

Paul A Lindahl

List of Publications by Year in descending order

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76196

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#	ARTICLE	IF	CITATIONS
1	Ni-Zn-[Fe ₄ S ₄] and Ni-Ni-[Fe ₄ S ₄] clusters in closed and open $\hat{\pm}$ subunits of acetyl-CoA synthase/carbon monoxide dehydrogenase. <i>Nature Structural and Molecular Biology</i> , 2003, 10, 271-279.	3.6	418
2	The Ni-Containing Carbon Monoxide Dehydrogenase Family:Â Light at the End of the Tunnel?â€. <i>Biochemistry</i> , 2002, 41, 2097-2105.	1.2	197
3	Nature of the C-Cluster in Ni-Containing Carbon Monoxide Dehydrogenases. <i>Journal of the American Chemical Society</i> , 1996, 118, 830-845.	6.6	131
4	LdpA: a component of the circadian clock senses redox state of the cell. <i>EMBO Journal</i> , 2005, 24, 1202-1210.	3.5	119
5	Methylation of Carbon Monoxide Dehydrogenase from <i>Clostridium thermoaceticum</i> and Mechanism of Acetyl Coenzyme A Synthesis. <i>Journal of the American Chemical Society</i> , 1997, 119, 3959-3970.	6.6	114
6	Effects of sulfur site modification on the redox potentials of derivatives of [N,N'-bis(2-mercaptoethyl)-1,5-diazacyclooctanato]nickel(II). <i>Journal of the American Chemical Society</i> , 1993, 115, 4665-4674.	6.6	110
7	Nickel and iron EXAFS of F420-reducing hydrogenase from <i>Methanobacterium thermoautotrophicum</i> . <i>Journal of the American Chemical Society</i> , 1984, 106, 3062-3064.	6.6	98
8	Acetyl-coenzyme A synthase: the case for a NipO-based mechanism of catalysis. <i>Journal of Biological Inorganic Chemistry</i> , 2004, 9, 516-524.	1.1	97
9	Evidence of a Molecular Tunnel Connecting the Active Sites for CO ₂ Reduction and Acetyl-CoA Synthesis in Acetyl-CoA Synthase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 1999, 121, 9221-9222.	6.6	95
10	Metalâ€metal bonds in biology. <i>Journal of Inorganic Biochemistry</i> , 2012, 106, 172-178.	1.5	93
11	MÃssbauer and EPR Study of the Ni-Activated $\hat{\pm}$ -Subunit of Carbon Monoxide Dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 1997, 119, 8301-8312.	6.6	91
12	A Multinuclear ENDOR Study of the C-Cluster in CO Dehydrogenase from <i>Clostridium thermoaceticum</i> :â€ Evidence for H ₂ O and Histidine Coordination to the [Fe ₄ S ₄] Center. <i>Journal of the American Chemical Society</i> , 1998, 120, 8767-8776.	6.6	91
13	Stoichiometric reductive titrations of <i>Desulfovibrio gigas</i> hydrogenase. <i>Journal of the American Chemical Society</i> , 1995, 117, 2565-2572.	6.6	82
14	Biophysical Characterization of the Iron in Mitochondria from Atm1p-Depleted <i>Saccharomyces cerevisiae</i> . <i>Biochemistry</i> , 2009, 48, 9556-9568.	1.2	80
15	Inactivation of Acetyl-CoA Synthase/Carbon Monoxide Dehydrogenase by Copper. <i>Journal of the American Chemical Society</i> , 2003, 125, 9316-9317.	6.6	75
16	2,4,6-Trinitrotoluene Reduction by Carbon Monoxide Dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Applied and Environmental Microbiology</i> , 2000, 66, 1474-1478.	1.4	72
17	Spectroscopic, Redox, and Structural Characterization of the Ni-Labile and Nonlabile Forms of the Acetyl-CoA Synthase Active Site of Carbon Monoxide Dehydrogenase. <i>Journal of the American Chemical Society</i> , 1998, 120, 7502-7510.	6.6	67
