

Michikazu Tanio

List of Publications by Year in descending order

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34
papers

915
citations

516215

16
h-index

454577

30
g-index

34
all docs

34
docs citations

34
times ranked

616
citing authors

#	ARTICLE	IF	CITATIONS
1	Existence of a Proton Transfer Chain in Bacteriorhodopsin: A Participation of Glu-194 in the Release of Protons to the Extracellular Surface. <i>Biochemistry</i> , 1998, 37, 2496-2506.	1.2	173
2	Location of a Cation-Binding Site in the Loop between Helices F and G of Bacteriorhodopsin as Studied by ¹³ C NMR. <i>Biophysical Journal</i> , 1999, 76, 1523-1531.	0.2	72
3	Direct Evidence of Interaction of a Green Tea Polyphenol, Epigallocatechin Gallate, with Lipid Bilayers by Solid-state Nuclear Magnetic Resonance. <i>Bioscience, Biotechnology and Biochemistry</i> , 2004, 68, 1743-1747.	0.6	67
4	Conformation and backbone dynamics of bacteriorhodopsin revealed by ¹³ C-NMR. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2000, 1460, 39-48.	0.5	64
5	Trivalent Recognition Unit of Innate Immunity System. <i>Journal of Biological Chemistry</i> , 2007, 282, 3889-3895.	1.6	60
6	A structure-based mechanism for benzalacetone synthase from <i>Rheum palmatum</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 669-673.	3.3	48
7	Dynamic aspects of membrane proteins and membrane-associated peptides as revealed by ¹³ C NMR: Lessons from bacteriorhodopsin as an intact protein. <i>Annual Reports on NMR Spectroscopy</i> , 2002, 47, 39-108.	0.7	38
8	Irreversible Conformational Change of Bacterio-opsin Induced by Binding of Retinal during Its Reconstitution to Bacteriorhodopsin, as Studied by ¹³ NMR. <i>Journal of Biochemistry</i> , 2000, 127, 861-869.	0.9	36
9	Expression, purification and crystallization of a human tau-tubulin kinase 2 that phosphorylates tau protein. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007, 63, 602-604.	0.7	35
10	Long-Distance Effects of Site-Directed Mutations on Backbone Conformation in Bacteriorhodopsin from Solid State NMR of [¹³ C]Val-Labeled Proteins. <i>Biophysical Journal</i> , 1999, 77, 431-442.	0.2	34
11	Alteration of Conformation and Dynamics of Bacteriorhodopsin Induced by Protonation of Asp 85 and Deprotonation of Schiff Base as Studied by ¹³ C NMR. <i>Biochemistry</i> , 2000, 39, 14472-14480.	1.2	34
12	Conformational Changes of Bacteriorhodopsin along the Proton-Conduction Chain as Studied with ¹³ C NMR of [¹³ C]Ala-Labeled Protein: Arg82 May Function as an Information Mediator. <i>Biophysical Journal</i> , 1999, 77, 1577-1584.	0.2	31
13	Binding site of C-reactive protein on M-ficolin. <i>Molecular Immunology</i> , 2009, 47, 215-221.	1.0	31
14	Site-directed ¹³ C solid-state NMR studies on membrane proteins: strategy and goals toward revealing conformation and dynamics as illustrated for bacteriorhodopsin labeled with [¹³ C]amino acid residues. <i>Magnetic Resonance in Chemistry</i> , 2004, 42, 218-230.	1.1	28
15	Evidence of local conformational fluctuations and changes in bacteriorhodopsin, dependent on lipids, detergents and trimeric structure, as studied by ¹³ C NMR. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 1998, 1375, 84-92.	1.4	26
16	Histidine-regulated activity of M-ficolin. <i>Biochemical Journal</i> , 2009, 417, 485-491.	1.7	17
17	Significance of low-frequency local fluctuation motions in the transmembrane B and C α -helices of bacteriorhodopsin, to facilitate efficient proton uptake from the cytoplasmic surface, as revealed by site-directed solid-state ¹³ C NMR. <i>European Biophysics Journal</i> , 2004, 33, 580-588.	1.2	15
18	Surface and Dynamic Structures of Bacteriorhodopsin in a 2D Crystal, a Distorted or Disrupted Lattice, as Revealed by Site-directed Solid-state ¹³ C NMR. <i>Photochemistry and Photobiology</i> , 2007, 83, 253-262.	1.3	15

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19	Conformation and dynamics of membrane proteins and biologically active peptides as studied by high-resolution solid-state ¹³ C NMR. <i>Journal of Molecular Structure</i> , 1998, 441, 137-148.	1.8	12
20	Amino acid-selective isotope labeling of proteins for nuclear magnetic resonance study: Proteins secreted by <i>Brevibacillus choshinensis</i> . <i>Analytical Biochemistry</i> , 2009, 386, 156-160.	1.1	12
21	Trimeric structure and conformational equilibrium of M-ficolin fibrinogen-like domain. <i>Journal of Synchrotron Radiation</i> , 2008, 15, 243-245.	1.0	11
22	Overexpression, purification and preliminary crystallographic analysis of human M-ficolin fibrinogen-like domain. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006, 62, 652-655.	0.7	9
23	¹⁵ N isotope labeling of a protein secreted by <i>Brevibacillus choshinensis</i> for NMR study. <i>Analytical Biochemistry</i> , 2008, 373, 164-166.	1.1	9
24	Intramolecular allosteric interaction in the phospholipase C- $\hat{1}$ pleckstrin homology domain. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2013, 1834, 1034-1043.	1.1	9
25	Dynamic aspect of bacteriorhodopsin as viewed from ¹³ C NMR: Conformational elucidation, surface dynamics and information transfer from the surface to inner residues. <i>Spectroscopy</i> , 2002, 16, 107-120.	0.8	8
26	Analysis of the phospholipase C- $\hat{1}$ pleckstrin homology domain using native polyacrylamide gel electrophoresis. <i>Analytical Biochemistry</i> , 2012, 431, 106-114.	1.1	7
27	Crystallization and preliminary crystallographic analysis of a plant type III polyketide synthase that produces benzalacetone. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2008, 64, 304-306.	0.7	6
28	Suppressed or recovered intensities analysis in site-directed ¹³ C NMR: Assessment of low-frequency fluctuations in bacteriorhodopsin and D85N mutants revisited. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2010, 1798, 167-176.	1.4	5
29	Validation of HPLC Method for Determination of Histamine in Human Immunoglobulin Formulations. <i>Journal of AOAC INTERNATIONAL</i> , 2020, 103, 1223-1229.	0.7	2
30	¹ H, ¹³ C and ¹⁵ N backbone resonance assignments of the monomeric human M-ficolin fibrinogen-like domain secreted by <i>Brevibacillus choshinensis</i> . <i>Biomolecular NMR Assignments</i> , 2014, 8, 207-211.	0.4	1
31	Structure and Dynamics of Membrane-Bound Proteins. , 2018, , 669-681.		0
32	Structure and Dynamics of Membrane-Bound Proteins. , 2016, , 1-13.		0
33	Functional and structural characterization of membrane-binding proteins using NMR. <i>Annual Reports on NMR Spectroscopy</i> , 2022, , 47-131.	0.7	0
34	Calcium-dependent reversible coaggregation activity of C-reactive protein and M-ficolin. <i>Molecular Immunology</i> , 2022, 149, 157-164.	1.0	0