

Christopher Dennison

List of Publications by Year in descending order

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times ranked

2755
citing authors

#	ARTICLE	IF	CITATIONS
1	Cytosolic Copper Binding by a Bacterial Storage Protein and Interplay with Copper Efflux. <i>International Journal of Molecular Sciences</i> , 2019, 20, 4144.	4.1	8
2	The Coordination Chemistry of Copper Uptake and Storage for Methane Oxidation. <i>Chemistry - A European Journal</i> , 2019, 25, 74-86.	3.3	10
3	Bacterial copper storage proteins. <i>Journal of Biological Chemistry</i> , 2018, 293, 4616-4627.	3.4	48
4	Insight into Metal Removal from Peptides that Sequester Copper for Methane Oxidation. <i>Chemistry - A European Journal</i> , 2018, 24, 4515-4518.	3.3	16
5	Visualizing Biological Copper Storage: The Importance of Thiolate-Coordinated Tetranuclear Clusters. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 8697-8700.	13.8	16
6	Visualizing Biological Copper Storage: The Importance of Thiolate-Coordinated Tetranuclear Clusters. <i>Angewandte Chemie</i> , 2017, 129, 8823-8826.	2.0	3
7	Bacterial cytosolic proteins with a high capacity for Cu(I) that protect against copper toxicity. <i>Scientific Reports</i> , 2016, 6, 39065.	3.3	52
8	Methanobactin and the Link between Copper and Bacterial Methane Oxidation. <i>Microbiology and Molecular Biology Reviews</i> , 2016, 80, 387-409.	6.6	118
9	A four-helix bundle stores copper for methane oxidation. <i>Nature</i> , 2015, 525, 140-143.	27.8	83
10	The importance of Zn binding by the human copper metallochaperone for Cu,Zn-superoxide dismutase. <i>RSC Advances</i> , 2014, 4, 22542-22544.	3.6	5
11	How the Dynamics of the Metal-Binding Loop Region Controls the Acid Transition in Cupredoxins. <i>Biochemistry</i> , 2013, 52, 7397-7404.	2.5	5
12	Investigating the Role of Zinc and Copper Binding Motifs of Trafficking Sites in the Cyanobacterium <i>Synechocystis</i> PCC 6803. <i>Biochemistry</i> , 2013, 52, 6816-6823.	2.5	5
13	Crosstalk between Cu(i) and Zn(ii) homeostasis via Atx1 and cognate domains. <i>Chemical Communications</i> , 2013, 49, 8000.	4.1	16
14	The influence of protein folding on the copper affinities of trafficking and target sites. <i>Dalton Transactions</i> , 2013, 42, 3233-3239.	3.3	9
15	Refined Distances Between Paramagnetic Centers of a Multi-Copper Nitrite Reductase Determined by Pulsed EPR (DEER) Spectroscopy. <i>Angewandte Chemie - International Edition</i> , 2013, 52, 1990-1993.	13.8	27
16	Variations in methanobactin structure influences copper utilization by methane-oxidizing bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 8400-8404.	7.1	81
17	The influence of active site loop mutations on the thermal stability of azurin from <i>Pseudomonas aeruginosa</i> . <i>Archives of Biochemistry and Biophysics</i> , 2012, 521, 18-23.	3.0	4
18	Cu(I) Affinities of the Domain 1 and 3 Sites in the Human Metallochaperone for Cu,Zn-Superoxide Dismutase. <i>Biochemistry</i> , 2012, 51, 1439-1448.	2.5	34

