Peter Schmieder

List of Publications by Year
in descending order

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129	5,572	57719	98753
papers	citations	44 h-index	67 g-index

136	136	136	6829
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	A software tool for the prediction of Xaa-Pro peptide bond conformations in proteins based on 13C chemical shift statistics. Journal of Biomolecular NMR, 2002, 24, 149-154.	1.6	308
2	Specific interactions between the syntrophin PDZ domain and voltage-gated sodium channels. Nature Structural Biology, 1998, 5, 19-24.	9.7	217
3	Protonâ€Detected Solidâ€State NMR Spectroscopy of Fibrillar and Membrane Proteins. Angewandte Chemie - International Edition, 2011, 50, 4508-4512.	7.2	179
4	Origin and diagenesis of polyphosphate in lake sediments: A 31PßšNMR study. Limnology and Oceanography, 2004, 49, 1-10.	1.6	160
5	HETLOC, an Efficient Method for Determining Heteronuclear Long-Range Couplings with Heteronuclei in Natural Abundance. Angewandte Chemie International Edition in English, 1991, 30, 1329-1331.	4.4	144
6	Dual epitope recognition by the VASP EVH1 domain modulates polyproline ligand specificity and binding affinity. EMBO Journal, 2000, 19, 4903-4914.	3.5	120
7	WW domain sequence activity relationships identified using ligand recognition propensities of 42 WW domains. Protein Science, 2003, 12, 491-500.	3.1	119
8	Application of nonlinear sampling schemes to COSY-type spectra. Journal of Biomolecular NMR, 1993, 3, 569-76.	1.6	97
9	Hassallidin A, a Glycosylated Lipopeptide with Antifungal Activity from the CyanobacteriumHassalliasp.. Journal of Natural Products, 2005, 68, 695-700.	1.5	97
10	Structural changes of TasA in biofilm formation of Bacillus subtilis. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 3237-3242.	3.3	97
11	Improved resolution in triple-resonance spectra by nonlinear sampling in the constant-time domain. Journal of Biomolecular NMR, 1994, 4, 483-490.	1.6	96
12	Hunting the eagle killer: A cyanobacterial neurotoxin causes vacuolar myelinopathy. Science, 2021, 371, .	6.0	96
13	High Resolution ¹H-Detected Solid-State NMR Spectroscopy of Protein Aliphatic Resonances: Access to Tertiary Structure Information. Journal of the American Chemical Society, 2010, 132, 15133-15135.	6.6	95
14	Thiocyclosporins: Preparation, Solution and Crystal Structure, and Immunosuppressive Activity. Helvetica Chimica Acta, 1991, 74, 1953-1990.	1.0	92
15	Small Molecule AKAP-Protein Kinase A (PKA) Interaction Disruptors That Activate PKA Interfere with Compartmentalized cAMP Signaling in Cardiac Myocytes. Journal of Biological Chemistry, 2011, 286, 9079-9096.	1.6	92
16	The Helix-Destabilizing Propensity Scale ofd-Amino Acids:Â The Influence of Side Chain Steric Effects. Journal of the American Chemical Society, 2000, 122, 4865-4870.	6.6	88
17	MUSIC in Triple-Resonance Experiments: Amino Acid Type-Selective 1Hâ€“15N Correlations. Journal of Magnetic Resonance, 1999, 141, 34-43.	1.2	82
18	Hyperphosphorylation of Glucosyl C6 Carbons and Altered Structure of Glycogen in the Neurodegenerative Epilepsy Lafora Disease. Cell Metabolism, 2013, 17, 756-767.	7.2	80

#	Article	IF	Citations
19	Heteronuclear Solution-State NMR Studies of the Chromophore in Cyanobacterial Phytochrome Cph1. Biochemistry, 2005, 44, 8244-8250.	1.2	78
20	Solution structure of the receptor tyrosine kinase EphB2 SAM domain and identification of two distinct homotypic interaction sites. Protein Science, 1999, 8, 1954-1961.	3.1	73
21	Probing protein-chromophore interactions in Cph1 phytochrome by mutagenesis. FEBS Journal, 2006, 273, 1415-1429.	2.2	72
