## Frank D Sönnichsen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4568850/publications.pdf Version: 2024-02-01

		53794	62596
147	7,217	45	80
papers	citations	h-index	g-index
151	151	151	7603
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Effect of trifluoroethanol on protein secondary structure: an NMR and CD study using a synthetic actin peptide. Biochemistry, 1992, 31, 8790-8798.	2.5	638
2	Magnetic Bistability of Molecules in Homogeneous Solution at Room Temperature. Science, 2011, 331, 445-448.	12.6	489
3	Molecular architecture of human prion protein amyloid: A parallel, in-register β-structure. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 18946-18951.	7.1	302
4	Solution NMR of membrane proteins: practice and challenges. Magnetic Resonance in Chemistry, 2006, 44, S24-S40.	1.9	210
5	Solution Nuclear Magnetic Resonance Structure of Membrane-Integral Diacylglycerol Kinase. Science, 2009, 324, 1726-1729.	12.6	205
6	A Diminished Role for Hydrogen Bonds in Antifreeze Protein Binding to Iceâ€. Biochemistry, 1997, 36, 14652-14660.	2.5	204
7	Structure of KCNE1 and Implications for How It Modulates the KCNQ1 Potassium Channel. Biochemistry, 2008, 47, 7999-8006.	2.5	183
8	Refined solution structure of type III antifreeze protein: hydrophobic groups may be involved in the energetics of the protein–ice interaction. Structure, 1996, 4, 1325-1337.	3.3	177
9	The Entner–Doudoroff pathway is an overlooked glycolytic route in cyanobacteria and plants. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 5441-5446.	7.1	160
10	Intramolecular Quenching of Tryptophan Fluorescence by the Peptide Bond in Cyclic Hexapeptides. Journal of the American Chemical Society, 2002, 124, 9278-9286.	13.7	141
11	Phosphatidylserine exposure is required for ADAM17 sheddase function. Nature Communications, 2016, 7, 11523.	12.8	134
12	The Solution Structure of Type II Antifreeze Protein Reveals a New Member of the Lectin Family. Biochemistry, 1998, 37, 4712-4721.	2.5	125
13	Solution Structure of the E200K Variant of Human Prion Protein. Journal of Biological Chemistry, 2000, 275, 33650-33654.	3.4	120
14	Structureâ€function relationship in the globular type III antifreeze protein: Identification of a cluster of surface residues required for binding to ice. Protein Science, 1994, 3, 1760-1769.	7.6	119
15	The nonhelical structure of antifreeze protein type III. Science, 1993, 259, 1154-1157.	12.6	111
16	Photoswitchable Magnetic Resonance Imaging Contrast by Improved Light-Driven Coordination-Induced Spin State Switch. Journal of the American Chemical Society, 2015, 137, 7552-7555.	13.7	110
17	On choosing a detergent for solution NMR studies of membrane proteins. Journal of Biomolecular NMR, 1998, 11, 381-386.	2.8	107
18	Preferential Heterodimeric Parallel Coiled-coil Formation by Synthetic Max and c-Myc Leucine Zippers: A Description of Putative Electrostatic Interactions Responsible for the Specificity of Heterodimerization. Journal of Molecular Biology, 1995, 254, 505-520.	4.2	106

#	Article	IF	CITATIONS
19	Coordinationâ€Induced Spin Crossover (CISCO) through Axial Bonding of Substituted Pyridines to Nickel–Porphyrins: Ïfâ€Donor versus Ï€â€Acceptor Effects. Chemistry - A European Journal, 2010, 16, 10074-10083.	3.3	106
20	Nitrogen Bridged Diazocines: Photochromes Switching within the Near-Infrared Region with High Quantum Yields in Organic Solvents and in Water. Journal of the American Chemical Society, 2019, 141, 13592-13600.	13.7	101