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19	Understanding the Mechanism of Short-Range Electron Transfer Using an Immobilized Cupredoxin. <i>Journal of the American Chemical Society</i> , 2012, 134, 11848-11851.	13.7	34
20	Copper-Binding Properties and Structures of Methanobactins from <i>Methylosinus trichosporium</i> OB3b. <i>Inorganic Chemistry</i> , 2011, 50, 1378-1391.	4.0	76
21	Metal-Binding Loop Length Is a Determinant of the p <i>K_a</i> of a Histidine Ligand at a Type 1 Copper Site. <i>Inorganic Chemistry</i> , 2011, 50, 482-488.	4.0	13
22	Copper Trafficking Mechanism of CXXC-Containing Domains: Insight from the pH-Dependence of Their Cu(I) Affinities. <i>Journal of the American Chemical Society</i> , 2011, 133, 2983-2988.	13.7	96
23	Fluorescence Lifetime Analysis of Nitrite Reductase from <i>Alcaligenes xylosoxidans</i> at the Single-Molecule Level Reveals the Enzyme Mechanism. <i>Chemistry - A European Journal</i> , 2011, 17, 12015-12019.	3.3	22
24	Redox cycling and kinetic analysis of single molecules of solution-phase nitrite reductase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 17269-17274.	7.1	53
25	Thermodynamics of copper and zinc distribution in the cyanobacterium <i>Synechocystis</i> PCC 6803. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 13007-13012.	7.1	51
26	Visualizing the Metal-Binding Versatility of Copper Trafficking Sites. <i>Biochemistry</i> , 2010, 49, 7798-7810.	2.5	27
27	Factors Affecting the Electron Transfer Properties of an Immobilized Cupredoxin. <i>Journal of Physical Chemistry C</i> , 2010, 114, 22322-22329.	3.1	19
28	Can soaked-in scavengers protect metalloprotein active sites from reduction during data collection?. <i>Journal of Synchrotron Radiation</i> , 2009, 16, 191-204.	2.4	61
29	Active site loop dictates the thermodynamics of reduction and ligand protonation in cupredoxins. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2009, 1794, 995-1000.	2.3	8
30	Metal-binding loop length and not sequence dictates structure. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 5616-5621.	7.1	22
31	The Importance of the Long Type-1 Copper-Binding Loop of Nitrite Reductase for Structure and Function. <i>Chemistry - A European Journal</i> , 2008, 14, 5820-5828.	3.3	7
32	Thermal unfolding studies of a phycocyanin. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2008, 1784, 1997-2003.	2.3	9
33	Protein-folding location can regulate manganese-binding versus copper- or zinc-binding. <i>Nature</i> , 2008, 455, 1138-1142.	27.8	281
34	π-Interaction Tuning of the Active Site Properties of Metalloproteins. <i>Journal of the American Chemical Society</i> , 2008, 130, 15420-15428.	13.7	35
35	The role of ligand-containing loops at copper sites in proteins. <i>Natural Product Reports</i> , 2008, 25, 15-24.	10.3	24
36	Regulation of Protein Function: Crystal Packing Interfaces and Conformational Dimerization. <i>Biochemistry</i> , 2008, 47, 6583-6589.	2.5	20

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37	FutA2 Is a Ferric Binding Protein from <i>Synechocystis</i> PCC 6803. <i>Journal of Biological Chemistry</i> , 2008, 283, 12520-12527.	3.4	56
38	NMR Structural Analysis of Cadmium Sensing by Winged Helix Repressor CmtR. <i>Journal of Biological Chemistry</i> , 2007, 282, 30181-30188.	3.4	41
39	A Periplasmic Iron-binding Protein Contributes toward Inward Copper Supply. <i>Journal of Biological Chemistry</i> , 2007, 282, 3837-3846.	3.4	46
40	Engineering Copper Sites in Proteins: Δ Loops Confer Native Structures and Properties to Chimeric Cupredoxins. <i>Journal of the American Chemical Society</i> , 2007, 129, 709-718.	13.7	45
41	Influence of Loop Shortening on the Metal Binding Site of Cupredoxin Pseudoazurin. <i>Biochemistry</i> , 2007, 46, 9981-9991.	2.5	18
42	Transient Protein Interactions between Cytochrome <i>c</i> and the Phycocyanin Uromylin. <i>ChemBioChem</i> , 2007, 8, 732-735.	2.6	2
43	Thermodynamics of the alkaline transition in phycocyanins. <i>Journal of Biological Inorganic Chemistry</i> , 2007, 12, 895-900.	2.6	7
44	The Role of Hydrogen Bonding at the Active Site of a Cupredoxin: The Phe114Pro Azurin Variant. <i>Biochemistry</i> , 2006, 45, 8812-8822.	2.5	78
45	Ultrafast Pump-Probe Study of Excited-State Charge-Transfer Dynamics in Uromylin from Horseradish Root. <i>Journal of Physical Chemistry B</i> , 2006, 110, 17252-17259.	2.6	26
46	Active Site Comparison of Coll Blue and Green Nitrite Reductases. <i>Chemistry - A European Journal</i> , 2006, 12, 6647-6659.	3.3	11
47	Basic requirements for a metal-binding site in a protein: The influence of loop shortening on the cupredoxin azurin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 7258-7263.	7.1	40
48	Investigating the structure and function of cupredoxins. <i>Coordination Chemistry Reviews</i> , 2005, 249, 3025-3054.	18.8	146
49	Transient Homodimer Interactions Studied Using the Electron Self-exchange Reaction. <i>Journal of Biological Chemistry</i> , 2005, 280, 19281-19288.	3.4	14
50	BACE1 Cytoplasmic Domain Interacts with the Copper Chaperone for Superoxide Dismutase-1 and Binds Copper. <i>Journal of Biological Chemistry</i> , 2005, 280, 17930-17937.	3.4	111
51	Optical Spectroscopic Investigation of the Alkaline Transition in Uromylin from Horseradish Root. <i>Biochemistry</i> , 2005, 44, 16090-16097.	2.5	14
52	Investigating the Cause of the Alkaline Transition of Phycocyanins. <i>Biochemistry</i> , 2005, 44, 3056-3064.	2.5	14
53	Crystal Structures of Oxidized and Reduced Stellacyanin from Horseradish Roots. <i>Journal of the American Chemical Society</i> , 2005, 127, 158-166.	13.7	51
54	Ligand and loop variations at type 1 copper sites: influence on structure and reactivity. <i>Dalton Transactions</i> , 2005, , 3436.	3.3	21