18	Mixed-Valence Nickelâ€Iron Dithiolate Models of the [NiFe]-Hydrogenase Active Site. <i>Inorganic Chemistry</i> , 2012, 51, 2338-2348.	1.9	67

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19	EPR and Mössbauer Spectroscopy of Intact Mitochondria Isolated from Yah1p-Depleted <i>Saccharomyces cerevisiae</i> . <i>Biochemistry</i> , 2008, 47, 9888-9899.	1.2	64
20	Evidence for a Proton Transfer Network and a Required Persulfide-Bond-Forming Cysteine Residue in Ni-Containing Carbon Monoxide Dehydrogenases. <i>Biochemistry</i> , 2004, 43, 5728-5734.	1.2	62
21	Heterogeneous nickel-iron environments in carbon monoxide dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 1993, 115, 5522-5526.	6.6	61
22	Organization of Clusters and Internal Electron Pathways in CO Dehydrogenase from <i>Clostridium thermoaceticum</i> : Relevance to the Mechanism of Catalysis and Cyanide Inhibition. <i>Biochemistry</i> , 1994, 33, 8702-8711.	1.2	61
23	Spectroscopic States of the CO Oxidation/CO ₂ Reduction Active Site of Carbon Monoxide Dehydrogenase and Mechanistic Implications. <i>Biochemistry</i> , 1996, 35, 8371-8380.	1.2	60
24	Stopped-Flow Kinetics of Methyl Group Transfer between the Corrinoid-Iron-Sulfur Protein and Acetyl-Coenzyme A Synthase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 2002, 124, 6277-6284.	6.6	60
25	The evolution of acetyl-CoA synthase. , 2001, 31, 403-434.		59
26	Structures and Energetics of Models for the Active Site of Acetyl-Coenzyme A Synthase: A Role of Distal and Proximal Metals in Catalysis. <i>Journal of the American Chemical Society</i> , 2004, 126, 3410-3411.	6.6	59
27	Iron EXAFS of the iron-molybdenum cofactor of nitrogenase. <i>Journal of the American Chemical Society</i> , 1982, 104, 4703-4705.	6.6	58
28	EXAFS, EPR, and Electronic Absorption Spectroscopic Study of the α . Metallo Subunit of CO Dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 1995, 117, 7065-7070.	6.6	58
29	Analysis of Oxidative Titrations of <i>Desulfovibrio gigas</i> Hydrogenase; Implications for the Catalytic Mechanism. <i>Biochemistry</i> , 1994, 33, 14339-14350.	1.2	57
30	Discovery of a labile nickel ion required for CO/acetyl-CoA exchange activity in the NiFe complex of carbon monoxide dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 1992, 114, 9718-9719.	6.6	56
31	The Tunnel of Acetyl-Coenzyme A Synthase/Carbon Monoxide Dehydrogenase Regulates Delivery of CO to the Active Site. <i>Journal of the American Chemical Society</i> , 2005, 127, 5833-5839.	6.6	56
32	Biophysical Characterization of Iron in Mitochondria Isolated from Respiring and Fermenting Yeast. <i>Biochemistry</i> , 2010, 49, 5436-5444.	1.2	56
33	Evaluation of Multivalent Dendrimers Based on Melamine: Kinetics of Thiol-Disulfide Exchange Depends on the Structure of the Dendrimer. <i>Journal of the American Chemical Society</i> , 2003, 125, 5086-5094.	6.6	54
34	Stability of the Ni-C State and Oxidative Titrations of <i>Desulfovibrio gigas</i> Hydrogenase Monitored by EPR and Electronic Absorption Spectroscopies. <i>Journal of the American Chemical Society</i> , 1994, 116, 3442-3448.	6.6	52
