## Soung-Hun Roh

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9025157/publications.pdf

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27 1,021 15 21 g-index

29 29 29 1754

29 29 29 1754 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Cryo-EM structures of GroEL:ES2 with RuBisCO visualize molecular contacts of encapsulated substrates in a double-cage chaperonin. IScience, 2022, 25, 103704.	1.9	5
2	Oxidative stress protein Oxr1 promotes Vâ€ATPase holoenzyme disassembly in catalytic activityâ€independent manner. EMBO Journal, 2022, 41, e109360.	3.5	15
3	Structural insights into ClpP protease side exit poreâ€opening by a pH drop coupled with substrate hydrolysis. EMBO Journal, 2022, 41, e109755.	3.5	8
4	The N-terminus of varicella-zoster virus glycoprotein B has a functional role in fusion. PLoS Pathogens, 2021, 17, e1008961.	2.1	12
5	Mapping the catalytic conformations of an assembly-line polyketide synthase module. Science, 2021, 374, 729-734.	6.0	41
6	Integrated cryoEM imaging center at Seoul National University, Korea. Biodesign, 2021, 9, 72-77.	0.2	0
7	The N-terminus of varicella-zoster virus glycoprotein B has a functional role in fusion. , 2021, 17, e1008961.		O
8	The N-terminus of varicella-zoster virus glycoprotein B has a functional role in fusion. , 2021, 17, e1008961.		0
9	The N-terminus of varicella-zoster virus glycoprotein B has a functional role in fusion. , 2021, 17, e1008961.		O
10	The N-terminus of varicella-zoster virus glycoprotein B has a functional role in fusion. , 2021, 17, e1008961.		0
11	Cryo-EM and MD infer water-mediated proton transport and autoinhibition mechanisms of V <sub>o</sub> complex. Science Advances, 2020, 6, .	4.7	51
12	A glycoprotein B-neutralizing antibody structure at 2.8 à uncovers a critical domain for herpesvirus fusion initiation. Nature Communications, 2020, 11, 4141.	5.8	23
13	Cryo-EM structures of NPC1L1 reveal mechanisms of cholesterol transport and ezetimibe inhibition. Science Advances, 2020, 6, eabb1989.	4.7	49
14	Cryo-EM Structures of the Hsp104 Protein Disaggregase Captured in the ATP Conformation. Cell Reports, 2019, 26, 29-36.e3.	2.9	36
15	The 3.5-Ã CryoEM Structure of Nanodisc-Reconstituted Yeast Vacuolar ATPase Vo Proton Channel. Molecular Cell, 2018, 69, 993-1004.e3.	4.5	103
16	Structural principles of distinct assemblies of the human $\hat{l}\pm4\hat{l}^22$ nicotinic receptor. Nature, 2018, 557, 261-265.	13.7	177
17	Resolution of Heterogeneity in Nicotinic Receptor Assembly by Cryo-EM. Biophysical Journal, 2018, 114, 10a-11a.	0.2	0
18	Subunit conformational variation within individual GroEL oligomers resolved by Cryo-EM. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8259-8264.	3.3	86

#	ARTICLE	IF	CITATIONS
19	Chaperonin TRiC/CCT Recognizes Fusion Oncoprotein AML1-ETO through Subunit-Specific Interactions. Biophysical Journal, 2016, 110, 2377-2385.	0.2	12
20	Chaperonin TRiC/CCT Modulates the Folding and Activity of Leukemogenic Fusion Oncoprotein AML1-ETO. Journal of Biological Chemistry, 2016, 291, 4732-4741.	1.6	25
21	Contribution of the Type II Chaperonin, TRiC/CCT, to Oncogenesis. International Journal of Molecular Sciences, 2015, 16, 26706-26720.	1.8	65
22	Improved Peak Detection and Deconvolution of Native Electrospray Mass Spectra from Large Protein Complexes. Journal of the American Society for Mass Spectrometry, 2015, 26, 2141-2151.	1.2	49
23	Contribution of chaperones to STAT pathway signaling. Jak-stat, 2014, 3, e970459.	2.2	24
24	Modulation of STAT3 Folding and Function by TRiC/CCT Chaperonin. PLoS Biology, 2014, 12, e1001844.	2.6	84
25	Reprogramming an ATP-Driven Biological Machine into a Light-Gated Protein Nanocage. Biophysical Journal, 2014, 106, 439a.	0.2	0
26	Reprogramming an ATP-driven protein machine into a light-gated nanocage. Nature Nanotechnology, 2013, 8, 928-932.	15.6	55
27	Molecular Surface of Tarantula Toxins Interacting with Voltage Sensors in Kv Channels. Journal of General Physiology, 2004, 123, 455-467.	0.9	100