

# Rub n Fern ndez-Busnadiego

## List of Publications by Year in descending order

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33  
papers

2,323  
citations

361045

20  
h-index

454577

30  
g-index

42  
all docs

42  
docs citations

42  
times ranked

3297  
citing authors

#	ARTICLE	IF	CITATIONS
1	In Situ Structure of Neuronal C9orf72 Poly-GA Aggregates Reveals Proteasome Recruitment. <i>Cell</i> , 2018, 172, 696-705.e12.	13.5	311
2	In Situ Architecture and Cellular Interactions of PolyQ Inclusions. <i>Cell</i> , 2017, 171, 179-187.e10.	13.5	271
3	Quantitative analysis of the native presynaptic cytomatrix by cryoelectron tomography. <i>Journal of Cell Biology</i> , 2010, 188, 145-156.	2.3	209
4	Stress- and ubiquitylation-dependent phase separation of the proteasome. <i>Nature</i> , 2020, 578, 296-300.	13.7	204
5	Three-dimensional architecture of extended synaptotagmin-mediated endoplasmic reticulumâ€“plasma membrane contact sites. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E2004-13.	3.3	185
6	The cryo-electron microscopy structure of huntingtin. <i>Nature</i> , 2018, 555, 117-120.	13.7	125
7	Synucleins Have Multiple Effects on Presynaptic Architecture. <i>Cell Reports</i> , 2017, 18, 161-173.	2.9	120
8	Cryoâ€“electron tomography reveals a critical role of RIM1Î± in synaptic vesicle tethering. <i>Journal of Cell Biology</i> , 2013, 201, 725-740.	2.3	110
9	Epsin deficiency impairs endocytosis by stalling the actin-dependent invagination of endocytic clathrin-coated pits. <i>ELife</i> , 2014, 3, e03311.	2.8	101
10	Tricalbin-Mediated Contact Sites Control ER Curvature to Maintain Plasma Membrane Integrity. <i>Developmental Cell</i> , 2019, 51, 476-487.e7.	3.1	87
11	Molecular and structural architecture of polyQ aggregates in yeast. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E3446-E3453.	3.3	68
12	In situ architecture of neuronal Î±-Synuclein inclusions. <i>Nature Communications</i> , 2021, 12, 2110.	5.8	66
13	Conformation of Pseudoazurin in the 152ÅkDa Electron Transfer Complex with Nitrite Reductase Determined by Paramagnetic NMR. <i>Journal of Molecular Biology</i> , 2008, 375, 1405-1415.	2.0	64
14	Cryoâ€“electron tomographyâ€“the cell biology that came in from the cold. <i>FEBS Letters</i> , 2017, 591, 2520-2533.	1.3	56
15	Dynamic instability of clathrin assembly provides proofreading control for endocytosis. <i>Journal of Cell Biology</i> , 2019, 218, 3200-3211.	2.3	41
16	Insights into the molecular organization of the neuron by cryo-electron tomography. <i>Microscopy (Oxford, England)</i> , 2011, 60, S137-S148.	0.7	35
17	Deciphering the molecular architecture of membrane contact sites by cryo-electron tomography. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2017, 1864, 1507-1512.	1.9	29
18	Cnm1 mediates nucleusâ€“mitochondria contact site formation in response to phospholipid levels. <i>Journal of Cell Biology</i> , 2021, 220, .	2.3	29

#	ARTICLE	IF	CITATIONS
19	Gel-like inclusions of C-terminal fragments of TDP43 sequester stalled proteasomes in neurons. <i>EMBO Reports</i> , 2022, 23, e53890.	2.0	28
20	Expression of DNAJB12 or DNAJB14 Causes Coordinate Invasion of the Nucleus by Membranes Associated with a Novel Nuclear Pore Structure. <i>PLoS ONE</i> , 2014, 9, e94322.	1.1	26
21	Hierarchical detection and analysis of macromolecular complexes in cryo-electron tomograms using Pyto software. <i>Journal of Structural Biology</i> , 2016, 196, 503-514.	1.3	26
22	Investigating the Structure of Neurotoxic Protein Aggregates Inside Cells. <i>Trends in Cell Biology</i> , 2020, 30, 951-966.	3.6	24
23	Reliable estimation of membrane curvature for cryo-electron tomography. <i>PLoS Computational Biology</i> , 2020, 16, e1007962.	1.5	23
24	Supramolecular architecture of endoplasmic reticulum-plasma membrane contact sites. <i>Biochemical Society Transactions</i> , 2016, 44, 534-540.	1.6	13
25	Amyloid-like aggregating proteins cause lysosomal defects in neurons via gain-of-function toxicity. <i>Life Science Alliance</i> , 2022, 5, e202101185.	1.3	13
26	The evolution of the huntingtin-associated protein 40 (HAP40) in conjunction with huntingtin. <i>BMC Evolutionary Biology</i> , 2020, 20, 162.	3.2	11
27	Pathological polyQ expansion does not alter the conformation of the Huntingtin-HAP40 complex. <i>Structure</i> , 2021, 29, 804-809.e5.	1.6	8
28	Lipoprotein-like particles in a prokaryote: quinone droplets of <i>Thermoplasma acidophilum</i> . <i>FEMS Microbiology Letters</i> , 2016, 363, fnw169.	0.7	4
29	Cryo-Electron Tomography of the Mammalian Synapse. <i>Methods in Molecular Biology</i> , 2018, 1847, 217-224.	0.4	3
30	Quantitative Synaptic Biology: A Perspective on Techniques, Numbers and Expectations. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7298.	1.8	3
31	Tricalbin-Mediated Contact Sites Control ER Curvature to Maintain Plasma Membrane Integrity. <i>SSRN Electronic Journal</i> , 0, , .	0.4	2
32	The Cell at Molecular Resolution. , 2012, , 141-183.		0
33	High-Resolution Insights Into Neurodegeneration. , 2018, , .		0