21	Reversed-phase chromatography of synthetic amphipathic α-helical peptides as a model for ligand/receptor interactions Effect of changing hydrophobic environment on the relative hydrophilicity/hydrophobicity of amino acid side-chains. Journal of Chromatography A, 1994, 676, 139-153.	3.7	99
22	NMR Assignments for a Helical 40 kDa Membrane Protein. Journal of the American Chemical Society, 2004, 126, 5048-5049.	13.7	86
23	Interhelical Salt Bridges, Coiled-Coil Stability, and Specificity of Dimerization. Science, 1996, 271, 1136-1138.	12.6	85
24	Mass and NMR spectroscopic characterization of 3,4-methylenedioxypyrovalerone: A designer drug with α-pyrrolidinophenone structure. Forensic Science International, 2009, 190, 1-8.	2.2	78
25	Comprehensive evaluation of solution nuclear magnetic resonance spectroscopy sample preparation for helical integral membrane proteins. Journal of Structural and Functional Genomics, 2006, 7, 51-64.	1.2	77
26	Membrane-Proximal Domain of a Disintegrin and Metalloprotease-17 Represents the Putative Molecular Switch of Its Shedding Activity Operated by Protein-disulfide Isomerase. Journal of the American Chemical Society, 2013, 135, 5776-5781.	13.7	75
27	Structure Effects of Double D-Amino Acid Replacements: A Nuclear Magnetic Resonance and Circular Dichroism Study Using Amphipathic Model Helixes. Biochemistry, 1995, 34, 12954-12962.	2.5	73
28	Binding of an Oligopeptide to a Specific Plane of Ice. Journal of Biological Chemistry, 1998, 273, 11714-11718.	3.4	67
29	Fe <sup>III</sup> Spinâ€Crossover Complexes with Photoisomerizable Ligands: Experimental and Theoretical Studies on the Ligandâ€Driven Lightâ€Induced Spin Change Effect. European Journal of Inorganic Chemistry, 2012, 2012, 2776-2783.	2.0	66
30	Rational design of a room temperature molecular spin switch. The light-driven coordination induced spin state switch (LD-CISSS) approach. Dalton Transactions, 2014, 43, 17395-17405.	3.3	66
31	New polyphenolic compounds in commercial deodistillate and rapeseed oils. Food Chemistry, 2010, 123, 607-615.	8.2	64
32	NMR Characterization of Side Chain Flexibility and Backbone Structure in the Type I Antifreeze Protein at Near Freezing Temperaturesâ€. Biochemistry, 1996, 35, 16698-16704.	2.5	56
33	High molecular weight poly(N-methyl-B-vinylazaborine) – a semi-inorganic B–N polystyrene analogue. Chemical Communications, 2017, 53, 7258-7261.	4.1	56
34	High Resolution Solution Structure of the 1.3S Subunit of Transcarboxylase from Propionibacterium shermaniiâ€. Biochemistry, 2000, 39, 2509-2516.	2.5	55
35	Use of proline mutants to help solve the NMR solution structure of type III antifreeze protein. Protein Science, 1993, 2, 1411-1428.	7.6	54
36	Temperature coefficients of amide proton NMR resonance frequencies in trifluoroethanol: A monitor of intramolecular hydrogen bonds in helical peptides?. Journal of Biomolecular NMR. 1996. 8. 93-97.	2.8	53

#	Article	IF	CITATIONS
37	A new class of hexahelical insect proteins revealed as putative carriers of small hydrophobic ligands. Structure, 1999, 7, 1325-1332.	3.3	51
38	Light-controlled switching of the spin state of iron(III). Nature Communications, 2018, 9, 4750.	12.8	51