35	Function and carbon monoxide binding properties of the nickel-iron complex in carbon monoxide dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Biochemistry</i> , 1992, 31, 12870-12875.	1.2	51
36	Reduction and Methyl Transfer Kinetics of the β Subunit from Acetyl Coenzyme A Synthase. <i>Journal of the American Chemical Society</i> , 2003, 125, 318-319.	6.6	51

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37	Identification of a cyanide binding site in CO dehydrogenase from <i>Clostridium thermoaceticum</i> using EPR and ENDOR spectroscopies. <i>Journal of the American Chemical Society</i> , 1993, 115, 12204-12205.	6.6	50
38	Iron Content of <i>Saccharomyces cerevisiae</i> Cells Grown under Iron-Deficient and Iron-Overload Conditions. <i>Biochemistry</i> , 2013, 52, 105-114.	1.2	50
39	Catalytic Coupling of the Active Sites in Acetyl-CoA Synthase, a Bifunctional CO-Channeling Enzyme. <i>Biochemistry</i> , 2001, 40, 13262-13267.	1.2	48
40	Kinetics of CO Insertion and Acetyl Group Transfer Steps, and a Model of the Acetyl-CoA Synthase Catalytic Mechanism. <i>Journal of the American Chemical Society</i> , 2006, 128, 12331-12338.	6.6	47
41	Labile Low-Molecular-Mass Metal Complexes in Mitochondria: Trials and Tribulations of a Burgeoning Field. <i>Biochemistry</i> , 2016, 55, 4140-4153.	1.2	44
42	Effect of Sodium Sulfide on Ni-Containing Carbon Monoxide Dehydrogenases. <i>Journal of the American Chemical Society</i> , 2004, 126, 9094-9100.	6.6	43
43	Biophysical Investigation of the Ironome of Human Jurkat Cells and Mitochondria. <i>Biochemistry</i> , 2012, 51, 5276-5284.	1.2	43
44	Antiferromagnetic coupling in the binuclear metal cluster of manganese-substituted phosphotriesterase. <i>Journal of the American Chemical Society</i> , 1993, 115, 12173-12174.	6.6	42
45	Catalytic Mechanism and Three-Dimensional Structure of Adenine Deaminase. <i>Biochemistry</i> , 2011, 50, 1917-1927.	1.2	42
46	Kinetic Modeling of the Assembly, Dynamic Steady State, and Contraction of the FtsZ Ring in Prokaryotic Cytokinesis. <i>PLoS Computational Biology</i> , 2008, 4, e1000102.	1.5	41
47	A Nonheme High-Spin Ferrous Pool in Mitochondria Isolated from Fermenting <i>Saccharomyces cerevisiae</i> . <i>Biochemistry</i> , 2010, 49, 4227-4234.	1.2	41
48	Assembly of an Exchange-Coupled [Ni:Fe ₄ S ₄] Cluster in the Î± Metallosubunit of Carbon Monoxide Dehydrogenase from <i>Clostridium thermoaceticum</i> with Spectroscopic Properties and CO-Binding Ability Mimicking Those of the Acetyl-CoA Synthase Active Site. <i>Journal of the American Chemical Society</i> , 1996, 118, 483-484.	6.6	40
49	Mössbauer and EPR Study of Iron in Vacuoles from Fermenting <i>Saccharomyces cerevisiae</i> . <i>Biochemistry</i> , 2011, 50, 10275-10283.	1.2	40
50	Mössbauer and EPR Study of Recombinant Acetyl-CoA Synthase from <i>Moorella thermoacetica</i> . <i>Biochemistry</i> , 2006, 45, 8674-8685.	1.2	38
51	Carbon Monoxide Dehydrogenase from <i>Clostridium thermoaceticum</i> : A Quaternary Structure, Stoichiometry of Its SDS-Induced Dissociation, and Characterization of the Faster-Migrating Form. <i>Biochemistry</i> , 1996, 35, 1965-1971.	1.2	37