22	NMR structure of the Wnt modulator protein Sclerostin. Biochemical and Biophysical Research Communications, 2009, 380, 160-165.	1.0	72
23	Real-Time Tracking of Phytochromeâ€™s Orientational Changes During Pr Photoisomerization. Journal of the American Chemical Society, 2012, 134, 1408-1411.	6.6	72
24	The companion of cellulose synthase 1 confers salt tolerance through a Tau-like mechanism in plants. Nature Communications, 2019, 10, 857.	5.8	71
25	Site-Specifically Phosphorylated Lysine Peptides. Journal of the American Chemical Society, 2014, 136, 13622-13628.	6.6	68
26	Enabling adoption of 2D-NMR for the higher order structure assessment of monoclonal antibody therapeutics. MAbs, 2019, 11, 94-105.	2.6	67
27	Chromophore Structure of Cyanobacterial Phytochrome Cph1 in the Pr State: Reconciling Structural and Spectroscopic Data by QM/MM Calculations. Biophysical Journal, 2009, 96, 4153-4163.	0.2	66
28	MUSIC, Selective Pulses, and Tuned Delays: Amino Acid Type-Selective 1Hâ€“15N Correlations, II. Journal of Magnetic Resonance, 2001, 148, 61-72.	1.2	64
29	Metalâ€Free, Regioselective Triazole Ligations that Deliver Locked cis Peptide Mimetics. Angewandte Chemie - International Edition, 2009, 48, 5042-5045.	7.2	64
30	Quantification of Maximum-Entropy Spectrum Reconstructions. Journal of Magnetic Resonance, 1997, 125, 332-339.	1.2	62
31	Hassallidin Bâ€”Second antifungal member of the Hassallidin family. Bioorganic and Medicinal Chemistry Letters, 2006, 16, 4220-4222.	1.0	57
32	High-Resolution Double-Quantum Deuterium Magic Angle Spinning Solid-State NMR Spectroscopy of Perdeuterated Proteins. Journal of the American Chemical Society, 2009, 131, 2-3.	6.6	56
33	Harnessing¹³C-labeledmyo-inositol to interrogate inositol phosphate messengers by NMR. Chemical Science, 2019, 10, 5267-5274.	3.7	56
34	Lightâ€Directed Protein Binding of a Biologically Relevant Î²â€Sheet. Angewandte Chemie - International Edition, 2009, 48, 6636-6639.	7.2	54
35	Photocontrol of Contracting Muscle Fibers. Angewandte Chemie - International Edition, 2011, 50, 7699-7702.	7.2	53
36	15N MAS NMR Studies of Cph1 Phytochrome:Â Chromophore Dynamics and Intramolecular Signal Transduction. Journal of Physical Chemistry B, 2006, 110, 20580-20585.	1.2	51

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37	Methods for Detection and Quantification of Polyphosphate and Polyphosphate Accumulating Microorganisms in Aquatic Sediments. International Review of Hydrobiology, 2008, 93, 1-30.	0.5	51
38	Conformational Analysis of thecis- andtrans-Isomers of FK506 by NMR and Molecular Dynamics. Helvetica Chimica Acta, 1991, 74, 1027-1047.	1.0	50
39	Determination of heteronuclear long-range couplings to heteronuclei in natural abundance by two- and three-dimensional NMR spectroscopy. Journal of Biomolecular NMR, 1991, 1, 403-420.	1.6	50
40	The solution structure of the N-terminal domain of E3L shows a tyrosine conformation that may explain its reduced affinity to Z-DNA in vitro. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 2712-2717.	3.3	50
41	Light-dependent dimerisation in the N-terminal sensory module of cyanobacterial phytochrome 1. FEBS Letters, 2005, 579, 3970-3974.	1.3	50
42	The structures of the active center in dark-adapted bacteriorhodopsin by solution-state NMR spectroscopy. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 9765-9770.	3.3	48
43	Vinylphosphonites for Staudinger-induced chemoselective peptide cyclization and functionalization. Chemical Science, 2019, 10, 6322-6329.	3.7	48