39	Entry to " <mml:math <br="" xmlns:mml="http://www.w3.org/1998/Math/MathML">altimg="si1.gif"&gt;<mml:mrow> <mml:mrow> <mml:mrow> <mml:mtext>HCO</mml:mtext> </mml:mrow> <mr Tunnel―Revealed by SLC4A4 Human Mutation and Structural Model. Journal of Biological Chemistry, 2008. 283. 18402-18410.</mr </mml:mrow></mml:mrow></mml:math>	nl:mtext>3	
40	Differences in heat stability and ligand binding among β-lactoglobulin genetic variants A, B and C using 1H NMR and fluorescence quenching. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 1083-1093.	2.3	50
41	Comparison of the solution structures of microcystin-LR and motuporin. Nature Structural and Molecular Biology, 1995, 2, 114-116.	8.2	49
42	Topology and Secondary Structure of the N-Terminal Domain of Diacylglycerol Kinaseâ€. Biochemistry, 2002, 41, 12876-12882.	2.5	49
43	CAMRA: chemical shift based computer aided protein NMR assignments. Journal of Biomolecular NMR, 1998, 12, 395-405.	2.8	46
44	The Heme Complex of Hmu O, a Bacterial Heme Degradation Enzyme from Corynebacterium diphtheriae. Journal of Biological Chemistry, 1999, 274, 24490-24496.	3.4	45
45	Comparative modeling of the threeâ€dimensional structure of Type II antifreeze protein. Protein Science, 1995, 4, 460-471.	7.6	45
46	Photochemical properties of multi-azobenzene compounds. Photochemical and Photobiological Sciences, 2013, 12, 511-518.	2.9	45
47	The Quiet Renaissance of Protein Nuclear Magnetic Resonance. Biochemistry, 2013, 52, 1303-1320.	2.5	45
48	Alternative Roles for Putative Ice-Binding Residues in Type I Antifreeze Proteinâ€. Biochemistry, 1999, 38, 4743-4749.	2.5	43
49	The Ice-Binding Site of Sea Raven Antifreeze Protein Is Distinct from the Carbohydrate-Binding Site of the Homologous C-Type Lectinâ€. Biochemistry, 1998, 37, 17745-17753.	2.5	41
50	A Structural Model for the Membrane-bound Form of the Juxtamembrane Domain of the Epidermal Growth Factor Receptor. Journal of Biological Chemistry, 2005, 280, 24043-24052.	3.4	41
51	Macin Family of Antimicrobial Proteins Combines Antimicrobial and Nerve Repair Activities. Journal of Biological Chemistry, 2012, 287, 14246-14258.	3.4	41
52	Identification of 1-butyl-3-(1-(4-methyl)naphthoyl)indole in a herbal mixture. Forensic Science International, 2012, 215, 8-13.	2.2	41
53	The Role of Transient Changes in Sample Susceptibility in Causing Apparent Multiple-Quantum Peaks in HOESY Spectra. Journal of Magnetic Resonance Series A, 1996, 121, 83-87.	1.6	39
54	The relative positions of alanine residues in the hydrophobic core control the formation of two-stranded or four-stranded α-helical coiled-coils. Protein Engineering, Design and Selection, 1996, 9, 353-363.	2.1	39

#	Article	IF	CITATIONS
55	Diversely halogenated spiropyrans - Useful synthetic building blocks for a versatile class of molecular switches. Dyes and Pigments, 2017, 136, 292-301.	3.7	39
56	Unexpected Advanced Generation Cephalosporinase Activity of the M69F Variant of SHV β-Lactamase. Journal of Biological Chemistry, 2002, 277, 47719-47723.	3.4	37
57	The α-Helical Peptide Nucleic Acid Concept: Merger of Peptide Secondary Structure and Codified Nucleic Acid Recognition. Journal of the American Chemical Society, 2004, 126, 4626-4640.	13.7	37
58	Source of the Ice-Binding Specificity of Antifreeze Protein Type I. Journal of Chemical Information and Computer Sciences, 2000, 40, 1276-1284.	2.8	36
59	Structure and function of a unique pore-forming protein from a pathogenic acanthamoeba. Nature Chemical Biology, 2013, 9, 37-42.	8.0	36
