## Toshimichi Fujiwara

List of Publications by Year in descending order

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102 papers 3,081 citations

218677 26 h-index 50 g-index

107 all docs

107 docs citations

107 times ranked

3989 citing authors

#	Article	IF	CITATIONS
1	Protein Data Bank: the single global archive for 3D macromolecular structure data. Nucleic Acids Research, 2019, 47, D520-D528.	14.5	671
2	3D structure of amyloid protofilaments of beta2-microglobulin fragment probed by solid-state NMR. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 18119-18124.	7.1	224
3	Structure of the light-harvesting bacteriochlorophyllcassembly in chlorosomes fromChlorobium limicoladetermined by solid-state NMR. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 790-795.	7.1	103
4	Photoinduced Changes of Adsorbed Water on a TiO2Photocatalytic Film As Studied by1H NMR Spectroscopy. Journal of Physical Chemistry B, 2003, 107, 12042-12044.	2.6	101
5	Characteristics of Water Adsorbed on TiO2Photocatalytic Systems with Increasing Temperature as Studied by Solid-State1H NMR Spectroscopy. Journal of Physical Chemistry B, 2004, 108, 9121-9125.	2.6	94
6	Current NMR Techniques for Structure-Based Drug Discovery. Molecules, 2018, 23, 148.	3.8	92
7	New tools and functions in dataâ€out activities at Protein Data Bank Japan (PDBj). Protein Science, 2018, 27, 95-102.	7.6	90
8	Structure of Tightly Membrane-Bound Mastoparan-X, a G-Protein-Activating Peptide, Determined by Solid-State NMR. Biophysical Journal, 2006, 91, 1368-1379.	0.5	74
9	Continuously Frequency Tunable High Power Sub-THz Radiation Source—Gyrotron FU CW VI for 600ÂMHz DNP-NMR Spectroscopy. Journal of Infrared, Millimeter, and Terahertz Waves, 2010, 31, 775-790.	2.2	72
10	Helium-cooling and -spinning dynamic nuclear polarization for sensitivity-enhanced solid-state NMR at 14T and 30K. Journal of Magnetic Resonance, 2012, 225, 1-9.	2.1	72
11	Dynamic nuclear polarization experiments at 14.1 T for solid-state NMR. Physical Chemistry Chemical Physics, 2010, 12, 5799.	2.8	68
12	Advanced instrumentation for DNP-enhanced MAS NMR for higher magnetic fields and lower temperatures. Journal of Magnetic Resonance, 2016, 264, 107-115.	2.1	64
13	Modern Technologies of Solution Nuclear Magnetic Resonance Spectroscopy for Three-dimensional Structure Determination of Proteins Open Avenues for Life Scientists. Computational and Structural Biotechnology Journal, 2017, 15, 328-339.	4.1	57
14	13C-13C and 13C-15N Dipolar Correlation NMR of Uniformly Labeled Organic Solids for the Complete Assignment of Their 13C and 15N Signals: An Application to Adenosine. Journal of the American Chemical Society, 1995, 117, 11351-11352.	13.7	47
15	How Far Can the Sensitivity of NMR Be Increased?. Annual Reports on NMR Spectroscopy, 2006, 58, 155-175.	1.5	44
16	Photocatalytic Reaction Sites at the TiO2Surface as Studied by Solid-State1H NMR Spectroscopy. Langmuir, 2003, 19, 1935-1937.	3.5	39
17	Signal assignments and chemical-shift structural analysis of uniformly13C,15N-labeled peptide, mastoparan-X, by multidimensional solid-state NMR under magic-angle spinning. Journal of Biomolecular NMR, 2004, 28, 311-325.	2.8	38
18	Application of Continuously Frequency-Tunable 0.4 THz Gyrotron to Dynamic Nuclear Polarization for 600ÂMHz Solid-State NMR. Journal of Infrared, Millimeter, and Terahertz Waves, 2012, 33, 745-755.	2.2	38

