

Stefan Bohn

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2010505/publications.pdf>

Version: 2024-02-01

16
papers

1,962
citations

687220

13
h-index

1058333

14
g-index

18
all docs

18
docs citations

18
times ranked

2632
citing authors

#	ARTICLE	IF	CITATIONS
1	High-resolution structure and biophysical characterization of the nucleocapsid phosphoprotein dimerization domain from the Covid-19 severe acute respiratory syndrome coronavirus 2. <i>Biochemical and Biophysical Research Communications</i> , 2021, 538, 54-62.	1.0	100
2	Protocol for RNA-seq Expression Analysis in Yeast. <i>Bio-protocol</i> , 2021, 11, e4161.	0.2	0
3	Structural insights into photosystem II assembly. <i>Nature Plants</i> , 2021, 7, 524-538.	4.7	102
4	Cryo-EM structure of the cetacean morbillivirus nucleoprotein-RNA complex. <i>Journal of Structural Biology</i> , 2021, 213, 107750.	1.3	12
5	Genetic interaction mapping informs integrative structure determination of protein complexes. <i>Science</i> , 2020, 370, .	6.0	24
6	The 26S Proteasome. , 2013, , 3691-3700.		0
7	Localization of the regulatory particle subunit Sem1 in the 26S proteasome. <i>Biochemical and Biophysical Research Communications</i> , 2013, 435, 250-254.	1.0	28
8	The proteasomal subunit Rpn6 is a molecular clamp holding the core and regulatory subcomplexes together. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 149-154.	3.3	136
9	Localization of the proteasomal ubiquitin receptors Rpn10 and Rpn13 by electron cryomicroscopy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 1479-1484.	3.3	114
10	Near-atomic resolution structural model of the yeast 26S proteasome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 14870-14875.	3.3	242
11	Expanding the Chemical Cross-Linking Toolbox by the Use of Multiple Proteases and Enrichment by Size Exclusion Chromatography. <i>Molecular and Cellular Proteomics</i> , 2012, 11, M111.014126.	2.5	264
12	Molecular architecture of the 26S proteasome holocomplex determined by an integrative approach. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 1380-1387.	3.3	422
13	False discovery rate estimation for cross-linked peptides identified by mass spectrometry. <i>Nature Methods</i> , 2012, 9, 901-903.	9.0	273
14	Positioning of large organelles by a membrane-associated cytoskeleton in <i>Plasmodium</i> sporozoites. <i>Cellular Microbiology</i> , 2010, 12, 362-371.	1.1	74
15	Structure of the 26S proteasome from <i>Schizosaccharomyces pombe</i> at subnanometer resolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 20992-20997.	3.3	130
16	The Vaccinia Virus F11L Gene Product Facilitates Cell Detachment and Promotes Migration. <i>Traffic</i> , 2008, 9, 1283-1298.	1.3	34