

Mei Hong

List of Publications by Year in descending order

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203
papers

13,404
citations

17776

65
h-index

35168

102
g-index

209
all docs

209
docs citations

209
times ranked

11073
citing authors

#	ARTICLE	IF	CITATIONS
1	From Angstroms to Nanometers: Measuring Interatomic Distances by Solid-State NMR. <i>Chemical Reviews</i> , 2022, 122, 9848-9879.	23.0	29
2	Cholesterol-Mediated Clustering of the HIV Fusion Protein gp41 in Lipid Bilayers. <i>Journal of Molecular Biology</i> , 2022, 434, 167345.	2.0	4
3	Binding Sites of a Positron Emission Tomography Imaging Agent in Alzheimer's β -Amyloid Fibrils Studied Using ^{19}F Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2022, 144, 1416-1430.	6.6	22
4	Age-dependent aggregation of β -synuclein in the nervous system of gut-brain axis is associated with caspase-1 activation. <i>Metabolic Brain Disease</i> , 2022, 37, 1669-1681.	1.4	2
5	High-pH structure of EmrE reveals the mechanism of proton-coupled substrate transport. <i>Nature Communications</i> , 2022, 13, 991.	5.8	15
6	pH- and Calcium-Dependent Aromatic Network in the SARS-CoV-2 Envelope Protein. <i>Journal of the American Chemical Society</i> , 2022, 144, 6839-6850.	6.6	21
7	Clustering of tetrameric influenza M2 peptides in lipid bilayers investigated by ^{19}F solid-state NMR. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2022, 1864, 183909.	1.4	8
8	Fluent molecular mixing of Tau isoforms in Alzheimer's disease neurofibrillary tangles. <i>Nature Communications</i> , 2022, 13, .	5.8	27
9	Golgi-localized putative S-adenosyl methionine transporters required for plant cell wall polysaccharide methylation. <i>Nature Plants</i> , 2022, 8, 656-669.	4.7	23
10	Solid-state NMR spectroscopy. <i>Nature Reviews Methods Primers</i> , 2021, 1, .	11.8	196
11	Direct Observation of Cholesterol Dimers and Tetramers in Lipid Bilayers. <i>Journal of Physical Chemistry B</i> , 2021, 125, 1825-1837.	1.2	25
12	Water orientation and dynamics in the closed and open influenza B virus M2 proton channels. <i>Communications Biology</i> , 2021, 4, 338.	2.0	20
13	Comparative analysis of ^{13}C chemical shifts of β -sheet amyloid proteins and outer membrane proteins. <i>Journal of Biomolecular NMR</i> , 2021, 75, 151-166.	1.6	2
14	Inclusion of the C-Terminal Domain in the β -Sheet Core of Heparin-Fibrillized Three-Repeat Tau Protein Revealed by Solid-State Nuclear Magnetic Resonance Spectroscopy. <i>Journal of the American Chemical Society</i> , 2021, 143, 7839-7851.	6.6	30
15	Structurally Based Design of Glucagon Mutants That Inhibit Fibril Formation. <i>Biochemistry</i> , 2021, 60, 2033-2043.	1.2	4
16	Xylan Structure and Dynamics in Native <i>Brachypodium</i> Grass Cell Walls Investigated by Solid-State NMR Spectroscopy. <i>ACS Omega</i> , 2021, 6, 15460-15471.	1.6	19
17	Off-resonance ^{13}C - ^2H REDOR NMR for site-resolved studies of molecular motion. <i>Journal of Biomolecular NMR</i> , 2021, 75, 335-345.	1.6	3
18	Interactions of HIV gp41's membrane-proximal external region and transmembrane domain with phospholipid membranes from ^{31}P NMR. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2021, 1863, 183723.	1.4	7