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19	Closed-cycle cold helium magic-angle spinning for sensitivity-enhanced multi-dimensional solid-state NMR. Journal of Magnetic Resonance, 2015, 259, 76-81.	2.1	38
20	Redox-coupled Conformational Alternations in Cytochrome c3 from D.vulgaris Miyazaki F on the Basis of its Reduced Solution Structure. Journal of Molecular Biology, 2002, 319, 767-778.	4.2	36
21	Multidimensional Solid-State Nuclear Magnetic Resonance for Correlating Anisotropic Interactions under Magic-Angle Spinning Conditions. Journal of Magnetic Resonance, 1997, 124, 147-153.	2.1	34
22	Molecular mobility of protein in lyophilized formulations linked to the molecular mobility of polymer excipients, as determined by high resolution 13C solid-state NMR. Pharmaceutical Research, 1999, 16, 1621-1625.	3.5	34
23	Diverse Structural Conversion and Aggregation Pathways of AlzheimerÊ⅓s Amyloid-β (1–40). ACS Nano, 2019, 13, 8766-8783.	14.6	33
24	Boosting Protein Dynamics Studies Using Quantitative Nonuniform Sampling NMR Spectroscopy. Journal of Physical Chemistry B, 2011, 115, 13740-13745.	2.6	30
25	The wobblingâ€inâ€aâ€cone analysis of internal motion in macromolecules. Journal of Chemical Physics, 1985, 83, 3110-3117.	3.0	29
26	Phase-modulated heteronuclear decoupling in NMR of solids. Journal of Magnetic Resonance, 2003, 162, 46-53.	2.1	27
27	Signal assignment and secondary structure analysis of a uniformly [13C, 15N]-labeled membrane protein, H+-ATP synthase subunit c, by magic-angle spinning solid-state NMR. Journal of Biomolecular NMR, 2006, 36, 279-293.	2.8	26
28	An automated system designed for large scale NMR data deposition and annotation: application to over 600 assigned chemical shift data entries to the BioMagResBank from the Riken Structural Genomics/Proteomics Initiative internal database. Journal of Biomolecular NMR, 2012, 53, 311-320.	2.8	26
29	Structure determination of uniformly 13C, 15N labeled protein using qualitative distance restraints from MAS solid-state 13C-NMR observed paramagnetic relaxation enhancement. Journal of Biomolecular NMR, 2016, 64, 87-101.	2.8	25
30	Development of second harmonic gyrotrons, Gyrotron FU CW GII and Gyrotron FU CW GIII, equipped with internal mode converters. Journal of Infrared, Millimeter, and Terahertz Waves, 2014, 35, 169-178.	2.2	24
31	Latest approaches for efficient protein production in drug discovery. Expert Opinion on Drug Discovery, 2014, 9, 1189-1204.	5.0	24
32	Robust folding of a de novo designed ideal protein even with most of the core mutated to valine. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 31149-31156.	7.1	23
33	Noise peak filtering in multi-dimensional NMR spectra using convolutional neural networks. Bioinformatics, 2018, 34, 4300-4301.	4.1	22
34	Photoinduced Changes of Surface and Adsorbed Water in TiO2Photocatalytic Systems as Studied by Solid State1H-NMR Spectroscopy. Chemistry Letters, 2002, 31, 420-421.	1.3	21
35	Phosphoinositide binding by the PH domain in ceramide transfer protein (CERT) is inhibited by hyperphosphorylation of an adjacent serine-repeat motif. Journal of Biological Chemistry, 2018, 293, 11206-11217.	3.4	21
36	Cooperative regulation of PBI1 and MAPKs controls WRKY45 transcription factor in rice immunity. Nature Communications, 2022, 13, 2397.	12.8	20