44	MUSIC and Aromatic Residues: Amino Acid Type-Selective 1Hâ€“15N Correlations, III. Journal of Magnetic Resonance, 2001, 153, 186-192.	1.2	46
45	Highly Functionalized Terpyridines as Competitive Inhibitors of AKAPâ€“PKA Interactions. Angewandte Chemie - International Edition, 2013, 52, 12187-12191.	7.2	46
46	Chemically Induced Vinylphosphonothiolate Electrophiles for Thiolâ€“Thiol Bioconjugations. Journal of the American Chemical Society, 2020, 142, 9544-9552.	6.6	46
47	Discovery of Low-Molecular-Weight Ligands for the AF6 PDZ Domain. Angewandte Chemie - International Edition, 2006, 45, 3790-3795.	7.2	41
48	Largeâ€scale purification of ribosomeâ€nascent chain complexes for biochemical and structural studies. FEBS Letters, 2009, 583, 2407-2413.	1.3	41
49	Stereochemical Elucidation of Natural Products from Residual Chemical Shift Anisotropies in a Liquid Crystalline Phase. Journal of the American Chemical Society, 2020, 142, 2301-2309.	6.6	41
50	Intradomain Allosteric Network Modulates Calcium Affinity of the C-Type Lectin Receptor Langerin. Journal of the American Chemical Society, 2016, 138, 12176-12186.	6.6	40
51	A modular toolkit to inhibit proline-rich motifâ€“mediated proteinâ€“protein interactions. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 5011-5016.	3.3	39
52	Interaction of the Antimicrobial Peptide Cyclo(RRWWRF) with Membranes by Molecular Dynamics Simulations. Biophysical Journal, 2005, 89, 2296-2306.	0.2	38
53	NMR spectroscopy reveals unexpected structural variation at the proteinâ€“protein interface in MHC class I molecules. Journal of Biomolecular NMR, 2013, 57, 167-178.	1.6	38
54	Heteronuclear Multidimensional NMR Spectroscopy of Solubilized Membrane Proteins: Resonance Assignment of Native Bacteriorhodopsin. ChemBioChem, 2002, 3, 1019-1023.	1.3	36

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55	Structure of the Antimicrobial, Cationic Hexapeptide Cyclo(RRWWRF) and Its Analogues in Solution and Bound to Detergent Micelles. ChemBioChem, 2005, 6, 1654-1662.	1.3	36
56	The Switch that Does Not Flip: The Blue-Light Receptor YtvA from Bacillus subtilis Adopts an Elongated Dimer Conformation Independent of the Activation State as Revealed by a Combined AUC and SAXS Study. Journal of Molecular Biology, 2010, 403, 78-87.	2.0	35
57	Active and silent chromophore isoforms for phytochrome Pr photoisomerization: An alternative evolutionary strategy to optimize photoreaction quantum yields. Structural Dynamics, 2014, 1, 014701.	0.9	35
58	Effects of Halide Ions on the Carbamidocyclophane Biosynthesis in Nostoc sp. CAVN2. Marine Drugs, 2016, 14, 21.	2.2	35
59	A convenient method for saponin isolation in tumour therapy. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 713-718.	1.2	34
60	The NMR structure of the 47-kDa dimeric enzyme 3,4-dihydroxy-2-butanone-4-phosphate synthase and ligand binding studies reveal the location of the active site. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 13025-13030.	3.3	33
61	Heteronuclear NMR Investigation on the Structure and Dynamics of the Chromophore Binding Pocket of the Cyanobacterial Phytochrome Cph1. Journal of the American Chemical Society, 2008, 130, 11170-11178.	6.6	33
62	Septin 9 negatively regulates ubiquitin-dependent downregulation of epidermal growth factor receptor. Journal of Cell Science, 2015, 128, 397-407.	1.2	32
63	Direct Experimental Evidence for Halogenâ€“Aryl Ï€â€…Interactions in Solution from Molecular Torsion Balances. Angewandte Chemie - International Edition, 2017, 56, 6454-6458.	7.2	32
64	Efficient Î±â€Helix Induction in a Linear Peptide Chain by Nâ€Capping with a Bridgedâ€tricyclic Diproline Analogue. Angewandte Chemie - International Edition, 2013, 52, 9539-9543.	7.2	31