60	Characterization and Function of the First Antibiotic Isolated from a Vent Organism: The Extremophile Metazoan Alvinella pompejana. PLoS ONE, 2014, 9, e95737.	2.5	36
61	NMR solution structure and flexibility of a peptide antigen representing the receptor binding domain of Pseudomonas aeruginosa. Biochemistry, 1993, 32, 13432-13440.	2.5	35
62	Membrane Protein Preparation for TROSY NMR Screening. Methods in Enzymology, 2005, 394, 321-334.	1.0	35
63	Parallel ultrafast E–C ring closure and E–Z isomerisation in a photochromic furylfulgide studied by femtosecond time-resolved spectroscopy. Physical Chemistry Chemical Physics, 2009, 11, 5952.	2.8	35
64	Purification, Characterization, and Structural Analysis of a Plant Low-Temperature-Induced Protein. Plant Physiology, 1997, 113, 367-376.	4.8	31
65	Identification of rosmarinic acid and sulfated flavonoids as inhibitors of microfouling on the surface of eelgrass <i>Zostera marina</i> . Biofouling, 2017, 33, 867-880.	2.2	31
66	Modeling Studies of Binding of Sea Raven Type II Antifreeze Protein to Ice. Journal of Chemical Information and Computer Sciences, 1997, 37, 1006-1010.	2.8	30
67	Seized designer supplement named "1-Androsterone― Identification as 3β-hydroxy-5α-androst-1-en-17-one and its urinary elimination. Steroids, 2011, 76, 540-547.	1.8	30
68	Occurrence of Fusarium Mycotoxins and Their Modified Forms in Forage Maize Cultivars. Toxins, 2021, 13, 110.	3.4	30
69	The first supramolecular ion triplet complex. New Journal of Chemistry, 2010, 34, 1247.	2.8	28
70	Thermodynamic Stabilization of the Folded Domain of Prion Protein Inhibits Prion Infection inÂVivo. Cell Reports, 2013, 4, 248-254.	6.4	28
71	Synthesis of Bifunctional Azobenzene Glycoconjugates for Cysteineâ€Based Photosensitive Cross‣inking with Bioactive Peptides. Chemistry - A European Journal, 2015, 21, 13723-13731.	3.3	28
72	A Paramagnetic NMR Spectroscopy Toolbox for the Characterisation of Paramagnetic/Spin rossover Coordination Complexes and Metal–Organic Cages. Angewandte Chemie - International Edition, 2020, 59, 19344-19351.	13.8	27

#	Article	IF	CITATIONS
73	Photochromism of Rotationâ€Hindered Furylfulgides Influenced by Steric Modifications. European Journal of Organic Chemistry, 2011, 2011, 1947-1955.	2.4	26
74	Different Secondary Metabolite Profiles of Phylogenetically almost Identical Streptomyces griseus Strains Originating from Geographically Remote Locations. Microorganisms, 2019, 7, 166.	3.6	25
75	Longâ€Distance Rate Acceleration by Bulk Gold. Angewandte Chemie - International Edition, 2019, 58, 6574-6578.	13.8	25
76	Continuous Radio Amplification by Stimulated Emission of Radiation using Parahydrogen Induced Polarization (PHIPâ€RASER) at 14 Tesla. ChemPhysChem, 2020, 21, 667-672.	2.1	25
77	Structural characterization of the entire 1.3S subunit of transcarboxylase from <i>Propionibacterium shermanii</i> . Protein Science, 1998, 7, 2156-2163.	7.6	24
78	Conformationally Specific Misfolding of an Integral Membrane Proteinâ€. Biochemistry, 2001, 40, 5111-5118.	2.5	24
79	Specificity Determinants of a Novel Nck Interaction with the Juxtamembrane Domain of the Epidermal Growth Factor Receptor,. Biochemistry, 2008, 47, 3096-3108.	2.5	24
80	Negishi's Reagent Versus Rosenthal's Reagent in the Formation of Zirconacyclopentadienes. Chemistry - A European Journal, 2019, 25, 13318-13328.	3.3	24
