

Rahul Das

List of Publications by Year in descending order

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32
papers

2,563
citations

361296

20
h-index

434063

31
g-index

34
all docs

34
docs citations

34
times ranked

3064
citing authors

#	ARTICLE	IF	CITATIONS
1	Mechanism for Activation of the EGF Receptor Catalytic Domain by the Juxtamembrane Segment. <i>Cell</i> , 2009, 137, 1293-1307.	13.5	506
2	Conformational Coupling across the Plasma Membrane in Activation of the EGF Receptor. <i>Cell</i> , 2013, 152, 543-556.	13.5	423
3	Architecture and Membrane Interactions of the EGF Receptor. <i>Cell</i> , 2013, 152, 557-569.	13.5	417
4	cAMP activation of PKA defines an ancient signaling mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 93-98.	3.3	113
5	Chemical genomics in <i>Escherichia coli</i> identifies an inhibitor of bacterial lipoprotein targeting. <i>Nature Chemical Biology</i> , 2009, 5, 849-856.	3.9	111
6	Understanding the Molecular Basis for the Inhibition of the Alzheimer's A β -Peptide Oligomerization by Human Serum Albumin Using Saturation Transfer Difference and Off-Resonance Relaxation NMR Spectroscopy. <i>Journal of the American Chemical Society</i> , 2007, 129, 4282-4290.	6.6	109
7	Regulation of the catalytic activity of the EGF receptor. <i>Current Opinion in Structural Biology</i> , 2011, 21, 777-784.	2.6	87
8	The Projection Analysis of NMR Chemical Shifts Reveals Extended EPAC Autoinhibition Determinants. <i>Biophysical Journal</i> , 2012, 102, 630-639.	0.2	83
9	Structural analysis of autoinhibition in the Ras-specific exchange factor RasGRP1. <i>ELife</i> , 2013, 2, e00813.	2.8	78
10	Analysis of the Role of the C-Terminal Tail in the Regulation of the Epidermal Growth Factor Receptor. <i>Molecular and Cellular Biology</i> , 2015, 35, 3083-3102.	1.1	74
11	Dynamically Driven Ligand Selectivity in Cyclic Nucleotide Binding Domains. <i>Journal of Biological Chemistry</i> , 2009, 284, 23682-23696.	1.6	71
12	The Auto-Inhibitory Role of the EPAC Hinge Helix as Mapped by NMR. <i>PLoS ONE</i> , 2012, 7, e48707.	1.1	63
13	Entropy-driven cAMP-dependent Allosteric Control of Inhibitory Interactions in Exchange Proteins Directly Activated by cAMP. <i>Journal of Biological Chemistry</i> , 2008, 283, 19691-19703.	1.6	59
14	Communication between Tandem cAMP Binding Domains in the Regulatory Subunit of Protein Kinase A-R1 \pm as Revealed by Domain-silencing Mutations. <i>Journal of Biological Chemistry</i> , 2010, 285, 15523-15537.	1.6	46
15	A Model for Agonism and Antagonism in an Ancient and Ubiquitous cAMP-binding Domain. <i>Journal of Biological Chemistry</i> , 2007, 282, 581-593.	1.6	41
16	Understanding cAMP-Dependent Allostery by NMR Spectroscopy: A Comparative Analysis of the EPAC1 cAMP-Binding Domain in Its Apo and cAMP-Bound States. <i>Journal of the American Chemical Society</i> , 2007, 129, 14482-14492.	6.6	41
17	Degradation of MAC13243 and studies of the interaction of resulting thiourea compounds with the lipoprotein targeting chaperone LolA. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2013, 23, 2426-2431.	1.0	39
18	Definition of an electrostatic relay switch critical for the cAMP-dependent activation of protein kinase A as revealed by the D170A mutant of R1 \pm . <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 69, 112-124.	1.5	37

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19	Mapping Allostery through Equilibrium Perturbation NMR Spectroscopy. <i>Journal of the American Chemical Society</i> , 2006, 128, 8406-8407.	6.6	34
20	A novel protein tyrosine phosphatase like phytase from <i>Lactobacillus fermentum</i> NKN51: Cloning, characterization and application in mineral release for food technology applications. <i>Bioresource Technology</i> , 2018, 249, 1000-1008.	4.8	21
21	Elucidating the regulation of glucose tolerance in a β -glucosidase from <i>Halothermothrix orenii</i> by active site pocket engineering and computational analysis. <i>International Journal of Biological Macromolecules</i> , 2020, 156, 621-632.	3.6	19
22	Mapping Polypeptide Self-Recognition through ^1H Off-Resonance Relaxation. <i>Journal of the American Chemical Society</i> , 2005, 127, 9358-9359.	6.6	18
23	Arginine-Modified Fluorescent Gold Nanoclusters for Förster Resonance Energy Transfer with a Hemicyanine Dye: A Biofriendly Approach. <i>ACS Applied Nano Materials</i> , 2021, 4, 305-312.	2.4	14
24	An allosteric hot spot in the tandem-SH2 domain of ZAP-70 regulates T-cell signaling. <i>Biochemical Journal</i> , 2020, 477, 1287-1308.	1.7	13
25	NMR assignment of the cAMP-binding domain A of the PKA regulatory subunit. <i>Journal of Biomolecular NMR</i> , 2006, 36, 64-64.	1.6	8
26	Dynamic unfolding of a regulatory subunit of cAMP-dependent protein kinase by capillary electrophoresis: Impact of cAMP dissociation on protein stability. <i>Electrophoresis</i> , 2006, 27, 4196-4204.	1.3	8
27	Development of Non-ionic Surfactant and Protein-Coated Ultrasmall Silver Nanoparticles: Increased Viscoelasticity Enables Potency in Biological Applications. <i>ACS Omega</i> , 2020, 5, 8999-9006.	1.6	8
28	Analysis and Parametric Optimization of ^1H Off-Resonance Relaxation NMR Experiments Designed to Map Polypeptide Self-Recognition and Other Noncovalent Interactions. <i>Journal of Physical Chemistry B</i> , 2006, 110, 20664-20670.	1.2	7
29	Cytotoxic Ruthenium(II) Complexes of Pyrazolylbenzimidazole Ligands That Inhibit VEGFR2 Phosphorylation. <i>Inorganic Chemistry</i> , 2021, 60, 18379-18394.	1.9	6
30	Regulating the discriminatory response to antigen by T-cell receptor. <i>Bioscience Reports</i> , 2022, 42, .	1.1	5
31	Selective targeting of the inactive state of hematopoietic cell kinase (Hck) with a stable curcumin derivative. <i>Journal of Biological Chemistry</i> , 2021, 296, 100449.	1.6	3
32	Backbone resonance assignment of the cAMP-binding domains of the protein kinase A regulatory subunit β . <i>Biomolecular NMR Assignments</i> , 2021, 15, 379-382.	0.4	0