## **Timothy Grant**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4720309/publications.pdf

Version: 2024-02-01

16 2,589 14 17
papers citations h-index g-index

24 24 24 4499
all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Three-dimensional structure determination of protein complexes using matrix-landing mass spectrometry. Nature Communications, 2022, 13, 2276.	5.8	31
2	Locating macromolecular assemblies in cells by 2D template matching with cisTEM. ELife, 2021, $10$ , .	2.8	55
3	Neural networks learn the motions of molecular machines. Nature Methods, 2021, 18, 869-871.	9.0	5
4	Structure and assembly of calcium homeostasis modulator proteins. Nature Structural and Molecular Biology, 2020, 27, 150-159.	3.6	55
5	In situ Structure of Rotavirus VP1 RNA-Dependent RNA Polymerase. Journal of Molecular Biology, 2019, 431, 3124-3138.	2.0	45
6	High-resolution cryo-EM structures of outbreak strain human norovirus shells reveal size variations. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 12828-12832.	3.3	53
7	Structural basis for Fullerene geometry in a human endogenous retrovirus capsid. Nature Communications, 2019, 10, 5822.	5.8	20
8	Structural Mechanism of Functional Modulation by Gene Splicing in NMDA Receptors. Neuron, 2018, 98, 521-529.e3.	3.8	57
9	cisTEM, user-friendly software for single-particle image processing. ELife, 2018, 7, .	2.8	792
10	CryoEM structures of open dimers of gyrase A in complex with DNA illuminate mechanism of strand passage. ELife, 2018, 7, .	2.8	26
11	Mechanism of ribosome rescue by ArfA and RF2. ELife, 2017, 6, .	2.8	50
12	Activation of NMDA receptors and the mechanism of inhibition by ifenprodil. Nature, 2016, 534, 63-68.	13.7	173
13	Ensemble cryo-EM uncovers inchworm-like translocation of a viral IRES through the ribosome. ELife, 2016, 5, .	2.8	116
14	Measuring the optimal exposure for single particle cryo-EM using a 2.6 $\tilde{A}$ reconstruction of rotavirus VP6. ELife, 2015, 4, e06980.	2.8	748
15	Structure of the L Protein of Vesicular Stomatitis Virus from Electron Cryomicroscopy. Cell, 2015, 162, 314-327.	13.5	211
16	Automatic estimation and correction of anisotropic magnification distortion in electron microscopes. Journal of Structural Biology, 2015, 192, 204-208.	1.3	124