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19	Structure and dynamics of the drug-bound bacterial transporter EmrE in lipid bilayers. <i>Nature Communications</i> , 2021, 12, 172.	5.8	40
20	Bacterial Phosphate Granules Contain Cyclic Polyphosphates: Evidence from ³¹ P Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2020, 142, 18407-18421.	6.6	28
21	Structure and drug binding of the SARS-CoV-2 envelope protein transmembrane domain in lipid bilayers. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 1202-1208.	3.6	294
22	Pulsed Third-Spin-Assisted Recoupling NMR for Obtaining Long-Range ¹³ C- ¹³ C and ¹⁵ N- ¹³ C Distance Restraints. <i>Journal of Physical Chemistry B</i> , 2020, 124, 7138-7151.	1.2	11
23	Hydration and Dynamics of Full-Length Tau Amyloid Fibrils Investigated by Solid-State Nuclear Magnetic Resonance. <i>Biochemistry</i> , 2020, 59, 2237-2248.	1.2	30
24	Cholesterol Interaction with the Trimeric HIV Fusion Protein gp41 in Lipid Bilayers Investigated by Solid-State NMR Spectroscopy and Molecular Dynamics Simulations. <i>Journal of Molecular Biology</i> , 2020, 432, 4705-4721.	2.0	21
25	Two-dimensional ¹⁹ F- ¹³ C correlation NMR for ¹⁹ F resonance assignment of fluorinated proteins. <i>Journal of Biomolecular NMR</i> , 2020, 74, 193-204.	1.6	21
26	Atomic structures of closed and open influenza B M2 proton channel reveal the conduction mechanism. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 160-167.	3.6	52
27	In vitro ON4R tau fibrils contain a monomorphic β -sheet core enclosed by dynamically heterogeneous fuzzy coat segments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 16357-16366.	3.3	76
28	Fast MAS ¹ H- ¹³ C correlation NMR for structural investigations of plant cell walls. <i>Journal of Biomolecular NMR</i> , 2019, 73, 661-674.	1.6	19
29	Fully hydrophobic HIV gp41 adopts a hemifusion-like conformation in phospholipid bilayers. <i>Journal of Biological Chemistry</i> , 2019, 294, 14732-14744.	1.6	11
30	The peptide hormone glucagon forms amyloid fibrils with two coexisting β -strand conformations. <i>Nature Structural and Molecular Biology</i> , 2019, 26, 592-598.	3.6	58
31	Elucidating Relayed Proton Transfer through a His-Trp-His Triad of a Transmembrane Proton Channel by Solid-State NMR. <i>Journal of Molecular Biology</i> , 2019, 431, 2554-2566.	2.0	16
32	High-Sensitivity Detection of Nanometer ¹ H- ¹⁹ F Distances for Protein Structure Determination by ¹ H-Detected Fast MAS NMR. <i>Journal of Physical Chemistry B</i> , 2019, 123, 4387-4391.	1.2	30
33	High-sensitivity protein solid-state NMR spectroscopy. <i>Current Opinion in Structural Biology</i> , 2019, 58, 183-190.	2.6	23
34	Elucidating ligand-bound structures of membrane proteins using solid-state NMR spectroscopy. <i>Current Opinion in Structural Biology</i> , 2019, 57, 103-109.	2.6	17
35	The Transmembrane Conformation of the Influenza B Virus M2 Protein in Lipid Bilayers. <i>Scientific Reports</i> , 2019, 9, 3725.	1.6	16
36	The molecular structure of plant sporopollenin. <i>Nature Plants</i> , 2019, 5, 41-46.	4.7	150