81	A novel dileucine lysosomal-sorting-signal mediates intracellular EGF-receptor retention independently of protein ubiquitylation. Journal of Cell Science, 2005, 118, 3959-3971.	2.0	23
82	Adenovirus RIDα regulates endosome maturation by mimicking GTP-Rab7. Journal of Cell Biology, 2007, 179, 965-980.	5.2	23
83	Synthesis of glycocluster peptides. Carbohydrate Research, 2008, 343, 1665-1674.	2.3	22
84	Singly and Doubly Twisted [36]Annulenes: Synthesis and Calculations. Chemistry - A European Journal, 2010, 16, 7767-7772.	3.3	22
85	Synthesis and Photochromic Properties of Configurationally Varied Azobenzene Glycosides. ChemistryOpen, 2014, 3, 99-108.	1.9	19
86	Evaluation of High-Resolution Mass Spectrometry for the Quantitative Analysis of Mycotoxins in Complex Feed Matrices. Toxins, 2019, 11, 531.	3.4	19
87	Towards a light driven molecular assembler. Communications Chemistry, 2019, 2, .	4.5	19
88	Absence of Observable Biotinâ^'Protein Interactions in the 1.3S Subunit of Transcarboxylase:Â An NMR Studyâ€. Biochemistry, 1997, 36, 14676-14682.	2.5	18
89	ORB, a homology-based program for the prediction of protein NMR chemical shifts. Journal of Biomolecular NMR, 1997, 10, 165-179.	2.8	18
90	NMR structural studies on antifreeze proteins. Biochemistry and Cell Biology, 1998, 76, 284-293.	2.0	17

#	Article	IF	CITATIONS
91	A Membrane Setting for the Sorting Motifs Present in the Adenovirus E3-13.7 Protein Which Down-regulates the Epidermal Growth Factor Receptor. Journal of Biological Chemistry, 1998, 273, 17343-17350.	3.4	17
92	Hydrogen bond analysis of Type 1 antifreeze protein in water and the ice/water interface. PhysChemComm, 2001, 4, 32-36.	0.8	17
93	Smallâ€Molecule Activation with Molybdenum(0) Complexes Supported by Mixed Imidazolâ€2‥lidene/Phosphanyl Hybrid Ligands – Electronic and Structural Consequences of Substituting a Phosphane by a Carbene Group. European Journal of Inorganic Chemistry, 2013, 2013, 3943-3955.	2.0	17
94	Hidden Flexibility of Strychnine. European Journal of Organic Chemistry, 2014, 2014, 1147-1150.	2.4	17
95	Molybdenum(0)–carbonyl complexes supported by mixed benzimidazol-2-ylidene/phosphine ligands: Influence of benzannulation on the donor properties of the NHC groups. Journal of Organometallic Chemistry, 2014, 770, 61-68.	1.8	17
96	High‥ield Lithiation of Azobenzenes by Tin–Lithium Exchange. Chemistry - A European Journal, 2015, 21, 11165-11173.	3.3	17
97	Visible-light-driven photocontrol of the Trp-cage protein fold by a diazocine cross-linker. Organic and Biomolecular Chemistry, 2020, 18, 2650-2660.	2.8	17
98	Therapeutic Potential of the Peptide Leucine Arginine As a New Nonplant Bowman–Birk-Like Serine Protease Inhibitor. Journal of Medicinal Chemistry, 2013, 56, 6732-6744.	6.4	16
99	The dynamics and binding of a Type III antifreeze protein in water and on ice. Computational and Theoretical Chemistry, 1996, 388, 65-77.	1.5	15
100	New insights into the antioxidant activity of Trolox in o/w emulsions. Food Chemistry, 2011, 124, 781-787.	8.2	15
101	Spin Switching with Triazolate-Strapped Ferrous Porphyrins. Inorganic Chemistry, 2019, 58, 5265-5272.	4.0	15
102	Backbone Structure and Dynamics of a Hemolymph Protein from the Mealworm BeetleTenebrio molitorâ€. Biochemistry, 1997, 36, 13791-13801.	2.5	14
103	Antitumor Anthraquinones from an Easter Island Sea Anemone: Animal or Bacterial Origin?. Marine Drugs, 2019, 17, 154.	4.6	14
