

Kliment A Verba

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3467436/publications.pdf>

Version: 2024-02-01

13
papers

11,185
citations

933447

10
h-index

1199594

12
g-index

21
all docs

21
docs citations

21
times ranked

20827
citing authors

#	ARTICLE	IF	CITATIONS
1	Cryo-EM structures of the near full-length HER2/HER3 heterodimer reveal a novel allosteric mechanism of activation. <i>Biophysical Journal</i> , 2022, 121, 84a.	0.5	0
2	Efficient expression, purification, and visualization by cryo-EM of unliganded near full-length HER3. <i>Methods in Enzymology</i> , 2022, 667, 611-632.	1.0	0
3	Evolution of enhanced innate immune evasion by SARS-CoV-2. <i>Nature</i> , 2022, 602, 487-495.	27.8	237
4	An effective strategy for ligand-mediated pulldown of the HER2/HER3/NGR1 ^{Δ2} heterocomplex and cryo-EM structure determination at low sample concentrations. <i>Methods in Enzymology</i> , 2022, 667, 633-662.	1.0	2
5	Practical considerations for using K3 cameras in CDS mode for high-resolution and high-throughput single particle cryo-EM. <i>Journal of Structural Biology</i> , 2021, 213, 107745.	2.8	33
6	Drugging the "Undruggable" MYCN Oncogenic Transcription Factor: Overcoming Previous Obstacles to Impact Childhood Cancers. <i>Cancer Research</i> , 2021, 81, 1627-1632.	0.9	25
7	Structures of the HER2-HER3-NGR1 ^{Δ2} complex reveal a dynamic dimer interface. <i>Nature</i> , 2021, 600, 339-343.	27.8	48
8	Comparative host-coronavirus protein interaction networks reveal pan-viral disease mechanisms. <i>Science</i> , 2020, 370, .	12.6	508
9	A SARS-CoV-2 protein interaction map reveals targets for drug repurposing. <i>Nature</i> , 2020, 583, 459-468.	27.8	3,542
10	MotionCor2: anisotropic correction of beam-induced motion for improved cryo-electron microscopy. <i>Nature Methods</i> , 2017, 14, 331-332.	19.0	6,166
11	How Hsp90 and Cdc37 Lubricate Kinase Molecular Switches. <i>Trends in Biochemical Sciences</i> , 2017, 42, 799-811.	7.5	77
12	Atomic structure of Hsp90-Cdc37-Cdk4 reveals that Hsp90 traps and stabilizes an unfolded kinase. <i>Science</i> , 2016, 352, 1542-1547.	12.6	354
13	Cross-Monomer Substrate Contacts Reposition the Hsp90 N-Terminal Domain and Prime the Chaperone Activity. <i>Journal of Molecular Biology</i> , 2012, 415, 3-15.	4.2	45