

Juan C Fontecilla-Camps

List of Publications by Year in descending order

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97
papers

12,033
citations

50276

46
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94
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108
all docs

108
docs citations

108
times ranked

5774
citing authors

#	ARTICLE	IF	CITATIONS
1	Nickel and the origin and early evolution of life. <i>Metallomics</i> , 2022, 14, .	2.4	5
2	The Complex Roles of Adenosine Triphosphate in Bioenergetics. <i>ChemBioChem</i> , 2022, 23, e202200064.	2.6	5
3	Quinolate Synthase: An Example of the Roles of the Second and Outer Coordination Spheres in Enzyme Catalysis. <i>Chemical Reviews</i> , 2022, , .	47.7	10
4	Primordial bioenergy sources: The two facets of adenosine triphosphate. <i>Journal of Inorganic Biochemistry</i> , 2021, 216, 111347.	3.5	12
5	Transient Formation of a Second Active Site Cavity during Quinolinic Acid Synthesis by NadA. <i>ACS Chemical Biology</i> , 2021, 16, 2423-2433.	3.4	1
6	Structural basis for the catalytic activities of the multifunctional enzyme quinolate synthase. <i>Coordination Chemistry Reviews</i> , 2020, 417, 213370.	18.8	4
7	Electron and Proton Transfers Modulate DNA Binding by the Transcription Regulator RsrR. <i>Journal of the American Chemical Society</i> , 2020, 142, 5104-5116.	13.7	11
8	Design of specific inhibitors of quinolate synthase based on [4Fe-4S] cluster coordination. <i>Chemical Communications</i> , 2019, 55, 3725-3728.	4.1	4
9	Geochemical Continuity and Catalyst/Cofactor Replacement in the Emergence and Evolution of Life. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 42-48.	13.8	30
10	Geochemische Kontinuität und Katalysator/Cofaktor-Austausch für Ursprung und Evolution des Lebens. <i>Angewandte Chemie</i> , 2019, 131, 42-48.	2.0	5
11	Crystal Structure of the Transcription Regulator RsrR Reveals a [2Fe-2S] Cluster Coordinated by Cys, Glu, and His Residues. <i>Journal of the American Chemical Society</i> , 2019, 141, 2367-2375.	13.7	18
12	Crystallographic Trapping of Reaction Intermediates in Quinolinic Acid Synthesis by NadA. <i>ACS Chemical Biology</i> , 2018, 13, 1209-1217.	3.4	9
13	Crystallographic evidence for unexpected selective tyrosine hydroxylations in an aerated achiral Ru-papain conjugate. <i>Metallomics</i> , 2018, 10, 1452-1459.	2.4	1
14	X-ray structural, functional and computational studies of the O ₂ -sensitive E. coli hydrogenase-1 C19G variant reveal an unusual [4Fe-4S] cluster. <i>Chemical Communications</i> , 2018, 54, 7175-7178.	4.1	5
15	Crystal structures of the NO sensor NsrR reveal how its iron-sulfur cluster modulates DNA binding. <i>Nature Communications</i> , 2017, 8, 15052.	12.8	59
16	12. Iron-sulfur clusters and molecular oxygen: function, adaptation, degradation, and repair. , 2017, , 359-386.		2
17	Carbon-sulfur bond-forming reaction catalysed by the radical SAM enzyme HydE. <i>Nature Chemistry</i> , 2016, 8, 491-500.	13.6	72
18	Crystal Structures of Quinolate Synthase in Complex with a Substrate Analogue, the Condensation Intermediate, and Substrate-Derived Product. <i>Journal of the American Chemical Society</i> , 2016, 138, 11802-11809.	13.7	14