104	Synthese mittlerer und großer Ringe, XXIII. Photochemische Umlagerung von 3,6â€Alkanooxepinâ€4,5â€dicarbonsŤreestern. Chemische Berichte, 1989, 122, 199-207.	0.2	13
105	Towards Photoswitchable Contrast Agents for Absolute 3D Temperature MR Imaging. Angewandte Chemie - International Edition, 2021, 60, 8220-8226.	13.8	13
106	Effect of ensiling duration on the fate of deoxynivalenol, zearalenone and their derivatives in maize silage. Mycotoxin Research, 2020, 36, 127-136.	2.3	12
107	NMR structural studies on antifreeze proteins. Biochemistry and Cell Biology, 1998, 76, 284-293.	2.0	12
108	N-type calcium channel/syntaxin/snap-25 complex probed by antibodies to II–III intracellular loop of the α1B subunit. Neuroscience, 1999, 90, 665-676.	2.3	11

#	Article	IF	CITATIONS
109	Activity of a Two-Domain Antifreeze Protein Is Not Dependent on Linker Sequence. Biophysical Journal, 2007, 92, 541-546.	O.5	11
110	Partitioning of nitroxides in dispersed systems investigated by ultrafiltration, EPR and NMR spectroscopy. Journal of Colloid and Interface Science, 2015, 452, 15-23.	9.4	11
111	Synthese mittlerer und großer Ringe, XXV: Synthese funktionalisierter <i>trans</i> â€Hydrindane mit angularer αâ€Ketoestergruppe. Chemische Berichte, 1989, 122, 1969-1975.	0.2	10
112	Identification of structural traits that increase the antimicrobial activity of a chimeric peptide of human β-defensins 2 and 3. Biochemical and Biophysical Research Communications, 2012, 427, 207-211.	2.1	10
113	Two thiazolylindoles and a benzimidazole: Novel compounds on the designer drug market with potential cannabinoid receptor activity. Forensic Science International, 2015, 249, 133-147.	2.2	10
114	Transcarboxylase: One of Nature's Early Nanomachines. IUBMB Life, 2004, 56, 575-583.	3.4	9
115	Two Domains of RD3 Antifreeze Protein Diffuse Independently. Biochemistry, 2008, 47, 5935-5941.	2.5	9
116	Efficient reversible photoisomerisation with large solvodynamic size-switching of a main chain poly(azobenzene- <i>alt</i> -trisiloxane). Journal of Materials Chemistry C, 2020, 8, 1835-1845.	5.5	9
117	OnlyParahydrogen SpectrosopY (OPSY) pulse sequences – One does not fit all. Journal of Magnetic Resonance, 2018, 297, 86-95.	2.1	8
118	Longâ€Ðistance Rate Acceleration by Bulk Gold. Angewandte Chemie, 2019, 131, 6646-6650.	2.0	8
119	A new stereoselective approach to the trans-perhydroazulene skeleton. Tetrahedron: Asymmetry, 1993, 4, 281-284.	1.8	7
120	The solution structure of the kallikrein-related peptidases inhibitor SPINK6. Biochemical and Biophysical Research Communications, 2016, 471, 103-108.	2.1	7
121	Überbrückte verzweigte desoxy-methylfuranoside aus 3,6-hexanooxepinen. Tetrahedron Letters, 1992, 33, 8023-8026.	1.4	6
122	Photodimerisation of glycothymidines in solution and in micelles. Chemical Communications, 2011, 47, 9399.	4.1	6
123	In vitro singlet state and zero-quantum encoded magnetic resonance spectroscopy: Illustration with N-acetyl-aspartate. PLoS ONE, 2020, 15, e0239982.	2.5	6
124	Aqueous and Micelle-bound Structural Characterization of the Epidermal Growth Factor Receptor Juxtamembrane Domain Containing Basolateral Sorting Motifs. Journal of Biomolecular Structure and Dynamics, 2004, 21, 813-826.	3.5	5
125	Bioactive Abietane-Type Diterpenoid Glycosides from Leaves of Clerodendrum infortunatum (Lamiaceae). Molecules, 2021, 26, 4121.	3.8	5