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37	Impact of acidic pH on plant cell wall polysaccharide structure and dynamics: insights into the mechanism of acid growth in plants from solid-state NMR. <i>Cellulose</i> , 2019, 26, 291-304.	2.4	56
38	Interplay between membrane curvature and protein conformational equilibrium investigated by solid-state NMR. <i>Journal of Structural Biology</i> , 2019, 206, 20-28.	1.3	10
39	Fast Magic-Angle-Spinning ¹⁹ F Spin Exchange NMR for Determining Nanometer ¹⁹ F- ¹⁹ F Distances in Proteins and Pharmaceutical Compounds. <i>Journal of Physical Chemistry B</i> , 2018, 122, 2900-2911.	1.2	49
40	Structure and Dynamics of Membrane Proteins from Solid-State NMR. <i>Annual Review of Biophysics</i> , 2018, 47, 201-222.	4.5	105
41	Sodium butyrate alleviates LPS-induced acute lung injury in mice via inhibiting HMGB1 release. <i>International Immunopharmacology</i> , 2018, 56, 242-248.	1.7	50
42	Conformation and Trimer Association of the Transmembrane Domain of the Parainfluenza Virus Fusion Protein in Lipid Bilayers from Solid-State NMR: Insights into the Sequence Determinants of Trimer Structure and Fusion Activity. <i>Journal of Molecular Biology</i> , 2018, 430, 695-709.	2.0	16
43	Transport-Relevant Protein Conformational Dynamics and Water Dynamics on Multiple Time Scales in an Archetypal Proton Channel: Insights from Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2018, 140, 1514-1524.	6.6	25
44	Direct Determination of Hydroxymethyl Conformations of Plant Cell Wall Cellulose Using ¹ H Polarization Transfer Solid-State NMR. <i>Biomacromolecules</i> , 2018, 19, 1485-1497.	2.6	44
45	Structural factors affecting ¹³ C NMR chemical shifts of cellulose: a computational study. <i>Cellulose</i> , 2018, 25, 23-36.	2.4	45
46	Determining Cholesterol Binding to Membrane Proteins by Cholesterol ¹³ C Labeling in Yeast and Dynamic Nuclear Polarization NMR. <i>Journal of the American Chemical Society</i> , 2018, 140, 15437-15449.	6.6	48
47	Determination of Long-Range Distances by Fast Magic-Angle-Spinning Radiofrequency-Driven ¹⁹ F- ¹⁹ F Dipolar Recoupling NMR. <i>Journal of Physical Chemistry B</i> , 2018, 122, 9302-9313.	1.2	37
48	Rapid measurement of long-range distances in proteins by multidimensional ¹³ C- ¹⁹ F REDOR NMR under fast magic-angle spinning. <i>Journal of Biomolecular NMR</i> , 2018, 71, 31-43.	1.6	44
49	Efficient ¹⁵ N- ¹³ C Polarization Transfer by Third-Spin-Assisted Pulsed Cross-Polarization Magic-Angle-Spinning NMR for Protein Structure Determination. <i>Journal of Physical Chemistry B</i> , 2018, 122, 8367-8379.	1.2	12
50	Oligomeric Structure and Three-Dimensional Fold of the HIV gp41 Membrane-Proximal External Region and Transmembrane Domain in Phospholipid Bilayers. <i>Journal of the American Chemical Society</i> , 2018, 140, 8246-8259.	6.6	43
51	Water Distribution, Dynamics, and Interactions with Alzheimer's ¹²⁵ I ² -Amyloid Fibrils Investigated by Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2017, 139, 6242-6252.	6.6	77
52	Structural Basis for Asymmetric Conductance of the Influenza M2 Proton Channel Investigated by Solid-State NMR Spectroscopy. <i>Journal of Molecular Biology</i> , 2017, 429, 2192-2210.	2.0	29
53	Zinc-binding structure of a catalytic amyloid from solid-state NMR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 6191-6196.	3.3	102
54	Icephobic Surfaces Induced by Interfacial Nonfrozen Water. <i>ACS Applied Materials & Interfaces</i> , 2017, 9, 4202-4214.	4.0	138