52	Isolated <i>Saccharomyces cerevisiae</i> vacuoles contain low-molecular-mass transition-metal polyphosphate complexes. <i>Metallomics</i> , 2019, 11, 1298-1309.	1.0	37
53	COA6 Is Structurally Tuned to Function as a Thiol-Disulfide Oxidoreductase in Copper Delivery to Mitochondrial Cytochrome c Oxidase. <i>Cell Reports</i> , 2019, 29, 4114-4126.e5.	2.9	37
54	Kinetic Mechanism of Acetyl-CoA Synthase: A Steady-State Synthesis at Variable CO/CO ₂ Pressures. <i>Journal of the American Chemical Society</i> , 2001, 123, 4697-4703.	6.6	36

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55	Electron paramagnetic resonance and Mössbauer spectroscopy of intact mitochondria from respiring <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Inorganic Chemistry</i> , 2007, 12, 1029-1053.	1.1	35
56	Evidence that a respiratory shield in <i>Escherichia coli</i> protects a low-molecular-mass FeII pool from O ₂ -dependent oxidation. <i>Journal of Biological Chemistry</i> , 2019, 294, 50-62.	1.6	35
57	Spectroelectrochemical Characterization of the Metal Centers in Carbon Monoxide Dehydrogenase (CODH) and Nickel-deficient CODH from <i>Rhodospirillum rubrum</i> . <i>Journal of Biological Chemistry</i> , 1996, 271, 7973-7977.	1.6	34
58	CO/CO ₂ Potentiometric Titrations of Carbon Monoxide Dehydrogenase from <i>Clostridium thermoaceticum</i> and the Effect of CO ₂ . <i>Biochemistry</i> , 1998, 37, 10016-10026.	1.2	34
59	Identification and preliminary characterization of AcsF, a putative Ni-insertase used in the biosynthesis of acetyl-CoA synthase from <i>Clostridium thermoaceticum</i> . <i>Journal of Inorganic Biochemistry</i> , 2003, 93, 33-40.	1.5	34
60	EXAFS studies of the nitrogenase iron protein from <i>Azotobacter vinelandii</i> . <i>Inorganic Chemistry</i> , 1987, 26, 3912-3916.	1.9	32
61	Low spin quantitation of NiFeC EPR signal from carbon monoxide dehydrogenase is not due to damage incurred during protein purification. <i>BBA - Proteins and Proteomics</i> , 1993, 1161, 317-322.	2.1	32
62	Evidence for a Proposed Intermediate Redox State in the CO/CO ₂ Active Site of Acetyl-CoA Synthase (Carbon Monoxide Dehydrogenase) from <i>Clostridium thermoaceticum</i> . <i>Biochemistry</i> , 1999, 38, 15706-15711.	1.2	32
63	Function of the tunnel in acetylcoenzyme A synthase/carbon monoxide dehydrogenase. <i>Journal of Biological Inorganic Chemistry</i> , 2006, 11, 371-378.	1.1	31
64	Detection of Labile Low-Molecular-Mass Transition Metal Complexes in Mitochondria. <i>Biochemistry</i> , 2015, 54, 3442-3453.	1.2	31
65	Reactivities and biological functions of iron-sulfur clusters. <i>Journal of Cluster Science</i> , 1990, 1, 29-73.	1.7	30
66	Implications of a Carboxylate-Bound Fe Cluster Structure of Carbon Monoxide Dehydrogenase. <i>Angewandte Chemie - International Edition</i> , 2008, 47, 4054-4056.	7.2	30
67	Biophysical Investigation of the Iron in Aft1-1 ^{up} and Gal-YAH1 in <i>Saccharomyces cerevisiae</i> . <i>Biochemistry</i> , 2011, 50, 2660-2671.	1.2	30
68	Changing iron content of the mouse brain during development. <i>Metallomics</i> , 2012, 4, 761.	1.0	29
69	Redox titrations of carbon monoxide dehydrogenase from <i>Clostridium thermoaceticum</i> . <i>Biochemistry</i> , 1992, 31, 6003-6011.	1.2	28