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19	Fine-tuning of a radical-based reaction by radical <i>S</i> -adenosyl-L-methionine tryptophan lyase. <i>Science</i> , 2016, 351, 1320-1323.	12.6	53
20	CO and CN ^ˆ syntheses by [FeFe]-hydrogenase maturase HydG are catalytically differentiated events. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 104-109.	7.1	47
21	The crystal structure of the global anaerobic transcriptional regulator FNR explains its extremely fine-tuned monomer-dimer equilibrium. <i>Science Advances</i> , 2015, 1, e1501086.	10.3	37
22	5 Structure and Function of [NiFe]-Hydrogenases. , 2015, , 151-178.		0
23	Crystal Structure of HydG from <i>Carboxydotherrnus hydrogenoformans</i> : A Trifunctional [FeFe]-Hydrogenase Maturase. <i>ChemBioChem</i> , 2015, 16, 397-402.	2.6	41
24	Crystallographic studies of [NiFe]-hydrogenase mutants: towards consensus structures for the elusive unready oxidized states. <i>Journal of Biological Inorganic Chemistry</i> , 2015, 20, 11-22.	2.6	52
25	Tryptophan Lyase (NosL): Mechanistic Insights from Substrate Analogues and Mutagenesis. <i>Biochemistry</i> , 2015, 54, 4767-4769.	2.5	46
26	[NiFe]-hydrogenases revisited: nickel- ^ˆ carboxamido bond formation in a variant with accrued O ₂ -tolerance and a tentative re-interpretation of Ni-SI states. <i>Metallomics</i> , 2015, 7, 710-718.	2.4	19
27	IscS from <i>Archaeoglobus fulgidus</i> has no desulfurase activity but may provide a cysteine ligand for [Fe ₂ S ₂] cluster assembly. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2015, 1853, 1457-1463.	4.1	11
28	10. Iron-sulfur clusters and molecular oxygen: function, adaptation, degradation, and repair. , 2014, , 239-266.		3
29	The Stereochemical Basis of the Genetic Code and the (Mostly) Autotrophic Origin of Life. <i>Life</i> , 2014, 4, 1013-1025.	2.4	11
30	Structural Foundations for O ₂ Sensitivity and O ₂ Tolerance in [NiFe]-Hydrogenases. <i>Advances in Photosynthesis and Respiration</i> , 2014, , 23-41.	1.0	0
31	Crystal Structure of Tryptophan Lyase (NosL): Evidence for Radical Formation at the Amino Group of Tryptophan. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 11840-11844.	13.8	81
32	Artificial metalloenzymes derived from bovine β -lactoglobulin for the asymmetric transfer hydrogenation of an aryl ketone ^ˆ synthesis, characterization and catalytic activity. <i>Dalton Transactions</i> , 2014, 43, 5482-5489.	3.3	32
33	The Crystal Structure of Fe ₄ S ₄ Quinolate Synthase Unravels an Enzymatic Dehydration Mechanism That Uses Tyrosine and a Hydrolase-Type Triad. <i>Journal of the American Chemical Society</i> , 2014, 136, 5253-5256.	13.7	23
34	Crystal structure and functional studies of an unusual <i>l</i> -cysteine desulfurase from <i>Archaeoglobus fulgidus</i> . <i>Dalton Transactions</i> , 2013, 42, 3092-3099.	3.3	15
35	Electronic states of the O ₂ -tolerant [NiFe] hydrogenase proximal cluster. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E2538.	7.1	9
36	The binding mode of Ni-(L-His) ₂ in NikA revealed by X-ray crystallography. <i>Journal of Inorganic Biochemistry</i> , 2013, 121, 16-18.	3.5	41