65	Multicolor Caged dSTORM Resolves the Ultrastructure of Synaptic Vesicles in the Brain. Angewandte Chemie - International Edition, 2015, 54, 13230-13235.	7.2	31
66	Chemoselective synthesis and analysis of naturally occurring phosphorylated cysteine peptides. Nature Communications, 2016, 7, 12703.	5.8	31
67	The Oxidized Subunit B8 from Human Complex I Adopts a Thioredoxin Fold. Structure, 2004, 12, 1645-1654.	1.6	29
68	A General Oneâ€Pot Synthesis of 2Hâ€Indazoles Using an Organophosphorusâ€“Silane System. Chemistry - A European Journal, 2018, 24, 9090-9100.	1.7	29
69	Photoswitchable Click Amino Acids: Light Control of Conformation and Bioactivity. ChemBioChem, 2011, 12, 2555-2559.	1.3	28
70	Amino acid type-selective backbone 1H-15N-correlations for Arg and Lys. Journal of Biomolecular NMR, 2001, 20, 379-384.	1.6	27
71	Periplasmic Loop P2 of the MalF Subunit of the Maltose ATP Binding Cassette Transporter Is Sufficient To Bind the Maltose Binding Protein MalE. Biochemistry, 2009, 48, 2216-2225.	1.2	27
72	Dynamics of free versus complexed Î²2-microglobulin and the evolution of interfaces in MHC class I molecules. Immunogenetics, 2013, 65, 157-172.	1.2	27

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73	The helicase-binding domain of Escherichia coli DnaG primase interacts with the highly conserved C-terminal region of single-stranded DNA-binding protein. Nucleic Acids Research, 2013, 41, 4507-4517.	6.5	27
74	Structure of the competence pilus major pilin ComGC in Streptococcus pneumoniae. Journal of Biological Chemistry, 2017, 292, 14134-14146.	1.6	27
75	Measurements of H.alpha.-HN vicinal coupling constants in a protein with large line widths in a new 3D 1H-15N-13C quadruple resonance NMR experiment. Journal of the American Chemical Society, 1991, 113, 6323-6324.	6.6	26
76	Determination of the ? angle in a peptide backbone by NMR spectroscopy with a combination of homonuclear and heteronuclear coupling constants. Biopolymers, 1992, 32, 435-440.	1.2	26
77	Solution Structure, Backbone Dynamics, and Association Behavior of the C-Terminal BRCT Domain from the Breast Cancer-Associated Protein BRCA1â€‚â€¡. Biochemistry, 2004, 43, 15983-15995.	1.2	26
78	The depsipeptide technique applied to peptide segment condensation: Scope and limitations. Journal of Peptide Science, 2008, 14, 299-306.	0.8	25
79	Solid-Phase Synthesis of a Cyclodepsipeptide: Cotransin. Organic Letters, 2008, 10, 3857-3860.	2.4	25
80	Direct access to site-specifically phosphorylated-lysine peptides from a solid-support. Organic and Biomolecular Chemistry, 2015, 13, 6839-6843.	1.5	25
81	The E. coli Siderophores Enterobactin and Salmochelin Form Sixâ€Coordinate Silicon Complexes at Physiological pH. Angewandte Chemie - International Edition, 2011, 50, 4230-4233.	7.2	23
82	NMR quality control of fragment libraries for screening. Journal of Biomolecular NMR, 2020, 74, 555-563.	1.6	23
83	A Computational Modeling Approach Predicts Interaction of the Antifungal Protein AFP from Aspergillus giganteus with Fungal Membranes via Its Î³-Core Motif. MSphere, 2018, 3, .	1.3	22
84	3D Heteronuclear NMR techniques for carbon-13 in natural abundance. Journal of the American Chemical Society, 1990, 112, 8599-8600.	6.6	21
85	A modified strategy for sequence specific assignment of protein NMR spectra based on amino acid type selective experiments. Journal of Biomolecular NMR, 2005, 31, 115-128.	1.6	21
86	Solutionâ€State ¹⁵N NMR Spectroscopic Study of Î±â€Câ€Phycocyanin: Implications for the Structure of the Chromophoreâ€Binding Pocket of the Cyanobacterial Phytochrome Cph1. ChemBioChem, 2007, 8, 2249-2255.	1.3	21