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37	Development of a Compact sub-THz Gyrotron FU CW CI for Application to High Power THz Technologies. Journal of Infrared, Millimeter, and Terahertz Waves, 2012, 33, 724-744.	2.2	19
38	Characteristics of the mode converter of Gyrotron FU CW GII radiating Gaussian beams in both the fundamental and second harmonic frequency bands. Journal of Infrared, Millimeter, and Terahertz Waves, 2014, 35, 517-524.	2.2	19
39	Oxytocin solution structure changes upon protonation of the N-terminus in dimethyl sulfoxide. Journal of Biomolecular NMR, 1993, 3, 653-73.	2.8	18
40	Band-selective recoupling of homonuclear double-quantum dipolar interaction with a generalized composite $0\hat{A}^{\circ}$ pulse: application to aliphatic region-selective magnetization transfer in solids. Journal of Magnetic Resonance, 2003, 162, 54-66.	2.1	18
41	Detection of Peptideâ^'Phospholipid Interaction Sites in Bilayer Membranes by13C NMR Spectroscopy:Â Observation of2H/31P-Selective1H-Depolarization under Magic-Angle Spinning. Journal of the American Chemical Society, 2006, 128, 10654-10655.	13.7	18
42	Broadband Continuously Frequency Tunable Gyrotron for 600 MHz DNP-NMR Spectroscopy. Plasma and Fusion Research, 2014, 9, 1206058-1206058.	0.7	18
43	Utilization of lysine 13C-methylation NMR for protein–protein interaction studies. Journal of Biomolecular NMR, 2013, 55, 19-31.	2.8	17
44	Fluid Mechanical Matching of H+-ATP Synthase Subunit c-Ring with Lipid Membranes Revealed by 2H Solid-State NMR. Biophysical Journal, 2008, 94, 4339-4347.	0.5	16
45	Utilization of paramagnetic relaxation enhancements for high-resolution NMR structure determination of a soluble loop-rich protein with sparse NOE distance restraints. Journal of Biomolecular NMR, 2015, 61, 55-64.	2.8	16
46	Temperature- and composition-dependent conformational transitions of amphipathic peptide–phospholipid nanodiscs. Journal of Colloid and Interface Science, 2021, 588, 522-530.	9.4	16
47	Conformational study of 13C-enriched fibroin in the solid state, using the cross polarization nuclear magnetic resonance method. Journal of Molecular Biology, 1986, 187, 137-140.	4.2	14
48	Precision1Hâ€"1H distance measurement via13C NMR signals: utilization of1Hâ€"1H double-quantum dipolar interactions recoupled under magic angle spinning conditions. Magnetic Resonance in Chemistry, 2004, 42, 291-300.	1.9	14
49	Membrane Mediated Antimicrobial and Antitumor Activity of Cathelicidin 6: Structural Insights from Molecular Dynamics Simulation on Multi-Microsecond Scale. PLoS ONE, 2016, 11, e0158702.	2.5	14
50	Solution NMR structure and inhibitory effect against amyloid-β fibrillation of Humanin containing a d-isomerized serine residue. Biochemical and Biophysical Research Communications, 2016, 477, 647-653.	2.1	14
51	Protein 19F-labeling using transglutaminase for the NMR study of intermolecular interactions. Journal of Biomolecular NMR, 2017, 68, 271-279.	2.8	14
52	The Route from the Folded to the Amyloid State: Exploring the Potential Energy Surface of a Drugâ€Like Miniprotein. Chemistry - A European Journal, 2020, 26, 1968-1978.	3.3	14
53	Polyphenolâ€solubility alters amyloid fibril formation of αâ€synuclein. Protein Science, 2021, 30, 1701-1713.	7.6	14
54	Compound Radiofrequency-Driven Recoupling Pulse Sequences for Efficient Magnetization Transfer by Homonuclear Dipolar Interaction under Magic-Angle Spinning Conditions. Journal of Magnetic Resonance, 2000, 145, 73-83.	2.1	13