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37	Crystal Structure of the O ₂ -Tolerant Membrane-Bound Hydrogenase 1 from <i>Escherichia coli</i> in Complex with Its Cognate Cytochrome b. <i>Structure</i> , 2013, 21, 184-190.	3.3	93
38	Principles of Sustained Enzymatic Hydrogen Oxidation in the Presence of Oxygen – The Crucial Influence of High Potential Fe–S Clusters in the Electron Relay of [NiFe]-Hydrogenases. <i>Journal of the American Chemical Society</i> , 2013, 135, 2694-2707.	13.7	91
39	X-ray snapshots of possible intermediates in the time course of synthesis and degradation of protein-bound Fe ₄ S ₄ clusters. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 7188-7192.	7.1	62
40	Structural Basis for Enantioselectivity in the Transfer Hydrogenation of a Ketone Catalyzed by an Artificial Metalloenzyme. <i>European Journal of Inorganic Chemistry</i> , 2013, 2013, 3596-3600.	2.0	23
41	The Structural Plasticity of the Proximal [4Fe3S] Cluster is Responsible for the O ₂ Tolerance of Membrane-Bound [NiFe] Hydrogenases. <i>Angewandte Chemie - International Edition</i> , 2013, 52, 2002-2006.	13.8	31
42	X-ray crystallographic and computational studies of the O ₂ -tolerant [NiFe]-hydrogenase 1 from <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 5305-5310.	7.1	194
43	Structure-Function Relationships in [FeFe]-Hydrogenase Active Site Maturation. <i>Journal of Biological Chemistry</i> , 2012, 287, 13532-13540.	3.4	72
44	(IscA–IscU) ₂ Complex Structures Provide Insights into Fe ₂ S ₂ Biogenesis and Transfer. <i>Angewandte Chemie - International Edition</i> , 2012, 51, 5439-5442.	13.8	123
45	The structure of the periplasmic nickel-binding protein NikA provides insights for artificial metalloenzyme design. <i>Journal of Biological Inorganic Chemistry</i> , 2012, 17, 817-829.	2.6	27
46	The Evolutionary Relationship Between Complex I and [NiFe]-Hydrogenase. , 2012, , 109-121.		1
47	The quest for a functional substrate access tunnel in FeFe hydrogenase. <i>Faraday Discussions</i> , 2011, 148, 385-407.	3.2	70
48	Formaldehyde – A Rapid and Reversible Inhibitor of Hydrogen Production by [FeFe]-Hydrogenases. <i>Journal of the American Chemical Society</i> , 2011, 133, 1282-1285.	13.7	30
49	Carbon Monoxide Dehydrogenase Reaction Mechanism: A Likely Case of Abnormal CO ₂ Insertion to a Ni–H Bond. <i>Inorganic Chemistry</i> , 2011, 50, 1868-1878.	4.0	75
50	Histidine 416 of the periplasmic binding protein NikA is essential for nickel uptake in <i>Escherichia coli</i> . <i>FEBS Letters</i> , 2011, 585, 711-715.	2.8	22
51	Further Characterization of the [FeFe]-Hydrogenase Maturase HydG. <i>European Journal of Inorganic Chemistry</i> , 2011, 2011, 1121-1127.	2.0	23
52	A glycol free radical as the precursor in the synthesis of carbon monoxide and cyanide by the [FeFe]-hydrogenase maturase HydG. <i>FEBS Letters</i> , 2010, 584, 4197-4202.	2.8	51
53	Maturation of [FeFe]-hydrogenases: Structures and mechanisms. <i>International Journal of Hydrogen Energy</i> , 2010, 35, 10750-10760.	7.1	24
54	Unexpected electron transfer mechanism upon AdoMet cleavage in radical SAM proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 14867-14871.	7.1	84