126	Sesquiterpene lactones from Sonchus palustris L. (Asteraceae, Cichorieae). Phytochemistry, 2020, 170, 112196.	2.9	4

#	Article	IF	CITATIONS
127	Catalytic Hydrogenation of Trivinyl Orthoacetate: Mechanisms Elucidated by Parahydrogen Induced Polarization. ChemPhysChem, 2021, 22, 370-377.	2.1	4
128	Cytotoxic constituents and a new hydroxycinnamic acid derivative from Leontodon saxatilis (Asteraceae, Cichorieae). RSC Advances, 2021, 11, 10489-10496.	3.6	4
129	The Structure of Fish Antifreeze Proteins. Molecular Aspects of Fish and Marine Biology, 2002, , 109-138.	0.2	4
130	Assignment of the helical structure in neuropeptide Y by HPLC studies of methionine replacement analogues and 1H-NMR spectroscopy. Biopolymers, 1998, 39, 207-219.	2.4	3
131	Formation of solid bituminous matter in pegmatites: Constraints from experimentally formed organic matter on microporous silicate minerals. Chemie Der Erde, 2014, 74, 343-351.	2.0	3
132	Evaluation of NMR Based Structure Determination of Flexible Peptides: Application to Desmopressin. , 1993, , 569-575.		3
133	Insights into the leaves of Ceriscoides campanulata: Natural proanthocyanidins alleviate diabetes, inflammation, and esophageal squamous cell cancer via in vitro and in silico models. Fìtoterapìâ, 2022, 158, 105164.	2.2	3
134	Les protéines antigel. Biofutur, 1998, 1998, 52.	0.0	2
135	Characterization of the Carboxylate Delivery Module of Transcarboxylase:Â Following Spontaneous Decarboxylation of the 1.3S-CO2-Subunit by NMR and FTIR Spectroscopiesâ€. Biochemistry, 2002, 41, 2191-2197.	2.5	2
136	Thermodynamic and kinetic stabilization of divanadate in the monovanadate/divanadate equilibrium using a Zn-cyclene derivative: Towards a simple ATP synthase model. Beilstein Journal of Organic Chemistry, 2012, 8, 81-89.	2.2	2
137	Ultrafast electronic deactivation dynamics of the inosine dimer – a model case for H-bonded purine bases. Photochemical and Photobiological Sciences, 2013, 12, 1466.	2.9	2
138	Solution structure and functional studies of the highly potent equine antimicrobial peptide DEFA1. Biochemical and Biophysical Research Communications, 2015, 459, 668-672.	2.1	2
139	Resolving the excited state relaxation dynamics of guanosine monomers and hydrogen-bonded homodimers in chloroform solution. Chemical Physics, 2018, 515, 480-492.	1.9	2
140	Designed Trp-Cage Proteins with Antimicrobial Activity and Enhanced Stability. Biochemistry, 2021, 60, 3187-3199.	2.5	2
141	Conjugated oligomers with alternating heterocycles from a single monomer: synthesis and demonstration of electroluminescence. Organic Chemistry Frontiers, 2019, 6, 3636-3643.	4.5	1
142	Towards Photoswitchable Contrast Agents for Absolute 3D Temperature MR Imaging. Angewandte Chemie, 2021, 133, 8301-8307.	2.0	1
143	Metalâ€Dependent and Selective Crystallization of CAUâ€10 and MILâ€53 Frameworks through Linker Nitration. Chemistry - A European Journal, 2021, 27, 7696-7703.	3.3	0

A new paradigm for fish antifreeze protein binding to ice. , 2002, , 747-748.

#	Article	IF	CITATIONS
145	A Structure for Little Orphan Diacylglycerol Kinase. FASEB Journal, 2007, 21, A148.	0.5	0
146	NMR based structure and enzymatic insight into diacylglycerol kinase, an alphaâ€helical membrane protein. FASEB Journal, 2009, 23, LB223.	0.5	0
147	On the Use of NMR in Complex Biological Systems: NMR Studies of Calcium Sensitive Interactions amongst Muscle Proteins. , 1996, , 275-284.		0