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55	Protonation equilibria and pore-opening structure of the dual-histidine influenza B virus M2 transmembrane proton channel from solid-state NMR. <i>Journal of Biological Chemistry</i> , 2017, 292, 17876-17884.	1.6	22
56	A structural and mechanistic study of Î€-clamp-mediated cysteine perfluoroarylation. <i>Scientific Reports</i> , 2017, 7, 7954.	1.6	20
57	Effects of Pectin Molecular Weight Changes on the Structure, Dynamics, and Polysaccharide Interactions of Primary Cell Walls of <i>Arabidopsis thaliana</i> : Insights from Solid-State NMR. <i>Biomacromolecules</i> , 2017, 18, 2937-2950.	2.6	69
58	Cholesterol-binding site of the influenza M2 protein in lipid bilayers from solid-state NMR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 12946-12951.	3.3	85
59	² H- ¹³ C correlation solid-state NMR for investigating dynamics and water accessibilities of proteins and carbohydrates. <i>Journal of Biomolecular NMR</i> , 2017, 68, 257-270.	1.6	17
60	Gradients in Wall Mechanics and Polysaccharides along Growing Inflorescence Stems. <i>Plant Physiology</i> , 2017, 175, 1593-1607.	2.3	82
61	The Target of Î²-Expansin EXPB1 in Maize Cell Walls from Binding and Solid-State NMR Studies. <i>Plant Physiology</i> , 2016, 172, 2107-2119.	2.3	41
62	Cellulose Structural Polymorphism in Plant Primary Cell Walls Investigated by High-Field 2D Solid-State NMR Spectroscopy and Density Functional Theory Calculations. <i>Biomacromolecules</i> , 2016, 17, 2210-2222.	2.6	94
63	The Influenza M2 Ectodomain Regulates the Conformational Equilibria of the Transmembrane Proton Channel: Insights from Solid-State Nuclear Magnetic Resonance. <i>Biochemistry</i> , 2016, 55, 5387-5397.	1.2	10
64	Multidimensional solid-state NMR spectroscopy of plant cell walls. <i>Solid State Nuclear Magnetic Resonance</i> , 2016, 78, 56-63.	1.5	55
65	Structural Polymorphism of Alzheimer's Î²-Amyloid Fibrils as Controlled by an E22 Switch: A Solid-State NMR Study. <i>Journal of the American Chemical Society</i> , 2016, 138, 9840-9852.	6.6	79
66	Acid activation mechanism of the influenza A M2 proton channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E6955-E6964.	3.3	81
67	Solid-State Nuclear Magnetic Resonance Investigation of the Structural Topology and Lipid Interactions of a Viral Fusion Protein Chimera Containing the Fusion Peptide and Transmembrane Domain. <i>Biochemistry</i> , 2016, 55, 6787-6800.	1.2	12
68	Conformationally selective multidimensional chemical shift ranges in proteins from a PACSY database purged using intrinsic quality criteria. <i>Journal of Biomolecular NMR</i> , 2016, 64, 115-130.	1.6	28
69	Solid-State NMR Investigation of the Conformation, Proton Conduction, and Hydration of the Influenza B Virus M2 Transmembrane Proton Channel. <i>Journal of the American Chemical Society</i> , 2016, 138, 8143-8155.	6.6	49
70	Efficient DNP NMR of membrane proteins: sample preparation protocols, sensitivity, and radical location. <i>Journal of Biomolecular NMR</i> , 2016, 64, 223-237.	1.6	65
71	Solid-state NMR investigations of cellulose structure and interactions with matrix polysaccharides in plant primary cell walls. <i>Journal of Experimental Botany</i> , 2016, 67, 503-514.	2.4	167
72	Chemical ligation of the influenza M2 protein for solid-state NMR characterization of the cytoplasmic domain. <i>Protein Science</i> , 2015, 24, 1087-1099.	3.1	36

