

Ranabir Das

List of Publications by Year in descending order

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33
papers

738
citations

759233

12
h-index

552781

26
g-index

42
all docs

42
docs citations

42
times ranked

1130
citing authors

#	ARTICLE	IF	CITATIONS
1	Epidemiological and ES cell-based functional evaluation of BRCA2 variants identified in families with breast cancer. <i>Human Mutation</i> , 2021, 42, 200-212.	2.5	4
2	Non-covalent Interaction With SUMO Enhances the Activity of Human Cytomegalovirus Protein IE1. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 662522.	3.7	3
3	NEDD8 Deamidation Inhibits Cullin RING Ligase Dynamics. <i>Frontiers in Immunology</i> , 2021, 12, 695331.	4.8	5
4	An α -oriented methionine-aromatic structural motif in SUMO is critical for its stability and activity. <i>Journal of Biological Chemistry</i> , 2021, 297, 100970.	3.4	4
5	Destabilization of polar interactions in the prion protein triggers misfolding and oligomerization. <i>Protein Science</i> , 2021, 30, 2258-2271.	7.6	5
6	Rational Design of Protein-Specific Folding Modifiers. <i>Journal of the American Chemical Society</i> , 2021, 143, 18766-18776.	13.7	6
7	Stability of Begomoviral pathogenicity determinant \hat{I}^2C1 is modulated by mutually antagonistic SUMOylation and SIM interactions. <i>BMC Biology</i> , 2020, 18, 110.	3.8	12
8	Monitoring protein ubiquitination and SUMOylation in real-time by NMR. <i>Chemical Communications</i> , 2020, 56, 6735-6738.	4.1	2
9	A Fluorescence-Based Assay to Monitor SUMOylation in Real-Time. <i>Current Protocols in Protein Science</i> , 2020, 101, e111.	2.8	1
10	Amide temperature coefficients in characterizing the allosteric effects of ligand binding on local stability in proteins. <i>Biochemical and Biophysical Research Communications</i> , 2020, 524, 677-682.	2.1	3
11	The Viral SUMO-Targeted Ubiquitin Ligase ICPO is Phosphorylated and Activated by Host Kinase Chk2. <i>Journal of Molecular Biology</i> , 2020, 432, 1952-1977.	4.2	15
12	Genetically encoded live-cell sensor for tyrosinated microtubules. <i>Journal of Cell Biology</i> , 2020, 219, .	5.2	20
13	A Fyn biosensor reveals pulsatile, spatially localized kinase activity and signaling crosstalk in live mammalian cells. <i>ELife</i> , 2020, 9, .	6.0	14
14	A novel polyubiquitin chain linkage formed by viral Ubiquitin is resistant to host deubiquitinating enzymes. <i>Biochemical Journal</i> , 2020, 477, 2193-2219.	3.7	2
15	Casein kinase-2-mediated phosphorylation increases the SUMO-dependent activity of the cytomegalovirus transactivator IE2. <i>Journal of Biological Chemistry</i> , 2019, 294, 14546-14561.	3.4	12
16	A five-residue motif for the design of domain swapping in proteins. <i>Nature Communications</i> , 2019, 10, 452.	12.8	37
17	A conserved and buried edge-to-face aromatic interaction in small ubiquitin-like modifier (SUMO) has a role in SUMO stability and function. <i>Journal of Biological Chemistry</i> , 2019, 294, 6772-6784.	3.4	17
18	Deamidation disrupts native and transient contacts to weaken the interaction between UBC13 and RING-finger E3 ligases. <i>ELife</i> , 2019, 8, .	6.0	11

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19	Indomethacin elicits proteasomal dysfunctions develops apoptosis through mitochondrial abnormalities. <i>Journal of Cellular Physiology</i> , 2018, 233, 1685-1699.	4.1	11
20	Conformational Dynamics and Allostery in E2:E3 Interactions Drive Ubiquitination: gp78 and Ube2g2. <i>Structure</i> , 2017, 25, 794-805.e5.	3.3	24
21	Salt-Mediated Oligomerization of the Mouse Prion Protein Monitored by Real-Time NMR. <i>Journal of Molecular Biology</i> , 2017, 429, 1852-1872.	4.2	26
22	Amino acid composition after loop deletion drives domain swapping. <i>Protein Science</i> , 2017, 26, 1994-2002.	7.6	13
23	Structural and functional analysis of SMO-1, the SUMO homolog in <i>Caenorhabditis elegans</i> . <i>PLoS ONE</i> , 2017, 12, e0186622.	2.5	5
24	Observing a late folding intermediate of Ubiquitin at atomic resolution by NMR. <i>Protein Science</i> , 2016, 25, 1438-1450.	7.6	10
25	Partially Unfolded Forms of the Prion Protein Populated under Misfolding-promoting Conditions. <i>Journal of Biological Chemistry</i> , 2015, 290, 25227-25240.	3.4	42
26	Allosteric regulation of E2:E3 interactions promote a processive ubiquitination machine. <i>EMBO Journal</i> , 2013, 32, 2504-2516.	7.8	82
27	A Structurally Unique E2-Binding Domain Activates Ubiquitination by the ERAD E2, Ubc7p, through Multiple Mechanisms. <i>Molecular Cell</i> , 2013, 50, 516-527.	9.7	71
28	Functional evaluation of BRCA2 variants mapping to the PALB2-binding and C-terminal DNA-binding domains using a mouse ES cell-based assay. <i>Human Molecular Genetics</i> , 2012, 21, 3993-4006.	2.9	56
29	Promiscuous Interactions of gp78 E3 Ligase CUE Domain with Polyubiquitin Chains. <i>Structure</i> , 2012, 20, 2138-2150.	3.3	32
30	Allosteric Activation of E2-RING Finger-Mediated Ubiquitylation by a Structurally Defined Specific E2-Binding Region of gp78. <i>Molecular Cell</i> , 2009, 34, 674-685.	9.7	144
31	Structural Biophysics of the NusB:NusE Antitermination Complex. <i>Journal of Molecular Biology</i> , 2008, 376, 705-720.	4.2	21
32	Spectral implementation of some quantum algorithms by one- and two-dimensional nuclear magnetic resonance. <i>Journal of Chemical Physics</i> , 2004, 121, 7601.	3.0	9
33	Experimental implementation of Grover's search algorithm using efficient quantum state tomography. <i>Chemical Physics Letters</i> , 2003, 369, 8-15.	2.6	15