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55	Multidimensional solid-state nuclear magnetic resonance for determining the dihedral angle from the correlation of 13C–1H and 13C–13C dipolar interactions under magic-angle spinning conditions. Journal of Chemical Physics, 1998, 109, 2380-2393.	3.0	12
56	A novel chitinâ€binding mode of the chitinâ€binding domain of chitinase A1 from <i>Bacillus circulans </i> <scp>WL</scp> â€12 revealed by solidâ€state <scp>NMR</scp> . FEBS Letters, 2018, 592, 3173-3182.	2.8	11
57	Gadolinium Complexes as Contrast Agent for Cellular NMR Spectroscopy. International Journal of Molecular Sciences, 2020, 21, 4042.	4.1	11
58	The interactions of ferric and ferrous cytochrome c with cardiolipin in phospholipid membranes studied by solid-state 2H and 31P NMR. Journal of Molecular Structure, 1998, 441, 183-188.	3.6	10
59	Atomic structure of the bacteriochlorophyll c assembly in intact chlorosomes from Chlorobium limicola determined by solid-state NMR. Photosynthesis Research, 2010, 104, 221-231.	2.9	10
60	Purification, characterization and reconstitution into membranes of the oligomeric c-subunit ring of thermophilic FoF1-ATP synthase expressed in Escherichia coli. Protein Expression and Purification, 2012, 82, 396-401.	1.3	10
61	Active-Site Structure of the Thermophilic Foc-Subunit Ring in Membranes Elucidated by Solid-State NMR. Biophysical Journal, 2014, 106, 390-398.	0.5	10
62	Mechanistic and structural basis of bioengineered bovine Cathelicidin-5 with optimized therapeutic activity. Scientific Reports, 2017, 7, 44781.	3.3	10
63	Evolution of $\hat{I}\pm$ -synuclein conformation ensemble toward amyloid fibril via liquid-liquid phase separation (LLPS) as investigated by dynamic nuclear polarization-enhanced solid-state MAS NMR. Neurochemistry International, 2022, 157, 105345.	3.8	10
64	Modulation of the specific interaction of cardiolipin with cytochrome c by zwitterionic phospholipids in binary mixed bilayers; a 2H and 31P NMR study. Journal of Molecular Structure, 1995, 355, 47-53.	3.6	9
65	Structural analysis of pituitary adenylate cyclase-activating polypeptides bound to phospholipid membranes by magic angle spinning solid-state NMR. Biochimica Et Biophysica Acta - Biomembranes, 2007, 1768, 3001-3011.	2.6	9
66	Further Characterization of 394-GHz Gyrotron FU CW GII with Additional PID Control System for 600-MHz DNP-SSNMR Spectroscopy. Journal of Infrared, Millimeter, and Terahertz Waves, 2016, 37, 825-836.	2.2	9
67	Publication of nuclear magnetic resonance experimental data with semantic web technology and the application thereof to biomedical research of proteins. Journal of Biomedical Semantics, 2016, 7, 16.	1.6	9
68	Absolute 1H polarization measurement with a spin-correlated component of magnetization by hyperpolarized MAS-DNP solid-state NMR. Solid State Nuclear Magnetic Resonance, 2019, 99, 20-26.	2.3	9
69	In-cell NMR as a sensitive tool to monitor physiological condition of Escherichia coli. Scientific Reports, 2020, 10, 2466.	3.3	9
70	Efficiency analysis of helium-cooled MAS DNP: case studies of surface-modified nanoparticles and homogeneous small-molecule solutions. Physical Chemistry Chemical Physics, 2021, 23, 4919-4926.	2.8	9
71	Structure analysis of membrane-reconstituted subunit c-ring of E. coli H+-ATP synthase by solid-state NMR. Journal of Biomolecular NMR, 2010, 48, 1-11.	2.8	8
72	1 H-detected 1 $\mathrm{H\hat{a}}$ 1 H correlation spectroscopy of a stereo-array isotope labeled amino acid under fast magic-angle spinning. Journal of Magnetic Resonance, 2010, 203, 253-256.	2.1	8

