

Tapas K Mal

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2694027/publications.pdf>

Version: 2024-02-01

35
papers

2,082
citations

361413

20
h-index

414414

32
g-index

35
all docs

35
docs citations

35
times ranked

2637
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Pathogen-specific antimicrobials engineered de novo through membrane-protein biomimicry. <i>Nature Biomedical Engineering</i> , 2021, 5, 467-480. | 22.5 | 17 |
| 2 | T3P-Promoted Synthesis of a Series of 2-Aryl-3-phenyl-2,3-dihydro-4H-pyrido[3,2-e][1,3]thiazin-4-ones and Their Activity against the Kinetoplastid Parasite <i>Trypanosoma brucei</i> . <i>Molecules</i> , 2021, 26, 6099. | 3.8 | 3 |
| 3 | Sample Preparation and Data Analysis for NMR-Based Metabolomics. <i>Methods in Molecular Biology</i> , 2021, 2194, 301-313. | 0.9 | 7 |
| 4 | T3P-promoted synthesis of a series of novel 2-aryl-3-phenyl-2,3,5,6-tetrahydro-4H-1,3-thiazin-4-ones. <i>Tetrahedron Letters</i> , 2020, 61, 151836. | 1.4 | 5 |
| 5 | NMR Spectroscopy Analysis of Asphaltenes. <i>Energy & Fuels</i> , 2019, 33, 10391-10414. | 5.1 | 41 |
| 6 | Structural characterization of NRAS isoform 5. <i>Protein Science</i> , 2016, 25, 1069-1074. | 7.6 | 5 |
| 7 | Abstract 3093: 3D structural report of NRAS isoform 5. , , | | 0 |
| 8 | Molecular basis of photochromism of a fluorescent protein revealed by direct ¹³ C detection under laser illumination. <i>Journal of Biomolecular NMR</i> , 2010, 48, 237-246. | 2.8 | 15 |
| 9 | Structural Studies of Inositol 1,4,5-Trisphosphate Receptor. <i>Journal of Biological Chemistry</i> , 2010, 285, 36092-36099. | 3.4 | 48 |
| 10 | Light-dependent regulation of structural flexibility in a photochromic fluorescent protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 9227-9232. | 7.1 | 150 |
| 11 | NMR Investigation of Calmodulin. , 2008, , 503-516. | | 3 |
| 12 | Functional Silencing of TATA-binding Protein (TBP) by a Covalent Linkage of the N-terminal Domain of TBP-associated Factor 1. <i>Journal of Biological Chemistry</i> , 2007, 282, 22228-22238. | 3.4 | 11 |
| 13 | Ligand-induced Conformational Changes via Flexible Linkers in the Amino-terminal region of the Inositol 1,4,5-Trisphosphate Receptor. <i>Journal of Molecular Biology</i> , 2007, 373, 1269-1280. | 4.2 | 46 |
| 14 | Characterization of Dual Substrate Binding Sites in the Homodimeric Structure of <i>Escherichia coli</i> mRNA Interferase MazF. <i>Journal of Molecular Biology</i> , 2006, 357, 139-150. | 4.2 | 54 |
| 15 | Resonance Assignments of 30 kDa Complexes of TFIID Subunit TAF1 with TATA-binding Protein. <i>Journal of Biomolecular NMR</i> , 2005, 33, 76-76. | 2.8 | 2 |
| 16 | Structural Basis for the Activation of Microtubule Assembly by the EB1 and p150Glued Complex. <i>Molecular Cell</i> , 2005, 19, 449-460. | 9.7 | 121 |
| 17 | Cold-shock induced high-yield protein production in <i>Escherichia coli</i> . <i>Nature Biotechnology</i> , 2004, 22, 877-882. | 17.5 | 307 |
| 18 | Structural and Functional Characterization on the Interaction of Yeast TFIID Subunit TAF1 with TATA-binding Protein. <i>Journal of Molecular Biology</i> , 2004, 339, 681-693. | 4.2 | 37 |