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55	Reduction Potential Tuning at a Type 1 Copper Site Does Not Compromise Electron Transfer Reactivity. <i>Journal of the American Chemical Society</i> , 2005, 127, 16453-16459.	13.7	27
56	Loop-Contraction Mutagenesis of Type 1 Copper Sites. <i>Journal of the American Chemical Society</i> , 2004, 126, 15711-15719.	13.7	35
57	Characterization of Arabidopsis thaliana stellacyanin: A comparison with umecyanin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004, 55, 426-435.	2.6	11
58	An Axial Met Ligand at a Type 1 Copper Site is Preferable for Fast Electron Transfer. <i>ChemBioChem</i> , 2004, 5, 1579-1581.	2.6	15
59	Paramagnetic ¹ H NMR Spectrum of the Cobalt(II) Derivative of Spinach Plastocyanin. <i>Inorganic Chemistry</i> , 2004, 43, 1502-1510.	4.0	13
60	The Active-Site Structure of Umecyanin, the Stellacyanin from Horseradish Roots. <i>Journal of the American Chemical Society</i> , 2004, 126, 2481-2489.	13.7	23
61	Pseudospecificity of the Acidic Patch of Plastocyanin for the Interaction with Cytochrome f. <i>Journal of the American Chemical Society</i> , 2004, 126, 3028-3029.	13.7	14
62	The parsley plastocyanin-turnip cytochrome f complex: a structurally distorted but kinetically functional acidic patch. <i>Biochemical Journal</i> , 2004, 378, 45-51.	3.7	24
63	Loop-Contraction Mutagenesis of a Type 1 Copper Site. <i>Journal of the American Chemical Society</i> , 2003, 125, 4974-4975.	13.7	20
64	Introduction of a π - π Interaction at the Active Site of a Cupredoxin: Characterization of the Met16Phe Pseudoazurin Mutant. <i>Biochemistry</i> , 2003, 42, 6853-6862.	2.5	36
65	Active-Site Structure and Electron-Transfer Reactivity of Plastocyanins. <i>Journal of the American Chemical Society</i> , 2003, 125, 2101-2112.	13.7	60
66	Alkaline transition of phytocyanins: a comparison of stellacyanin and umecyanin. <i>Biochemical Journal</i> , 2003, 371, 377-383.	3.7	19
67	Paramagnetic ¹ H NMR Spectrum of Nickel(II) Pseudoazurin: Investigation of the Active Site Structure and the Acid and Alkaline Transitions. <i>Inorganic Chemistry</i> , 2002, 41, 6662-6672.	4.0	13
68	Effect of Histidine 6 Protonation on the Active Site Structure and Electron-Transfer Capabilities of Pseudoazurin from <i>Achromobacter cycloclastes</i> . <i>Biochemistry</i> , 2002, 41, 120-130.	2.5	31
69	Unusual Properties of Plastocyanin from the Fern <i>Dryopteris crassirhizoma</i> . <i>Biochemistry</i> , 2002, 41, 552-560.	2.5	33
70	UV resonance Raman and NMR spectroscopic studies on the pH dependent metal ion release from pseudoazurin. <i>Inorganica Chimica Acta</i> , 2002, 339, 383-392.	2.4	2
71	Effect of pH on the Self-Exchange Reactivity of the Plant Plastocyanin from Parsley. <i>Inorganic Chemistry</i> , 2001, 40, 354-360.	4.0	24
72	Investigations of the Alkaline and Acid Transitions of Umecyanin, a Stellacyanin from Horseradish Roots. <i>Biochemistry</i> , 2001, 40, 3158-3166.	2.5	35