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73	Combined Use of Replica-Exchange Molecular Dynamics and Magic-Angle-Spinning Solid-State NMR Spectral Simulations for Determining the Structure and Orientation of Membrane-Bound Peptide. Journal of Physical Chemistry B, 2011, 115, 9327-9336.	2.6	8
74	Secondary structural analysis of proteins based on 13C chemical shift assignments in unresolved solid-state NMR spectra enhanced by fragmented structure database. Journal of Biomolecular NMR, 2013, 55, 189-200.	2.8	8
75	Spectral fitting for signal assignment and structural analysis of uniformly 13C-labeled solid proteins by simulated annealing based on chemical shifts and spin dynamics. Journal of Biomolecular NMR, 2007, 38, 325-339.	2.8	7
76	A magic-angle-spinning NMR method for H1–H1 distance measurement using coherent polarization transfer in C13-labeled organic solids. Journal of Chemical Physics, 2008, 129, 154504.	3.0	7
77	Direct assignment of 13C solid-state NMR signals of TFoF1 ATP synthase subunit c-ring in lipid membranes and its implication for the ring structure. Journal of Biomolecular NMR, 2018, 70, 53-65.	2.8	7
78	Structural analysis of cross-linked poly(vinyl alcohol) using high-field DNP-NMR. RSC Advances, 2020, 10, 8039-8043.	3.6	7
79	Cryogenic signal amplification combined with helium-temperature MAS DNP toward ultimate NMR sensitivity at high field conditions. Journal of Magnetic Resonance, 2022, 335, 107139.	2.1	7
80	In support of the BMRB. Nature Structural and Molecular Biology, 2012, 19, 854-860.	8.2	6
81	1H, 15N and 13C resonance assignments of the conserved region in the middle domain of S. pombe Sin1 protein. Biomolecular NMR Assignments, 2015, 9, 89-92.	0.8	6
82	Plant-specific DUF1110 protein from <i>Oryza sativa </i> : expression, purification and crystallization. Acta Crystallographica Section F, Structural Biology Communications, 2016, 72, 480-484.	0.8	6
83	Transient antibody-antigen interactions mediate the strain-specific recognition of a conserved malaria epitope. Communications Biology, 2018, 1, 58.	4.4	6
84	Peptide Cyclization Mediated by Metalâ€Free Sâ€Arylation: Sâ€Protected Cysteine Sulfoxide as an Umpolung of the Cysteine Nucleophile. Chemistry - A European Journal, 2021, 27, 14092-14099.	3.3	6
85	Nanodisc-to-Nanofiber Transition of Noncovalent Peptide–Phospholipid Assemblies. ACS Omega, 2017, 2, 2935-2944.	3.5	5
86	Conformational states of HAMP domains interacting with sensory rhodopsin membrane systems: an integrated all-atom and coarse-grained molecular dynamics simulation approach. Molecular BioSystems, 2017, 13, 193-207.	2.9	5
87	Surface-Only Spectroscopy for Diffusion-Limited Systems Using Ultra-Low-Temperature DNP MAS NMR at 16.4 T. Journal of Physical Chemistry C, 2020, 124, 18609-18614.	3.1	5
88	A hybrid strategy combining solution NMR spectroscopy and isothermal titration calorimetry to characterize protein-nanodisc interaction. Analytical Biochemistry, 2022, 639, 114521.	2.4	5
89	Behavior of water molecules associated with the phase transitions in the binary system of dioctadecyldimethylammonium chloride and water studied by proton and deuterium magnetic resonances. Journal of Colloid and Interface Science, 1989, 127, 26-34.	9.4	4
90	Designing Analogues of Mini Atrial Natriuretic Peptide Based on Structural Analysis by NMR and Restrained Molecular Dynamics. Journal of Medicinal Chemistry, 2002, 45, 881-887.	6.4	4

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91	Phospholipid-Dependent Regulation of Cytochrome c3-Mediated Electron Transport across Membranes. Biophysical Journal, 2006, 90, 506-513.	0.5	4
92	Sensitivity enhancement by sequential data acquisition for 13C-direct detection NMR. Journal of Magnetic Resonance, 2021, 322, 106878.	2.1	4
93	Biochemical propensity mapping for structural and functional anatomy of importin α IBB domain. Genes To Cells, 2022, 27, 173-191.	1.2	4
94	Cold-Shock Expression System in E. coli for Protein NMR Studies. Methods in Molecular Biology, 2017, 1586, 345-357.	0.9	3
95	Advances in High-Field DNP Methods. , 2018, , 91-134.		3
96	Sequence requirements of the FFATâ€like motif for specific binding to VAPâ€A are revealed by NMR. FEBS Letters, 2021, 595, 2248-2256.	2.8	3
97	Veratridine binding to a transmembrane helix of sodium channel Nav1.4 determined by solid-state NMR. Bioorganic and Medicinal Chemistry, 2018, 26, 5644-5653.	3.0	2
98	Structural dynamics of the chromo-shadow domain and chromodomain of HP1 bound to histone H3K9 methylated peptide, as measured by site-directed spin-labeling EPR spectroscopy. Biochemical and Biophysical Research Communications, 2021, 567, 42-48.	2.1	2
99	Osmotic Stability of Muramyl Dipeptide-Bearing Liposomes and Molecular Miscibility in Their Membranes. Bulletin of the Chemical Society of Japan, 1999, 72, 541-548.	3.2	1
100	Corrugated transmission line systems for 395 GHz/600 MHz and 460 GHz/700 MHz DNP-NMR spectroscopy. , 2014, , .		1
101	Microdomain formation in phosphatidylethanolamine bilayers detected by 2H-NMR. Chemistry and Physics of Lipids, 1995, 76, 55-62.	3.2	0
102	3P-086 Solid-state NMR measurement of H^+-ATP synthase subunit c-ring reconstituted into lipid bilayers(The 46th Annual Meeting of the Biophysical Society of Japan). Seibutsu Butsuri, 2008, 48, S140.	0.1	O