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 19 | Probing Zn ²⁺ -binding effects on the zinc-ribbon domain of human general transcription factor TFIIIB. <i>Biochemical Journal</i> , 2004, 378, 317-324. | 3.7 | 16 |
| 20 | Nuclear Magnetic Resonance-Based Dissection of a Glycosyltransferase Specificity for the Mucin MUC1 Tandem Repeat. <i>Biochemistry</i> , 2003, 42, 13817-13825. | 2.5 | 17 |
| 21 | Structural Basis for Simultaneous Binding of Two Carboxy-terminal Peptides of Plant Glutamate Decarboxylase to Calmodulin. <i>Journal of Molecular Biology</i> , 2003, 328, 193-204. | 4.2 | 100 |
| 22 | Photo-Induced Peptide Cleavage in the Green-to-Red Conversion of a Fluorescent Protein. <i>Molecular Cell</i> , 2003, 12, 1051-1058. | 9.7 | 276 |
| 23 | Protein Structure Calculation from NMR 267. , 2002, 173, 267-283. | | 2 |
| 24 | The ATCUN Domain as a Probe of Intermolecular Interactions: Application to Calmodulin~Peptide Complexes. <i>Journal of the American Chemical Society</i> , 2002, 124, 14002-14003. | 13.7 | 36 |
| 25 | Characterization of the ATP-Binding Domain of the Sarco(endo)plasmic Reticulum Ca ²⁺ -ATPase: Probing Nucleotide Binding by Multidimensional NMR. <i>Biochemistry</i> , 2002, 41, 1156-1164. | 2.5 | 32 |
| 26 | Detecting Protein Kinase Recognition Modes of Calmodulin by Residual Dipolar Couplings in Solution NMR. <i>Biochemistry</i> , 2002, 41, 12899-12906. | 2.5 | 32 |
| 27 | Structure of the inositol 1,4,5-trisphosphate receptor binding core in complex with its ligand. <i>Nature</i> , 2002, 420, 696-700. | 27.8 | 309 |
| 28 | FRET-based in vivo Ca ²⁺ imaging by a new calmodulin-GFP fusion molecule. <i>Nature Structural Biology</i> , 2001, 8, 1069-1073. | 9.7 | 196 |
| 29 | Calexitin B Is a New Member of the Sarcoplasmic Calcium-binding Protein Family. <i>Journal of Biological Chemistry</i> , 2001, 276, 22529-22536. | 3.4 | 17 |
| 30 | TFIIA-TAF regulatory interplay: NMR evidence for overlapping binding sites on TBP. <i>FEBS Letters</i> , 2000, 468, 149-154. | 2.8 | 34 |
| 31 | Some NMR experiments and a structure determination employing a [15N,2H] enriched protein. <i>Journal of Biomolecular NMR</i> , 1998, 12, 259-276. | 2.8 | 34 |
| 32 | The Influence of a Scalar-Coupled Deuterium upon the Relaxation of a ¹⁵ N Nucleus and Its Possible Exploitation as a Probe for Side-Chain Interactions in Proteins. <i>Journal of Magnetic Resonance</i> , 1997, 124, 61-71. | 2.1 | 17 |
| 33 | Synthesis, and crystal and molecular structure of the 310-helical α -dehydro pentapeptide Boc-Leu-Phe-Ala- α -Phe-Leu-OMe. <i>Biopolymers</i> , 1995, 35, 141-147. | 2.4 | 36 |
| 34 | Assignment of Disulfide Bonds in the X Protein (HBx) of Hepatitis B Virus. <i>Biochemical and Biophysical Research Communications</i> , 1995, 212, 919-924. | 2.1 | 27 |
| 35 | Synthesis, Crystal and Molecular Structure of Boc-Pro- α -Phe-Ala- β -Phe- Ala-OMe; A Pentapeptide with a Novel β -Bend Ribbon Structure. <i>Angewandte Chemie International Edition in English</i> , 1994, 33, 970-973. | 4.4 | 29 |