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73	Paramagnetic NMR studies of copper-containing nitrite reductases. <i>Chemical Communications</i> , 2000, , 751-752.	4.1	4
74	Loop-Directed Mutagenesis of the Blue Copper Protein Amicyanin from <i>Paracoccus versutus</i> and Its Effect on the Structure and the Activity of the Type-1 Copper Site. <i>Journal of the American Chemical Society</i> , 2000, 122, 204-211.	13.7	83
75	The Met99Gln Mutant of Amicyanin from <i>Paracoccus versutus</i> . <i>Biochemistry</i> , 2000, 39, 9551-9560.	2.5	46
76	Alkaline Transition of Pseudoazurin from <i>Achromobacter cycloclastes</i> Studied by Paramagnetic NMR and Its Effect on Electron Transfer. <i>Inorganic Chemistry</i> , 1999, 38, 1491-1497.	4.0	24
77	Electron Transfer in Ruthenium-Modified Plastocyanin. <i>Journal of the American Chemical Society</i> , 1998, 120, 7551-7556.	13.7	72
78	¹ H NMR Study of the Paramagnetic Active Site of the CuA Variant of Amicyanin. <i>Biochemistry</i> , 1997, 36, 3262-3269.	2.5	47
79	¹ H NMR studies of the paramagnetic CuA center of cytochrome oxidase. <i>FEBS Letters</i> , 1996, 394, 340-344.	2.8	33
80	Analysis of the Paramagnetic Copper(II) Site of Amicyanin by ¹ H NMR Spectroscopy. <i>Biochemistry</i> , 1996, 35, 3085-3092.	2.5	77
81	Loop-Directed Mutagenesis Converts Amicyanin from <i>Thiobacillus versutus</i> into a Novel Blue Copper Protein. <i>Journal of the American Chemical Society</i> , 1996, 118, 7406-7407.	13.7	57
82	Electron Transfer Properties and Active Site Structure of the Type 1 (Blue) Copper Protein Umecyanin. <i>Chemistry - A European Journal</i> , 1996, 2, 104-109.	3.3	22
83	The Cu _A site of cytochrome <i>c</i> oxidase. <i>Recueil Des Travaux Chimiques Des Pays-Bas</i> , 1996, 115, 345-351.	0.0	11
84	Spectroscopic and Electrochemical Studies on Active-site Transitions of the Type 1 Copper Protein Pseudoazurin from <i>Achromobacter cycloclastes</i> . <i>Journal of Biological Chemistry</i> , 1995, 270, 25733-25738.	3.4	79
85	Engineered Cupredoxins and Bacterial Cytochrome <i>c</i> Oxidases Have Similar CuA Sites: Evidence from Resonance Raman Spectroscopy. <i>Journal of the American Chemical Society</i> , 1995, 117, 10759-10760.	13.7	34
86	Determination of the Self-Exchange Rate Constant for Rusticyanin from <i>Thiobacillus ferrooxidans</i> and a Comparison with Values for Other Type 1 Blue Copper Proteins. <i>Inorganic Chemistry</i> , 1995, 34, 5370-5374.	4.0	47
87	Introduction of a CuA site into the blue copper protein amicyanin from <i>Thiobacillus versutus</i> . <i>FEBS Letters</i> , 1995, 365, 92-94.	2.8	90
88	Type 1 blue copper protein amicyanin from <i>Thiobacillus versutus</i> : line-broadening effects of chromium(III) complexes and related studies. <i>Journal of the Chemical Society Dalton Transactions</i> , 1995, , 3395.	1.1	5
89	Heterogeneity of the covalent structure of the blue copper protein umecyanin from horseradish roots. <i>Protein Science</i> , 1995, 4, 209-227.	7.6	30
90	Reversible active site protonation and electron-transfer properties of <i>Achromobacter cycloclastes</i> pseudoazurin: comparisons with other type 1 copper proteins. <i>Journal of the Chemical Society Chemical Communications</i> , 1994, , 581.	2.0	18

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91	Redox reactivity of the type 1 (blue) copper protein amicyanin from <i>Thiobacillus versutus</i> with inorganic complexes. <i>Journal of the Chemical Society Dalton Transactions</i> , 1994, , 3017.	1.1	6
92	Proton nuclear magnetic resonance assignments and the electron self-exchange rate constant for pseudoazurin from <i>Achromobacter cycloclastes</i> . <i>Journal of the Chemical Society Dalton Transactions</i> , 1994, , 437.	1.1	9
93	Reactivity of Pseudoazurin from <i>Achromobacter cycloclastes</i> with Inorganic Redox Partners and Related NMR and Electrochemical Studies. <i>Inorganic Chemistry</i> , 1994, 33, 3299-3305.	4.0	50
94	Reactions of five spinach plastocyanin PCu(I) mutants with $[Fe(CN)_6]^{3-}$ and $[Co(phen)_3]^{3+}$ (phen = 2,2,6,6-tetramethylpiperidine-1-oxyl). <i>Journal of the Chemical Society Dalton Transactions</i> , 1994, , 2289-2296.	1.1	7
95	Determination of the self-exchange rate constant for plastocyanin from <i>Anabaena variabilis</i> by nuclear magnetic resonance line broadening. <i>Journal of the Chemical Society Dalton Transactions</i> , 1993, , 1959.	1.1	13