87	Blue Flickers of Hope: Secondary Structure, Dynamics, and Putative Dimerization Interface of the Blue-Light Receptor YtvA from Bacillus subtilis. Biochemistry, 2011, 50, 8163-8171.	1.2	21
88	Designed nanomolar small-molecule inhibitors of Ena/VASP EVH1 interaction impair invasion and extravasation of breast cancer cells. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 29684-29690.	3.3	21
89	Structures of cyclic, antimicrobial peptides in a membraneâ€mimicking environment define requirements for activity. Journal of Peptide Science, 2008, 14, 524-527.	0.8	20
90	Bridging the gap: A set of selective 1H-15N-correlations to link sequential neighbors of prolines. Journal of Biomolecular NMR, 2000, 17, 331-335.	1.6	18

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91	The blue-light receptor YtvA from Bacillus subtilis is permanently incorporated into the stressosome independent of the illumination state. Biochemical and Biophysical Research Communications, 2013, 432, 499-503.	1.0	18
92	NMR Spectroscopic Investigation of Mobility and Hydrogen Bonding of the Chromophore in the Binding Pocket of Phytochrome Proteins. ChemPhysChem, 2010, 11, 1248-1257.	1.0	17
93	Controlled thioamide vs. amide formation in the thioacidâ€“azide reaction under acidic aqueous conditions. Chemical Communications, 2014, 50, 4603.	2.2	17
94	Fast Heteronuclear 3D NMR Spectroscopy. Angewandte Chemie International Edition in English, 1990, 29, 546-548.	4.4	16
95	Backbone and sidechain 1H, 13C and 15N resonance assignments of the Bright/ARID domain from the human JARID1C (SMCX) protein. Biomolecular NMR Assignments, 2008, 2, 9-11.	0.4	16
96	pHâ€Ðependent Protonation of Surface Carboxylate Groups in PsbO Enables Local Buffering and Triggers Structural Changes. ChemBioChem, 2020, 21, 1597-1604.	1.3	16
97	Design of antimicrobial compounds based on peptide structures. Bioorganic and Medicinal Chemistry Letters, 2007, 17, 2334-2337.	1.0	12
98	Design, synthesis, structure and binding properties of PDZ binding, cyclic Î²-finger peptides. Biochemical and Biophysical Research Communications, 2010, 395, 535-539.	1.0	12
99	Sclerotiorin Stabilizes the Assembly of Nonfibrillar Abeta42 Oligomers with Low Toxicity, Seeding Activity, and Beta-sheet Content. Journal of Molecular Biology, 2020, 432, 2080-2098.	2.0	12
100	Unraveling the existence of dynamic water channels in light-harvesting proteins: alpha-C-phycocyanobilin in vitro. Chemical Science, 2013, 4, 755-763.	3.7	11
101	LOV Takes a Pick: Thermodynamic and Structural Aspects of the Flavin-LOV-Interaction of the Blue-Light Sensitive Photoreceptor YtvA from Bacillus subtilis. PLoS ONE, 2013, 8, e81268.	1.1	11
102	J-Deconvolution Using Maximum Entropy Reconstruction Applied to13Câ^'13C Solid-State Cross-Polarization Magic-Angle-Spinning NMR of Proteins. Journal of the American Chemical Society, 2007, 129, 6682-6683.	6.6	10
103	Structural and dynamic features of HLA-B27 subtypes. Current Opinion in Rheumatology, 2013, 25, 411-418.	2.0	10
104	Schnelle Heterokernâ€3Dâ€NMRâ€Spektroskopie. Angewandte Chemie, 1990, 102, 588-589.	1.6	9
105	Isolation of Microcystins from the Cyanobacterium Planktothrix rubescens Strain No80. Natural Products and Bioprospecting, 2014, 4, 37-45.	2.0	9
106	Sapofectosid â€“ Ensuring non-toxic and effective DNA and RNA delivery. International Journal of Pharmaceutics, 2017, 534, 195-205.	2.6	9
107	High yield expression and purification of isotopically labelled human endothelin-1 for use in NMR studies. Protein Expression and Purification, 2006, 48, 253-260.	0.6	8