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73	Relaxation-compensated difference spin diffusion NMR for detecting ^{13}C - ^{13}C long-range correlations in proteins and polysaccharides. <i>Journal of Biomolecular NMR</i> , 2015, 61, 97-107.	1.6	36
74	Cellulose-Pectin Spatial Contacts Are Inherent to Never-Dried Arabidopsis Primary Cell Walls: Evidence from Solid-State Nuclear Magnetic Resonance. <i>Plant Physiology</i> , 2015, 168, 871-884.	2.3	197
75	Investigation of the Curvature Induction and Membrane Localization of the Influenza Virus M2 Protein Using Static and Off-Magic-Angle Spinning Solid-State Nuclear Magnetic Resonance of Oriented Bicelles. <i>Biochemistry</i> , 2015, 54, 2214-2226.	1.2	29
76	The Influenza M2 Cytoplasmic Tail Changes the Proton-Exchange Equilibria and the Backbone Conformation of the Transmembrane Histidine Residue to Facilitate Proton Conduction. <i>Journal of the American Chemical Society</i> , 2015, 137, 6067-6077.	6.6	50
77	Distinguishing Bicontinuous Lipid Cubic Phases from Isotropic Membrane Morphologies Using ^{31}P Solid-State NMR Spectroscopy. <i>Journal of Physical Chemistry B</i> , 2015, 119, 4993-5001.	1.2	36
78	Aromatic spectral editing techniques for magic-angle-spinning solid-state NMR spectroscopy of uniformly ^{13}C -labeled proteins. <i>Solid State Nuclear Magnetic Resonance</i> , 2015, 72, 118-126.	1.5	16
79	Viral fusion protein transmembrane domain adopts β -strand structure to facilitate membrane topological changes for virus-cell fusion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 10926-10931.	3.3	54
80	^{15}N and ^1H Solid-State NMR Investigation of a Canonical Low-Barrier Hydrogen-Bond Compound: 1,8-Bis(dimethylamino)naphthalene. <i>Journal of Physical Chemistry B</i> , 2015, 119, 11581-11589.	1.2	12
81	De novo design of a transmembrane Zn^{2+} -transporting four-helix bundle. <i>Science</i> , 2014, 346, 1520-1524.	6.0	275
82	Probing membrane protein structure using water polarization transfer solid-state NMR. <i>Journal of Magnetic Resonance</i> , 2014, 247, 118-127.	1.2	58
83	Conformation and Lipid Interaction of the Fusion Peptide of the Paramyxovirus PIV5 in Anionic and Negative-Curvature Membranes from Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2014, 136, 2611-2624.	6.6	43
84	Cryoprotection of lipid membranes for high-resolution solid-state NMR studies of membrane peptides and proteins at low temperature. <i>Journal of Biomolecular NMR</i> , 2014, 59, 263-277.	1.6	22
85	Water-Polysaccharide Interactions in the Primary Cell Wall of <i>Arabidopsis thaliana</i> from Polarization Transfer Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2014, 136, 10399-10409.	6.6	111
86	Structure and Dynamics of <i>Brachypodium</i> Primary Cell Wall Polysaccharides from Two-Dimensional ^{13}C Solid-State Nuclear Magnetic Resonance Spectroscopy. <i>Biochemistry</i> , 2014, 53, 2840-2854.	1.2	68
87	Practical use of chemical shift databases for protein solid-state NMR: 2D chemical shift maps and amino-acid assignment with secondary-structure information. <i>Journal of Biomolecular NMR</i> , 2013, 56, 155-167.	1.6	55
88	Cationic membrane peptides: atomic-level insight of structure-activity relationships from solid-state NMR. <i>Amino Acids</i> , 2013, 44, 821-833.	1.2	57
89	pH-Dependent Conformation, Dynamics, and Aromatic Interaction of the Gating Tryptophan Residue of the Influenza M2 Proton Channel from Solid-State NMR. <i>Biophysical Journal</i> , 2013, 104, 1698-1708.	0.2	64
90	Sensitivity-enhanced solid-state NMR detection of expansin's target in plant cell walls. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 16444-16449.	3.3	196