70	Mitochondria Export Sulfur Species Required for Cytosolic tRNA Thiolation. <i>Cell Chemical Biology</i> , 2018, 25, 738-748.e3.	2.5	28
71	Carbon Monoxide Dehydrogenase from <i>Rhodospirillum rubrum</i> : Effect of Redox Potential on Catalysis. <i>Biochemistry</i> , 2004, 43, 1552-1559.	1.2	27
72	Mössbauer Evidence for an Exchange-Coupled {[Fe ₄ S ₄] ¹⁺ Ni _p ¹⁺ } A-Cluster in Isolated \pm Subunits of Acetyl-Coenzyme A Synthase/Carbon Monoxide Dehydrogenase. <i>Journal of the American Chemical Society</i> , 2008, 130, 6712-6713.	6.6	27

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73	Low-molecular-mass iron in healthy blood plasma is not predominately ferric citrate. <i>Metallomics</i> , 2018, 10, 802-817.	1.0	27
74	Structures, Interconversions, and Spectroscopy of Iron Carbonyl Clusters with an Interstitial Carbide: Localized Metal Center Reduction by Overall Cluster Oxidation. <i>Inorganic Chemistry</i> , 2017, 56, 5998-6012.	1.9	26
75	Insights into the iron-ome and manganese-ome of <i>mtm1</i> <i>Saccharomyces cerevisiae</i> mitochondria. <i>Metallomics</i> , 2013, 5, 656.	1.0	24
76	Mössbauer Spectra of Mouse Hearts Reveal Age-dependent Changes in Mitochondrial and Ferritin Iron Levels. <i>Journal of Biological Chemistry</i> , 2017, 292, 5546-5554.	1.6	24
77	The thermally induced decarboxylation mechanism of a mixed-oxidation state carboxylate-based iron metal-organic framework. <i>Chemical Communications</i> , 2019, 55, 12769-12772.	2.2	24
78	Effect of Zn on Acetyl Coenzyme A Synthase: Evidence for a Conformational Change in the β Subunit during Catalysis. <i>Journal of the American Chemical Society</i> , 2004, 126, 5954-5955.	6.6	22
79	Genetic Construction of Truncated and Chimeric Metalloproteins Derived from the β Subunit of Acetyl-CoA Synthase from <i>Clostridium thermoaceticum</i> . <i>Journal of the American Chemical Society</i> , 2002, 124, 8667-8672.	6.6	21
80	A framework for whole-cell mathematical modeling. <i>Journal of Theoretical Biology</i> , 2004, 231, 581-596.	0.8	21
81	Biophysical probes of iron metabolism in cells and organelles. <i>Current Opinion in Chemical Biology</i> , 2011, 15, 342-346.	2.8	21
82	Mitochondrial Iron-Sulfur Cluster Activity and Cytosolic Iron Regulate Iron Traffic in <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 2015, 290, 26968-26977.	1.6	21
83	Decomposition of Carbon Monoxide Dehydrogenase into α Metallosubunits and a Catalytically-Active Form Consisting Primarily of β Metallosubunits. <i>Biochemistry</i> , 1995, 34, 6037-6042.	1.2	19
84	Stoichiometric CO Reductive Titrations of Acetyl-CoA Synthase (Carbon Monoxide Dehydrogenase) from <i>Clostridium thermoaceticum</i> . <i>Biochemistry</i> , 1999, 38, 15697-15705.	1.2	19
85	Mathematical Model of a Cell Size Checkpoint. <i>PLoS Computational Biology</i> , 2010, 6, e1001036.	1.5	19
86	Chapter 15 Isolation of <i>Saccharomyces Cerevisiae</i> Mitochondria for Mössbauer, Epr, and Electronic Absorption Spectroscopic Analyses. <i>Methods in Enzymology</i> , 2009, 456, 267-285.	0.4	18
87	Speciation of iron in mouse liver during development, iron deficiency, IRP2 deletion and inflammatory hepatitis. <i>Metallomics</i> , 2015, 7, 93-101.	1.0	18