108	The Structure of MESD45â€“184 Brings Light into the Mechanism of LDLR Family Folding. Structure, 2011, 19, 337-348.	1.6	8

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109	Metal-triggered conformational reorientation of a self-peptide bound to a disease-associated HLA-B*27 subtype. Journal of Biological Chemistry, 2019, 294, 13269-13279.	1.6	8
110	Ambigols from the Cyanobacterium Fischerella ambigua Increase Prodigiosin Production in Serratia spp. ACS Chemical Biology, 2020, 15, 2929-2936.	1.6	8
111	The solution structure of the core of mesoderm development (MESD), a chaperone for members of the LDLR-family. Journal of Structural and Functional Genomics, 2007, 7, 131-138.	1.2	7
112	Smallâ€Molecule Inhibitors of AF6 PDZâ€Mediated Proteinâ€“Protein Interactions. ChemMedChem, 2014, 9, 1458-1462.	1.6	7
113	Synthesis and Evaluation of Nonâ€Hydrolyzable Phosphoâ€Lysine Peptide Mimics. Chemistry - A European Journal, 2021, 27, 2326-2331.	1.7	7
114	Blue News Update: BODIPY-GTP Binds to the Blue-Light Receptor YtvA While GTP Does Not. PLoS ONE, 2012, 7, e29201.	1.1	7
115	Chemical synthesis of the third WW domain of TCERG 1 by native chemical ligation. Journal of Peptide Science, 2011, 17, 644-649.	0.8	6
116	Determination of glucan phosphorylation using heteronuclear ¹H,¹³C double and ¹H,¹³C,³¹P tripleâ€resonance NMR spectra. Magnetic Resonance in Chemistry, 2013, 51, 655-661.	1.1	6
117	Plant derived triterpenes from Gypsophila elegans M.Bieb. enable non-toxic delivery of gene loaded nanoplexes. Journal of Biotechnology, 2018, 284, 131-139.	1.9	6
118	A new acetylated triterpene saponin from Agrostemma githago L. modulates gene delivery efficiently and shows a high cellular tolerance. International Journal of Pharmaceutics, 2020, 589, 119822.	2.6	5
119	The solution structure of an N-terminally truncated version of the yeast CDC24p PB1 domain shows a different Î²-sheet topology. FEBS Letters, 2005, 579, 3534-3538.	1.3	4
120	Direct Experimental Evidence for Halogenâ€“Aryl Ï€â€…Interactions in Solution from Molecular Torsion Balances. Angewandte Chemie, 2017, 129, 6554-6558.	1.6	3
121	NMR assignments of the periplasmic loop P2 of the MalF subunit of the maltose ATP binding cassette transporter. Biomolecular NMR Assignments, 2009, 3, 21-23.	0.4	2
122	Assignment of phycocyanobilin in HMPT using triple resonance experiments. Magnetic Resonance in Chemistry, 2011, 49, 543-548.	1.1	2
123	Crystal structure of Q4D6Q6, a conserved kinetoplastid-specific protein from Trypanosoma cruzi. Journal of Structural Biology, 2020, 211, 107536.	1.3	2
124	Synthesis and Evaluation of Nonâ€Hydrolyzable Phosphoâ€Lysine Peptide Mimics. Chemistry - A European Journal, 2021, 27, 2223-2223.	1.7	2
125	Chemical shift assignments and secondary structure prediction for Q4DY78, a conserved kinetoplastid-specific protein from Trypanosoma cruzi. Biomolecular NMR Assignments, 2016, 10, 325-328.	0.4	1
126	Resonance assignment of the RGS domain of human RGS10. Journal of Biomolecular NMR, 2007, 38, 191-191.	1.6	0

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127	Backbone and sidechain 1H, 13C and 15N resonance assignments of the RGS domain from human RGS14. Biomolecular NMR Assignments, 2007, 1, 95-97.	0.4	0
128	NMR structure and dynamics of Q4DY78, a conserved kinetoplasid-specific protein from Trypanosoma cruzi. Journal of Structural Biology, 2021, 213, 107715.	1.3	0
129	How solvent-free crosslinking conditions alter the chemistry and topology of hemiketal based polymer networks. Polymer, 2021, 229, 123986.	1.8	0