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91	A 2H Solid-State NMR Study of Lipid Clustering by Cationic Antimicrobial and Cell-Penetrating Peptides in Model Bacterial Membranes. <i>Biophysical Journal</i> , 2013, 105, 2333-2342.	0.2	31
92	Resonance assignment of the NMR spectra of disordered proteins using a multi-objective non-dominated sorting genetic algorithm. <i>Journal of Biomolecular NMR</i> , 2013, 57, 281-296.	1.6	16
93	Conformational analysis of the full-length M2 protein of the influenza A virus using solid-state NMR. <i>Protein Science</i> , 2013, 22, 1623-1638.	3.1	51
94	Magic-Angle-Spinning NMR Techniques for Measuring Long-Range Distances in Biological Macromolecules. <i>Accounts of Chemical Research</i> , 2013, 46, 2154-2163.	7.6	63
95	Spectrally edited 2D ¹³ C ¹³ C NMR spectra without diagonal ridge for characterizing ¹³ C-enriched low-temperature carbon materials. <i>Journal of Magnetic Resonance</i> , 2013, 234, 112-124.	1.2	40
96	Membrane-Dependent Conformation, Dynamics, and Lipid Interactions of the Fusion Peptide of the Paramyxovirus PIV5 from Solid-State NMR. <i>Journal of Molecular Biology</i> , 2013, 425, 563-576.	2.0	36
97	Drug-Induced Conformational and Dynamical Changes of the S31N Mutant of the Influenza M2 Proton Channel Investigated by Solid-State NMR. <i>Journal of the American Chemical Society</i> , 2013, 135, 9885-9897.	6.6	63
98	Structural basis for proton conduction and inhibition by the influenza M2 protein. <i>Protein Science</i> , 2012, 21, 1620-1633.	3.1	124
99	Spectral editing of two-dimensional magic-angle-spinning solid-state NMR spectra for protein resonance assignment and structure determination. <i>Journal of Biomolecular NMR</i> , 2012, 54, 343-353.	1.6	25
100	Pectin-Cellulose Interactions in the <i>Arabidopsis</i> Primary Cell Wall from Two-Dimensional Magic-Angle-Spinning Solid-State Nuclear Magnetic Resonance. <i>Biochemistry</i> , 2012, 51, 9846-9856.	1.2	135
101	Aggregation and Dynamics of Oligocholate Transporters in Phospholipid Bilayers Revealed by Solid-State NMR Spectroscopy. <i>Langmuir</i> , 2012, 28, 17071-17078.	1.6	13
102	Membrane Protein Structure and Dynamics from NMR Spectroscopy. <i>Annual Review of Physical Chemistry</i> , 2012, 63, 1-24.	4.8	179
103	NMR Determination of Protein Partitioning into Membrane Domains with Different Curvatures and Application to the Influenza M2 Peptide. <i>Biophysical Journal</i> , 2012, 102, 787-794.	0.2	58
104	Intramolecular ¹ H- ¹³ C distance measurement in uniformly ¹³ C, ¹⁵ N labeled peptides by solid-state NMR. <i>Solid State Nuclear Magnetic Resonance</i> , 2012, 45-46, 51-58.	1.5	4
105	Hydrogen-Bonding Partner of the Proton-Conducting Histidine in the Influenza M2 Proton Channel Revealed From ¹ H Chemical Shifts. <i>Journal of the American Chemical Society</i> , 2012, 134, 14753-14755.	6.6	65
106	NMR Detection of pH-Dependent Histidine-Water Proton Exchange Reveals the Conduction Mechanism of a Transmembrane Proton Channel. <i>Journal of the American Chemical Society</i> , 2012, 134, 3703-3713.	6.6	143
107	Paramagnetic Cu(II) for Probing Membrane Protein Structure and Function: Inhibition Mechanism of the Influenza M2 Proton Channel. <i>Journal of the American Chemical Society</i> , 2012, 134, 8693-8702.	6.6	46
108	Multidimensional solid-state NMR studies of the structure and dynamics of pectic polysaccharides in uniformly ¹³ C-labeled <i>Arabidopsis</i> primary cell walls. <i>Magnetic Resonance in Chemistry</i> , 2012, 50, 539-550.	1.1	74

