

K Tanuj Sapra

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/930797/publications.pdf>

Version: 2024-02-01

39
papers

1,863
citations

304368

22
h-index

360668

35
g-index

40
all docs

40
docs citations

40
times ranked

2206
citing authors

#	ARTICLE	IF	CITATIONS
1	The molecular architecture of lamins in somatic cells. <i>Nature</i> , 2017, 543, 261-264.	13.7	339
2	Differentiation of Cytoplasmic and Meiotic Spindle Assembly MCAK Functions by Aurora B-dependent Phosphorylation. <i>Molecular Biology of the Cell</i> , 2004, 15, 2895-2906.	0.9	202
3	Deciphering Molecular Interactions of Native Membrane Proteins by Single-Molecule Force Spectroscopy. <i>Annual Review of Biophysics and Biomolecular Structure</i> , 2007, 36, 233-260.	18.3	124
4	Single-molecule studies of membrane proteins. <i>Current Opinion in Structural Biology</i> , 2006, 16, 489-495.	2.6	102
5	Characterizing Molecular Interactions in Different Bacteriorhodopsin Assemblies by Single-molecule Force Spectroscopy. <i>Journal of Molecular Biology</i> , 2006, 355, 640-650.	2.0	93
6	Structural analysis of multicellular organisms with cryo-electron tomography. <i>Nature Methods</i> , 2015, 12, 634-636.	9.0	85
7	Detecting Molecular Interactions that Stabilize Native Bovine Rhodopsin. <i>Journal of Molecular Biology</i> , 2006, 358, 255-269.	2.0	71
8	Multi-compartment encapsulation of communicating droplets and droplet networks in hydrogel as a model for artificial cells. <i>Scientific Reports</i> , 2017, 7, 45167.	1.6	66
9	Stabilizing Effect of Zn ²⁺ in Native Bovine Rhodopsin. <i>Journal of Biological Chemistry</i> , 2007, 282, 11377-11385.	1.6	61
10	Roll, adhere, spread and contract: Structural mechanics of platelet function. <i>European Journal of Cell Biology</i> , 2015, 94, 129-138.	1.6	56
11	Point Mutations in Membrane Proteins Reshape Energy Landscape and Populate Different Unfolding Pathways. <i>Journal of Molecular Biology</i> , 2008, 376, 1076-1090.	2.0	52
12	Construction and Manipulation of Functional Three-Dimensional Droplet Networks. <i>ACS Nano</i> , 2014, 8, 771-779.	7.3	52
13	An engineered dimeric protein pore that spans adjacent lipid bilayers. <i>Nature Communications</i> , 2013, 4, 1725.	5.8	44
14	From Valleys to Ridges: Exploring the Dynamic Energy Landscape of Single Membrane Proteins. <i>ChemPhysChem</i> , 2008, 9, 954-966.	1.0	43
15	Mechanical Properties of Bovine Rhodopsin and Bacteriorhodopsin: Possible Roles in Folding and Function. <i>Langmuir</i> , 2008, 24, 1330-1337.	1.6	43
16	Nonlinear mechanics of lamin filaments and the meshwork topology build an emergent nuclear lamina. <i>Nature Communications</i> , 2020, 11, 6205.	5.8	40
17	Imaging and detecting molecular interactions of single transmembrane proteins. <i>Neurobiology of Aging</i> , 2006, 27, 546-561.	1.5	38
18	One Hairpin after the Other: Exploring Mechanical Unfolding Pathways of the Transmembrane Barrel Protein OmpG. <i>Angewandte Chemie - International Edition</i> , 2009, 48, 8306-8308.	7.2	38

#	ARTICLE	IF	CITATIONS
19	Lipid-coated hydrogel shapes as components of electrical circuits and mechanical devices. <i>Scientific Reports</i> , 2012, 2, 848.	1.6	37
20	Modulation of Molecular Interactions and Function by Rhodopsin Palmitylation. <i>Biochemistry</i> , 2009, 48, 4294-4304.	1.2	31
21	A novel pattern recognition algorithm to classify membrane protein unfolding pathways with high-throughput single-molecule force spectroscopy. <i>Bioinformatics</i> , 2007, 23, e231-e236.	1.8	30
22	Role of Extracellular Glutamic Acids in the Stability and Energy Landscape of Bacteriorhodopsin. <i>Biophysical Journal</i> , 2008, 95, 3407-3418.	0.2	23
23	Toward correlating structure and mechanics of platelets. <i>Cell Adhesion and Migration</i> , 2016, 10, 568-575.	1.1	23
24	Developments in cryo-electron tomography for in situ structural analysis. <i>Archives of Biochemistry and Biophysics</i> , 2015, 581, 78-85.	1.4	22
25	Single-Molecule Force Spectroscopy of Transmembrane β -Barrel Proteins. <i>Annual Review of Analytical Chemistry</i> , 2018, 11, 375-395.	2.8	21
26	The macromolecular architecture of platelet-derived microparticles. <i>Journal of Structural Biology</i> , 2016, 193, 181-187.	1.3	19
27	Evolving protocells to prototissues: rational design of a missing link. <i>Biochemical Society Transactions</i> , 2013, 41, 1159-1165.	1.6	18
28	Seeing and sensing single G protein-coupled receptors by atomic force microscopy. <i>Current Opinion in Cell Biology</i> , 2019, 57, 25-32.	2.6	18
29	Dual energy landscape: The functional state of the β -barrel outer membrane protein G molds its unfolding energy landscape. <i>Proteomics</i> , 2010, 10, 4151-4162.	1.3	16
30	Structural Properties of the Human Protease-Activated Receptor 1 Changing by a Strong Antagonist. <i>Structure</i> , 2018, 26, 829-838.e4.	1.6	13
31	Bend, Push, Stretch: Remarkable Structure and Mechanics of Single Intermediate Filaments and Meshworks. <i>Cells</i> , 2021, 10, 1960.	1.8	13
32	Profilin α -mediated cytoskeletal rearrangements regulate integrin function in mouse platelets. <i>Blood Advances</i> , 2018, 2, 1040-1045.	2.5	12
33	Conformational Plasticity of Human Protease-Activated Receptor 1 upon Antagonist- and Agonist-Binding. <i>Structure</i> , 2019, 27, 1517-1526.e3.	1.6	8
34	Complex Stability of Single Proteins Explored by Forced Unfolding Experiments. <i>Biophysical Journal</i> , 2005, 88, L37-L39.	0.2	5
35	Atomic Force Microscopy and Spectroscopy to Probe Single Membrane Proteins in Lipid Bilayers. <i>Methods in Molecular Biology</i> , 2013, 974, 73-110.	0.4	3
36	Imaging and Force Spectroscopy of Single Transmembrane Proteins with the Atomic Force Microscope. <i>Methods in Molecular Biology</i> , 2019, 2003, 107-144.	0.4	0

#	ARTICLE	IF	CITATIONS
37	A Structure-Based Analysis of Single Molecule Force Spectroscopy (SMFS) Data for Bacteriorhodopsin and Four Mutants. Lecture Notes in Computer Science, 2006, , 162-172.	1.0	0
38	Probing Single Membrane Proteins by Atomic Force Microscopy. , 2009, , 449-485.		0
39	Pattern Recognition of Single-Molecule Force Spectroscopy Data. , 2007, , 3-13.		0