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55	The role of the maturase HydG in [FeFe]-hydrogenase active site synthesis and assembly. FEBS Letters, 2009, 583, 506-511.	2.8	134
56	Structure-function relationships of anaerobic gas-processing metalloenzymes. Nature, 2009, 460, 814-822.	27.8	231
57	Visible Light-Driven H ₂ Production by Hydrogenases Attached to Dye-Sensitized TiO ₂ Nanoparticles. Journal of the American Chemical Society, 2009, 131, 18457-18466.	13.7	407
58	Novel Domain Arrangement in the Crystal Structure of a Truncated Acetyl-CoA Synthase from <i>Moorella thermoacetica</i> . Biochemistry, 2009, 48, 7916-7926.	2.5	15
59	Catalytic electrochemistry of a [NiFeSe]-hydrogenase on TiO ₂ and demonstration of its suitability for visible-light driven H ₂ production. Chemical Communications, 2009, , 550-552.	4.1	160
60	Electrochemical Kinetic Investigations of the Reactions of [FeFe]-Hydrogenases with Carbon Monoxide and Oxygen: Comparing the Importance of Gas Tunnels and Active-Site Electronic/Redox Effects. Journal of the American Chemical Society, 2009, 131, 14979-14989.	13.7	167
61	Structure and Function of [NiFe]-Hydrogenases. Metal Ions in Life Sciences, 2009, 6, 151-78.	2.8	2
62	A QM/MM study of proton transport pathways in a [NiFe] hydrogenase. Proteins: Structure, Function and Bioinformatics, 2008, 73, 195-203.	2.6	58
63	The Enamine Intermediate May Not Be Universal to Thiamine Catalysis. Angewandte Chemie - International Edition, 2008, 47, 628-628.	13.8	0
64	The Difference a Se Makes? Oxygen-Tolerant Hydrogen Production by the [NiFeSe]-Hydrogenase from <i>Desulfomicrobium baculatum</i> . Journal of the American Chemical Society, 2008, 130, 13410-13416.	13.7	172
65	Structural Characterization of a Putative Endogenous Metal Chelator in the Periplasmic Nickel Transporter NikA. Biochemistry, 2008, 47, 9937-9943.	2.5	67
66	X-ray Structure of the [FeFe]-Hydrogenase Maturase HydE from <i>Thermotoga maritima</i> . Journal of Biological Chemistry, 2008, 283, 18861-18872.	3.4	119
67	Experimental approaches to kinetics of gas diffusion in hydrogenase. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 11188-11193.	7.1	150
68	Structure/Function Relationships of [NiFe]- and [FeFe]-Hydrogenases. Chemical Reviews, 2007, 107, 4273-4303.	47.7	1,234
69	The Enamine Intermediate May Not Be Universal to Thiamine Catalysis. Angewandte Chemie - International Edition, 2007, 46, 9019-9022.	13.8	4
70	Electrochemical Investigations of the Interconversions between Catalytic and Inhibited States of the [FeFe]-Hydrogenase from <i>Desulfovibrio desulfuricans</i> . Journal of the American Chemical Society, 2006, 128, 16808-16815.	13.7	78
71	Function of the tunnel in acetylcoenzyme A synthase/carbon monoxide dehydrogenase. Journal of Biological Inorganic Chemistry, 2006, 11, 371-378.	2.6	31
72	Flexibility of Thiamine Diphosphate Revealed by Kinetic Crystallographic Studies of the Reaction of Pyruvate-Ferredoxin Oxidoreductase with Pyruvate. Structure, 2006, 14, 217-224.	3.3	40