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109	Cellulose microfibril crystallinity is reduced by mutating C-terminal transmembrane region residues CESA1 ^{A903V} and CESA3 ^{T942I} of cellulose synthase. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 4098-4103.	3.3	165
110	Membrane-Dependent Effects of a Cytoplasmic Helix on the Structure and Drug Binding of the Influenza Virus M2 Protein. Journal of the American Chemical Society, 2011, 133, 11572-11579.	6.6	80
111	Conformational Disorder of Membrane Peptides Investigated from Solid-State NMR Line Widths and Line Shapes. Journal of Physical Chemistry B, 2011, 115, 10758-10767.	1.2	36
112	Specific Binding of Adamantane Drugs and Direction of Their Polar Amines in the Pore of the Influenza M2 Transmembrane Domain in Lipid Bilayers and Dodecylphosphocholine Micelles Determined by NMR Spectroscopy. Journal of the American Chemical Society, 2011, 133, 4274-4284.	6.6	100
113	Molecular Dynamics Simulation Directed Rational Design of Inhibitors Targeting Drug-Resistant Mutants of Influenza A Virus M2. Journal of the American Chemical Society, 2011, 133, 12834-12841.	6.6	127
114	Structures of \hat{I}^2 -Hairpin Antimicrobial Protegrin Peptides in Lipopolysaccharide Membranes: Mechanism of Gram Selectivity Obtained from Solid-State Nuclear Magnetic Resonance. Biochemistry, 2011, 50, 2072-2083.	1.2	43
115	Conformational plasticity of the influenza A M2 transmembrane helix in lipid bilayers under varying pH, drug binding, and membrane thickness. Biochimica Et Biophysica Acta - Biomembranes, 2011, 1808, 415-423.	1.4	77
116	Protonation, Tautomerization, and Rotameric Structure of Histidine: A Comprehensive Study by Magic-Angle-Spinning Solid-State NMR. Journal of the American Chemical Society, 2011, 133, 1534-1544.	6.6	228
117	Structure and Interactions of Plant Cell-Wall Polysaccharides by Two- and Three-Dimensional Magic-Angle-Spinning Solid-State NMR. Biochemistry, 2011, 50, 989-1000.	1.2	290
118	Structure and dynamics of cationic membrane peptides and proteins: Insights from solid-state NMR. Protein Science, 2011, 20, 641-655.	3.1	87
119	3D ^{13}C - ^{13}C - ^{13}C correlation NMR for de novo distance determination of solid proteins and application to a human \hat{I}^\pm -defensin. Journal of Magnetic Resonance, 2010, 202, 203-210.	1.2	30
120	Structure of the amantadine binding site of influenza M2 proton channels in lipid bilayers. Nature, 2010, 463, 689-692.	13.7	590
121	Mechanisms of Proton Conduction and Gating in Influenza M2 Proton Channels from Solid-State NMR. Science, 2010, 330, 505-508.	6.0	294
122	Orientation, Dynamics, and Lipid Interaction of an Antimicrobial Arylamide Investigated by ¹⁹ F and ³¹ P Solid-State NMR Spectroscopy. Journal of the American Chemical Society, 2010, 132, 9197-9205.	6.6	39
123	Membrane-Bound Dynamic Structure of an Arginine-Rich Cell-Penetrating Peptide, the Protein Transduction Domain of HIV TAT, from Solid-State NMR. Biochemistry, 2010, 49, 6009-6020.	1.2	92
124	Water-Protein Interactions of an Arginine-Rich Membrane Peptide in Lipid Bilayers Investigated by Solid-State Nuclear Magnetic Resonance Spectroscopy. Journal of Physical Chemistry B, 2010, 114, 4063-4069.	1.2	74
125	Resonance Assignment and Three-Dimensional Structure Determination of a Human \hat{I}^\pm -Defensin, HNP-1, by Solid-State NMR. Journal of Molecular Biology, 2010, 397, 408-422.	2.0	43
126	High-Resolution Orientation and Depth of Insertion of the Voltage-Sensing S4 Helix of a Potassium Channel in Lipid Bilayers. Journal of Molecular Biology, 2010, 401, 642-652.	2.0	34

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127	The Membrane-Bound Structure and Topology of a Human α -Defensin Indicate a Dimer Pore Mechanism for Membrane Disruption. <i>Biochemistry</i> , 2010, 49, 9770-9782.	1.2	76
128	Conformational Changes of an Ion Channel Detected Through Water-Protein Interactions Using Solid-State NMR Spectroscopy. <i>Journal of the American Chemical Society</i> , 2010, 132, 2378-2384.	6.6	73
129	$2D\ ^1H$ - ^{31}P solid-state NMR studies of the dependence of inter-bilayer water dynamics on lipid headgroup structure and membrane peptides. <i>Journal of Magnetic Resonance</i> , 2009, 196, 39-47.	1.2	30
130	Effects of amantadine on the dynamics of membrane-bound influenza A M2 transmembrane peptide studied by NMR relaxation. <i>Journal of Biomolecular NMR</i> , 2009, 45, 185-196.	1.6	38
131	High-resolution solid-state NMR of anisotropically mobile molecules under very low-power 1H decoupling and moderate magic-angle spinning. <i>Journal of Magnetic Resonance</i> , 2009, 199, 225-232.	1.2	5
132	Structure and Function of the Influenza A M2 Proton Channel. <i>Biochemistry</i> , 2009, 48, 7356-7364.	1.2	125
133	Accurate Measurement of Methyl ^{13}C Chemical Shifts by Solid-State NMR for the Determination of Protein Side Chain Conformation: The Influenza A M2 Transmembrane Peptide as an Example. <i>Journal of the American Chemical Society</i> , 2009, 131, 7806-7816.	6.6	21
134	Immobilization of the Influenza A M2 Transmembrane Peptide in Virus Envelope-Mimetic Lipid Membranes: A Solid-State NMR Investigation. <i>Biochemistry</i> , 2009, 48, 6361-6368.	1.2	61
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136	Effects of arginine density on the membrane-bound structure of a cationic antimicrobial peptide from solid-state NMR. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2009, 1788, 514-521.	1.4	21
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