

Qiang Guo

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19
papers

758
citations

11
h-index

24
g-index

24
ext. papers

1,070
ext. citations

17.5
avg. IF

3.69
L-index

#	Paper	IF	Citations
19	A feature-guided, focused 3D signal permutation method for subtomogram averaging.. <i>Journal of Structural Biology</i> , 2022 , 107851	3.4	0
18	Gel-like inclusions of C-terminal fragments of TDP-43 sequester stalled proteasomes in neurons.. <i>EMBO Reports</i> , 2022 , e53890	6.5	1
17	In situ architecture of neuronal β synuclein inclusions. <i>Nature Communications</i> , 2021 , 12, 2110	17.4	24
16	Pathological polyQ expansion does not alter the conformation of the Huntingtin-HAP40 complex. <i>Structure</i> , 2021 , 29, 804-809.e5	5.2	3
15	The evolution of the huntingtin-associated protein 40 (HAP40) in conjunction with huntingtin. <i>BMC Evolutionary Biology</i> , 2020 , 20, 162	3	4
14	Stress- and ubiquitylation-dependent phase separation of the proteasome. <i>Nature</i> , 2020 , 578, 296-300	50.4	92
13	The cryo-electron microscopy structure of huntingtin. <i>Nature</i> , 2018 , 555, 117-120	50.4	70
12	Plant G proteins interact with endoplasmic reticulum luminal protein receptors to regulate endoplasmic reticulum retrieval. <i>Journal of Integrative Plant Biology</i> , 2018 , 60, 541-561	8.3	5
11	In Situ Structure of Neuronal C9orf72 Poly-GA Aggregates Reveals Proteasome Recruitment. <i>Cell</i> , 2018 , 172, 696-705.e12	56.2	196
10	Feature Decomposition Based Saliency Detection in Electron Cryo-Tomograms 2018 , 2018, 2467-2473	0.8	4
9	An integration of fast alignment and maximum-likelihood methods for electron subtomogram averaging and classification. <i>Bioinformatics</i> , 2018 , 34, i227-i236	7.2	4
8	Cryo-EM structures of the 80S ribosomes from human parasites <i>Trichomonas vaginalis</i> and <i>Toxoplasma gondii</i> . <i>Cell Research</i> , 2017 , 27, 1275-1288	24.7	12
7	CapZ regulates autophagosomal membrane shaping by promoting actin assembly inside the isolation membrane. <i>Nature Cell Biology</i> , 2015 , 17, 1112-23	23.4	84
6	Structural insights into the assembly of the 30S ribosomal subunit in vivo: functional role of S5 and location of the 17S rRNA precursor sequence. <i>Protein and Cell</i> , 2014 , 5, 394-407	7.2	20
5	Structural and functional insights into the mode of action of a universally conserved Obg GTPase. <i>PLoS Biology</i> , 2014 , 12, e1001866	9.7	91
4	Dissecting the in vivo assembly of the 30S ribosomal subunit reveals the role of RimM and general features of the assembly process. <i>Nucleic Acids Research</i> , 2013 , 41, 2609-20	20.1	56
3	Cryo-EM structures of the late-stage assembly intermediates of the bacterial 50S ribosomal subunit. <i>Nucleic Acids Research</i> , 2013 , 41, 7073-83	20.1	42

2 Structural basis for the function of a small GTPase RsgA on the 30S ribosomal subunit maturation revealed by cryoelectron microscopy. *Proceedings of the National Academy of Sciences of the United States of America*, **2011**, 108, 13100-5 11.5 47

1 Gel-like inclusions of C-terminal fragments of TDP-43 sequester and inhibit proteasomes in neurons 1