88	Recovery of <i>Saccharomyces cerevisiae</i> Cells under Iron-Sufficient Conditions and the Role of Fe ⁵⁸⁰ . <i>Biochemistry</i> , 2018, 57, 672-683.	1.2	18
89	A comprehensive mechanistic model of iron metabolism in <i>Saccharomyces cerevisiae</i> . <i>Metallomics</i> , 2019, 11, 1779-1799.	1.0	17
90	High-Spin Ferric Ions in <i>Saccharomyces cerevisiae</i> Vacuoles Are Reduced to the Ferrous State during Adenine-Precursor Detoxification. <i>Biochemistry</i> , 2014, 53, 3940-3951.	1.2	16

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91	Low-molecular-mass labile metal pools in <i>Escherichia coli</i> : advances using chromatography and mass spectrometry. <i>Journal of Biological Inorganic Chemistry</i> , 2021, 26, 479-494.	1.1	16
92	Stepwise Evolution of Nonliving to Living Chemical Systems. <i>Origins of Life and Evolution of Biospheres</i> , 2004, 34, 371-389.	0.8	15
93	Whole-cell modeling framework in which biochemical dynamics impact aspects of cellular geometry. <i>Journal of Theoretical Biology</i> , 2007, 244, 154-166.	0.8	15
94	Mathematical modeling of a minimal protocell with coordinated growth and division. <i>Journal of Theoretical Biology</i> , 2009, 260, 422-429.	0.8	15
95	Novel Domain Arrangement in the Crystal Structure of a Truncated Acetyl-CoA Synthase from <i>Moorella thermoacetica</i> . <i>Biochemistry</i> , 2009, 48, 7916-7926.	1.2	15
96	The catalase activity of diiron adenine deaminase. <i>Protein Science</i> , 2011, 20, 2080-2094.	3.1	14
97	Chromatographic detection of low-molecular-mass metal complexes in the cytosol of <i>Saccharomyces cerevisiae</i> . <i>Metallomics</i> , 2020, 12, 1094-1105.	1.0	14
98	Thermal decarboxylation for the generation of hierarchical porosity in isostructural metal-organic frameworks containing open metal sites. <i>Materials Advances</i> , 2021, 2, 5487-5493.	2.6	14
99	Acetyl-coenzyme A Synthases and Nickel-Containing Carbon Monoxide Dehydrogenases. , 2007, , 357-415.		13
100	The Lack of Synchronization between Iron Uptake and Cell Growth Leads to Iron Overload in <i>Saccharomyces cerevisiae</i> during Post-exponential Growth Modes. <i>Biochemistry</i> , 2013, 52, 9413-9425.	1.2	13
101	Mössbauer, EPR, and Modeling Study of Iron Trafficking and Regulation in <i>Yeast</i> and <i>CCC1-up Saccharomyces cerevisiae</i> . <i>Biochemistry</i> , 2014, 53, 2926-2940.	1.2	13
102	Dynamic responses of protein homeostatic regulatory mechanisms to perturbations from steady state. <i>Journal of Theoretical Biology</i> , 2003, 222, 407-423.	0.8	11
103	Mössbauer Study and Modeling of Iron Import and Trafficking in Human Jurkat Cells. <i>Biochemistry</i> , 2013, 52, 7926-7942.	1.2	11
104	Kinetics of Iron Import into Developing Mouse Organs Determined by a Pup-swapping Method*. <i>Journal of Biological Chemistry</i> , 2015, 290, 520-528.	1.6	11
105	Nickel-Dependent Oligomerization of the Alpha Subunit of Acetyl-Coenzyme A Synthase/Carbon Monoxide Dehydrogenase. <i>Biochemistry</i> , 2007, 46, 11606-11613.	1.2	10
106	A mathematical model of iron import and trafficking in wild-type and <i>Mrs3/4^Δ</i> yeast cells. <i>BMC Systems Biology</i> , 2019, 13, 23.	3.0	10
