

Takashi Fujishiro

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

922
citations

430874

18
h-index

454955

30
g-index

41
all docs

41
docs citations

41
times ranked

693
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural diversity of cysteine desulfurases involved in iron-sulfur cluster biosynthesis. <i>Biophysics and Physicobiology</i> , 2022, 19, n/a.	1.0	6
2	Cycloserine enantiomers inhibit PLP-dependent cysteine desulfurase SufS via distinct mechanisms. <i>FEBS Journal</i> , 2022, 289, 5947-5970.	4.7	2
3	Evidence for dynamic in vivo interconversion of the conformational states of IscU during iron-sulfur cluster biosynthesis. <i>Molecular Microbiology</i> , 2021, 115, 807-818.	2.5	6
4	Sulfur-mobilizing Enzymes Involved in Iron-sulfur Cluster Biosynthesis: Shared Structural Features and Functional Diversity. <i>Seibutsu Butsuri</i> , 2021, 61, 180-182.	0.1	0
5	A cyclic lipopeptide surfactin is a species-selective Hsp90 inhibitor that suppresses cyanobacterial growth. <i>Journal of Biochemistry</i> , 2021, 170, 255-264.	1.7	8
6	The Structure of the Dimeric State of IscU Harboring Two Adjacent [2Fe-2S] Clusters Provides Mechanistic Insights into Cluster Conversion to [4Fe-4S]. <i>Biochemistry</i> , 2021, 60, 1569-1572.	2.5	17
7	Crystal structure of <i>Escherichia coli</i> class II hybrid cluster protein, HCP, reveals a [4Fe-4S] cluster at the N-terminal protrusion. <i>FEBS Journal</i> , 2021, 288, 6752-6768.	4.7	6
8	The nickel-sirohydrochlorin formation mechanism of the ancestral class II chelatase CfbA in coenzyme F430 biosynthesis. <i>Chemical Science</i> , 2021, 12, 2172-2180.	7.4	2
9	Snapshots of PLP-substrate and PLP-product external aldimines as intermediates in two types of cysteine desulfurase enzymes. <i>FEBS Journal</i> , 2020, 287, 1138-1154.	4.7	19
10	2. Hydrogen development. , 2020, , 13-136.		0
11	Identification of IscU residues critical for de novo iron-sulfur cluster assembly. <i>Molecular Microbiology</i> , 2019, 112, 1769-1783.	2.5	13
12	Structure of sirohydrochlorin ferrocyclase SirB: the last of the structures of the class II chelatase family. <i>Dalton Transactions</i> , 2019, 48, 6083-6090.	3.3	5
13	Distinct roles for U-type proteins in iron-sulfur cluster biosynthesis revealed by genetic analysis of the <i>Bacillus subtilis</i> sufCDSUB operon. <i>Molecular Microbiology</i> , 2018, 107, 688-703.	2.5	20
14	Mapping the key residues of SufB and SufD essential for biosynthesis of iron-sulfur clusters. <i>Scientific Reports</i> , 2017, 7, 9387.	3.3	31
15	Zinc-Ligand Swapping Mediated Complex Formation and Sulfur Transfer between SufS and SufU for Iron-Sulfur Cluster Biogenesis in <i>Bacillus subtilis</i> . <i>Journal of the American Chemical Society</i> , 2017, 139, 18464-18467.	13.7	26
16	Towards artificial methanogenesis: biosynthesis of the [Fe]-hydrogenase cofactor and characterization of the semi-synthetic hydrogenase. <i>Faraday Discussions</i> , 2017, 198, 37-58.	3.2	29
17	A substrate-binding-state mimic of H ₂ O ₂ -dependent cytochrome P450 produced by one-point mutagenesis and peroxygenation of non-native substrates. <i>Catalysis Science and Technology</i> , 2016, 6, 5806-5811.	4.1	49
18	Identification of HcgC as a SAM-Dependent Pyridinol Methyltransferase in [Fe]-Hydrogenase Cofactor Biosynthesis. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 9648-9651.	13.8	18

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19	Identification of HcgC as a SAM-Dependent Pyridinol Methyltransferase in [Fe]-Hydrogenase Cofactor Biosynthesis. <i>Angewandte Chemie</i> , 2016, 128, 9800-9803.	2.0	9
20	Towards a functional identification of catalytically inactive [Fe]-hydrogenase paralogs. <i>FEBS Journal</i> , 2015, 282, 3412-3423.	4.7	7
21	Protein-pyridinol thioester precursor for biosynthesis of the organometallic acyl-iron ligand in [Fe]-hydrogenase cofactor. <i>Nature Communications</i> , 2015, 6, 6895.	12.8	26
22	Reconstitution of [Fe]-hydrogenase using model complexes. <i>Nature Chemistry</i> , 2015, 7, 995-1002.	13.6	92
23	A possible iron delivery function of the dinuclear iron center of HcgD in [Fe]-hydrogenase cofactor biosynthesis. <i>FEBS Letters</i> , 2014, 588, 2789-2793.	2.8	21
24	Crystal Structures of [Fe]-Hydrogenase in Complex with Inhibitory Isocyanides: Implications for the H ₂ O ₂ -Activation Site. <i>Angewandte Chemie - International Edition</i> , 2013, 52, 9656-9659.	13.8	50
25	Identification of the HcgB Enzyme in [Fe]-Hydrogenase-Cofactor Biosynthesis. <i>Angewandte Chemie - International Edition</i> , 2013, 52, 12555-12558.	13.8	25
26	Construction of biocatalysts using the myoglobin scaffold for the synthesis of indigo from indole. <i>Catalysis Science and Technology</i> , 2012, 2, 739-744.	4.1	21
27	Chiral-Substrate-Assisted Stereoselective Epoxidation Catalyzed by H ₂ O ₂ -Dependent Cytochrome P450 _{SP±} . <i>Chemistry - an Asian Journal</i> , 2012, 7, 2286-2293.	3.3	26
28	Non-covalent modification of the active site of cytochrome P450 for inverting the stereoselectivity of monooxygenation. <i>Tetrahedron Letters</i> , 2011, 52, 395-397.	1.4	23
29	Crystal Structure of H ₂ O ₂ -dependent Cytochrome P450 _{SP±} with Its Bound Fatty Acid Substrate. <i>Journal of Biological Chemistry</i> , 2011, 286, 29941-29950.	3.4	103
30	Aromatic C-H bond hydroxylation by P450 peroxygenases: a facile colorimetric assay for monooxygenation activities of enzymes based on Russig's blue formation. <i>Journal of Biological Inorganic Chemistry</i> , 2010, 15, 1109-1115.	2.6	37
31	Understanding substrate misrecognition of hydrogen peroxide dependent cytochrome P450 from <i>Bacillus subtilis</i> . <i>Journal of Biological Inorganic Chemistry</i> , 2010, 15, 1331-1339.	2.6	35
32	Hydrogen Peroxide Dependent Monooxygenations by Tricking the Substrate Recognition of Cytochrome P450 _{BS±} . <i>Angewandte Chemie - International Edition</i> , 2007, 46, 3656-3659.	13.8	132
33	Inside Cover: Hydrogen Peroxide Dependent Monooxygenations by Tricking the Substrate Recognition of Cytochrome P450 _{BS±} (<i>Angew. Chem. Int. Ed.</i> 20/2007). <i>Angewandte Chemie - International Edition</i> , 2007, 46, 3592-3592.	13.8	0
34	6 Structure and function of [Fe]-hydrogenase and biosynthesis of the FeGP cofactor. , 0, , .		1