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73	Structure–function relationships of nickel–iron sites in hydrogenase and a comparison with the active sites of other nickel–iron enzymes. <i>Coordination Chemistry Reviews</i> , 2005, 249, 1609-1619.	18.8	113
74	Structural differences between the ready and unready oxidized states of [NiFe] hydrogenases. <i>Journal of Biological Inorganic Chemistry</i> , 2005, 10, 239-249.	2.6	291
75	Crystallographic and Spectroscopic Evidence for High Affinity Binding of FeEDTA(H ₂ O)-to the Periplasmic Nickel Transporter NikA. <i>Journal of the American Chemical Society</i> , 2005, 127, 10075-10082.	13.7	74
76	Electrochemical Definitions of O ₂ Sensitivity and Oxidative Inactivation in Hydrogenases. <i>Journal of the American Chemical Society</i> , 2005, 127, 18179-18189.	13.7	208
77	Structural bases for the catalytic mechanism of Ni-containing carbon monoxide dehydrogenases. <i>Dalton Transactions</i> , 2005, , 3443.	3.3	50
78	Crystallographic evidence for a CO/CO ₂ tunnel gating mechanism in the bifunctional carbon monoxide dehydrogenase/acetyl coenzyme A synthase from <i>Moorella thermoacetica</i> . <i>Journal of Biological Inorganic Chemistry</i> , 2004, 9, 525-532.	2.6	44
79	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.	8.2	418
80	The active site and catalytic mechanism of NiFe hydrogenases. <i>Dalton Transactions</i> , 2003, , 4030-4038.	3.3	123
81	Density Functional Calculations for Modeling the Active Site of Nickel–Iron Hydrogenases. 2. Predictions for the Unready and Ready States and the Corresponding Activation Processes. <i>Inorganic Chemistry</i> , 2002, 41, 4424-4434.	4.0	68
82	High-resolution crystallographic analysis of <i>Desulfovibrio fructosovorans</i> 6NiFe ₉ hydrogenase. <i>International Journal of Hydrogen Energy</i> , 2002, 27, 1449-1461.	7.1	140
83	Crystallographic and FTIR Spectroscopic Evidence of Changes in Fe Coordination Upon Reduction of the Active Site of the Fe-Only Hydrogenase from <i>Desulfovibriodesulfuricans</i> . <i>Journal of the American Chemical Society</i> , 2001, 123, 1596-1601.	13.7	761
84	Crystal Structure of the Free Radical Intermediate of Pyruvate:ferredoxin Oxidoreductase. <i>Science</i> , 2001, 294, 2559-2563.	12.6	143
85	A novel FeS cluster in Fe-only hydrogenases. <i>Trends in Biochemical Sciences</i> , 2000, 25, 138-143.	7.5	401
86	Structural Model of the Fe-Hydrogenase/Cytochrome c 553 Complex Combining Transverse Relaxation-optimized Spectroscopy Experiments and Soft Docking Calculations. <i>Journal of Biological Chemistry</i> , 2000, 275, 23204-23210.	3.4	39
87	Combination of methods used in the structure solution of pyruvate:ferredoxin oxidoreductase from two crystal forms. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1999, 55, 1546-1554.	2.5	6
88	Crystal structures of the key anaerobic enzyme pyruvate:ferredoxin oxidoreductase, free and in complex with pyruvate. <i>Nature Structural Biology</i> , 1999, 6, 182-190.	9.7	175
89	<i>Desulfovibrio desulfuricans</i> iron hydrogenase: the structure shows unusual coordination to an active site Fe binuclear center. <i>Structure</i> , 1999, 7, 13-23.	3.3	1,320
90	The crystal structure of a reduced [NiFeSe] hydrogenase provides an image of the activated catalytic center. <i>Structure</i> , 1999, 7, 557-566.	3.3	448

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91	Nickel-iron-sulfur Active Sites: Hydrogenase and Co Dehydrogenase. <i>Advances in Inorganic Chemistry</i> , 1999, 47, 283-333.	1.0	55
92	Carboxy-Terminal Processing of the Large Subunit of [Fe] Hydrogenase from <i>Desulfovibrio desulfuricans</i> ATCC 7757. <i>Journal of Bacteriology</i> , 1999, 181, 2947-2952.	2.2	39
93	[3Fe-4S] to [4Fe-4S] cluster conversion in <i>Desulfovibrio fructosovorans</i> [NiFe] hydrogenase by site-directed mutagenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 11625-11630.	7.1	203
94	Gas access to the active site of Ni-Fe hydrogenases probed by X-ray crystallography and molecular dynamics. <i>Nature Structural Biology</i> , 1997, 4, 523-526.	9.7	325
95	Crystal structure of the nickel-iron hydrogenase from <i>Desulfovibrio gigas</i> . <i>Nature</i> , 1995, 373, 580-587.	27.8	1,532
96	Crystallization and Preliminary X-ray Diffraction Study of the Nickel-binding Protein NikA of <i>Escherichia coli</i> . <i>Journal of Molecular Biology</i> , 1994, 243, 353-355.	4.2	13
97	Catalytic Nickel-iron-sulfur Clusters: From Minerals to Enzymes. , 0, , 57-82.		24