107	A Sec14-like phosphatidylinositol transfer protein paralog defines a novel class of heme-binding proteins. <i>ELife</i> , 2020, 9, .	2.8	10
108	Tunnel mutagenesis and Ni-dependent reduction and methylation of the α subunit of acetyl coenzyme A synthase/carbon monoxide dehydrogenase. <i>Journal of Biological Inorganic Chemistry</i> , 2008, 13, 771-778.	1.1	9

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109	Ferric ions accumulate in the walls of metabolically inactivating <i>Saccharomyces cerevisiae</i> cells and are reductively mobilized during reactivation. <i>Metallomics</i> , 2016, 8, 692-708.	1.0	9
110	Low-molecular-mass iron complexes in blood plasma of iron-deficient pigs do not originate directly from nutrient iron. <i>Metallomics</i> , 2019, 11, 1900-1911.	1.0	9
111	Mössbauer and LC-ICP-MS investigation of iron trafficking between vacuoles and mitochondria in <i>vma21^Δ</i> <i>Saccharomyces cerevisiae</i> . <i>Journal of Biological Chemistry</i> , 2021, 296, 100141.	1.6	8
112	Low-molecular-mass metal complexes in the mouse brain. <i>Metallomics</i> , 2013, 5, 232.	1.0	7
113	Analysis of Protein Homeostatic Regulatory Mechanisms in Perturbed Environments at Steady State. <i>Journal of Theoretical Biology</i> , 2002, 215, 151-167.	0.8	6
114	Mathematical model for positioning the FtsZ contractile ring in <i>Escherichia coli</i> . <i>Journal of Mathematical Biology</i> , 2014, 68, 911-930.	0.8	5
115	Identification of the CO oxidation site of CO dehydrogenase by EPR and ENDOR studies of the cyanide-inhibited state.. <i>Journal of Inorganic Biochemistry</i> , 1993, 51, 204.	1.5	4
116	Direct Detection of the Labile Nickel Pool in <i>Escherichia coli</i> : New Perspectives on Labile Metal Pools. <i>Journal of the American Chemical Society</i> , 2021, 143, 18571-18580.	6.6	4
117	Autocatalytic activation of acetyl-CoA synthase. <i>Journal of Biological Inorganic Chemistry</i> , 2004, 9, 316-322.	1.1	3
118	4. The utility of Mössbauer spectroscopy in eukaryotic cell biology and animal physiology. , 2014, , 49-76.		3
119	Nickel-carbon bonds in acetyl-coenzyme a synthases/carbon monoxide dehydrogenases. <i>Metal Ions in Life Sciences</i> , 2009, 6, 133-50.	2.8	3
120	Stoichiometric Redox Titrations of Complex Metalloenzymes. <i>Methods in Enzymology</i> , 2002, 354, 296-309.	0.4	2
121	The <i>Pyrococcus furiosus</i> ironome is dominated by [Fe4S4] ²⁺ clusters or thioferrate-like iron depending on the availability of elemental sulfur. <i>Journal of Biological Chemistry</i> , 2021, 296, 100710.	1.6	2
122	6. The utility of Mössbauer spectroscopy in eukaryotic cell biology and animal physiology. , 2017, , 163-190.		1
123	Cis-Divacant Octahedral Fe(II) in a Dimensionally Reduced Family of 2-(Pyridin-2-yl)pyrrolide Complexes. <i>Inorganic Chemistry</i> , 2021, 60, 15617-15626.	1.9	1
124	Yeast cells depleted of the frataxin homolog Yfh1 redistribute cellular iron: Studies using Mössbauer spectroscopy and mathematical modeling. <i>Journal of Biological Chemistry</i> , 2022, 298, 101921.	1.6	1
125	4 Nickel-Carbon Bonds in Acetyl-Coenzyme A Synthases/Carbon Monoxide Dehydrogenases. , 2015, , 133-150.		0