	2
17	A Time-Resolved CRYO-EM Study of Ribosome Subunit Association by Mixing-Spraying. <i>Biophysical Journal</i> , 2014, 106, 598a.	0.5	0
18	Time-Resolved cryo-EM Study of Ribosome Subunit Association by Mixing-Spraying. <i>Biophysical Journal</i> , 2015, 108, 619a.	0.5	0

#	ARTICLE	IF	CITATIONS
19	Key Intermediates in Ribosome Recycling Visualized by Time-Resolved Cryoelectron Microscopy. journal of hand surgery Asian-Pacific volume, The, 2018, , 516-525.	0.4	0