

Yoshitsugu Y Shiro

List of Publications by Year in descending order

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215
papers

8,846
citations

36303

51
h-index

60623

81
g-index

222
all docs

222
docs citations

222
times ranked

7033
citing authors

#	ARTICLE	IF	CITATIONS
1	Capacity of extracellular globins to reduce liver fibrosis via scavenging reactive oxygen species and promoting MMP-1 secretion. <i>Redox Biology</i> , 2022, 52, 102286.	9.0	3
2	Structural basis for heme detoxification by an ATP-binding cassette-type efflux pump in gram-positive pathogenic bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	10
3	Spatially restricted substrate-binding site of cortisol-synthesizing CYP11B1 limits multiple hydroxylations and hinders aldosterone synthesis. <i>Current Research in Structural Biology</i> , 2021, 3, 192-205.	2.2	1
4	NO Dynamics in Microbial Denitrification System. <i>Chemistry Letters</i> , 2021, 50, 280-288.	1.3	6
5	Impact of membrane protein-lipid interactions on formation of bilayer lipid membranes on SAM-modified gold electrode. <i>Electrochimica Acta</i> , 2021, 373, 137888.	5.2	8
6	Heme controls the structural rearrangement of its sensor protein mediating the hemolytic bacterial survival. <i>Communications Biology</i> , 2021, 4, 467.	4.4	8
7	Short-lived intermediate in N ₂ O generation by P450 NO reductase captured by time-resolved IR spectroscopy and XFEL crystallography. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	21
8	Regulatory Switching by Concerted Motions on the Microsecond Time Scale of the Oxygen Sensor Protein FixL. <i>Journal of Physical Chemistry B</i> , 2021, 125, 6847-6856.	2.6	6
9	Timing of NO Binding and Protonation in the Catalytic Reaction of Bacterial Nitric Oxide Reductase as Established by Time-Resolved Spectroscopy. <i>Bulletin of the Chemical Society of Japan</i> , 2020, 93, 825-833.	3.2	15
10	Crystals in Minutes: Instant On-Site Microcrystallisation of Various Flavours of the CYP102A1 (P450BM3) Haem Domain. <i>Angewandte Chemie - International Edition</i> , 2020, 59, 7611-7618.	13.8	13
11	Kristalle in Minutenschnelle: Sofortige Mikrokristallisation verschiedenster Varianten der CYP102A1-(P450BM3)-Haemdomäne. <i>Angewandte Chemie</i> , 2020, 132, 7681-7689.	2.0	6
12	The active form of quinol-dependent nitric oxide reductase from <i>Neisseria meningitidis</i> is a dimer. <i>IUCr</i> , 2020, 7, 404-415.	2.2	10
13	Functional Studies on Hemoproteins and Heme-enzymes Based on Their Molecular Structures. <i>Bulletin of Japan Society of Coordination Chemistry</i> , 2020, 75, 51-56.	0.2	0
14	Hijacking the Heme Acquisition System of <i>Pseudomonas aeruginosa</i> for the Delivery of Phthalocyanine as an Antimicrobial. <i>ACS Chemical Biology</i> , 2019, 14, 1637-1642.	3.4	27
15	UV Resonance Raman Characterization of a Substrate Bound to Human Indoleamine 2,3-Dioxygenase 1. <i>Biophysical Journal</i> , 2019, 117, 706-716.	0.5	1
16	Highly malleable haem-binding site of the haemoprotein HasA permits stable accommodation of bulky tetraphenylporphycenes. <i>RSC Advances</i> , 2019, 9, 18697-18702.	3.6	13
17	Dimeric structures of quinol-dependent nitric oxide reductases (qNORs) revealed by cryo-electron microscopy. <i>Science Advances</i> , 2019, 5, eaax1803.	10.3	14
18	Mechanistic Insights into the Activation of Soluble Guanylate Cyclase by Carbon Monoxide: A Multistep Mechanism Proposed for the BAY 41-2272 Induced Formation of 5-Coordinate CO-Heme. <i>Biochemistry</i> , 2018, 57, 1620-1631.	2.5	8

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19	<i>Pseudomonas aeruginosa</i> overexpression system of nitric oxide reductase for in vivo and in vitro mutational analyses. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2018, 1859, 333-341.	1.0	5
20	Architecture of the complete oxygen-sensing FixL-FixJ two-component signal transduction system. <i>Science Signaling</i> , 2018, 11, .	3.6	38
21	Protein engineering of CYP105s for their industrial uses. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2018, 1866, 23-31.	2.3	16
22	$\hat{\pm}$ -Oxidative decarboxylation of fatty acids catalysed by cytochrome P450 peroxygenases yielding shorter-alkyl-chain fatty acids. <i>Catalysis Science and Technology</i> , 2018, 8, 434-442.	4.1	27
23	Roles of N- and C-terminal domains in the ligand-binding properties of cytoglobin. <i>Journal of Inorganic Biochemistry</i> , 2018, 179, 1-9.	3.5	15
24	Reconstitution of full-length P450BM3 with an artificial metal complex by utilising the transpeptidase Sortase A. <i>Chemical Communications</i> , 2018, 54, 7892-7895.	4.1	23
25	Characterization of the quinol-dependent nitric oxide reductase from the pathogen <i>Neisseria meningitidis</i> , an electrogenic enzyme. <i>Scientific Reports</i> , 2018, 8, 3637.	3.3	22
26	Surface-Enhanced Infrared Absorption Spectroscopy of Bacterial Nitric Oxide Reductase under Electrochemical Control Using a Vibrational Probe of Carbon Monoxide. <i>Journal of Physical Chemistry Letters</i> , 2018, 9, 5196-5200.	4.6	17
27	Direct Hydroxylation of Benzene to Phenol by Cytochrome P450BM3 Triggered by Amino Acid Derivatives. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 10324-10329.	13.8	62
28	Direct Hydroxylation of Benzene to Phenol by Cytochrome P450BM3 Triggered by Amino Acid Derivatives. <i>Angewandte Chemie</i> , 2017, 129, 10460-10465.	2.0	23
29	Production of an active form of vitamin D 2 by genetically engineered CYP105A1. <i>Biochemical and Biophysical Research Communications</i> , 2017, 486, 336-341.	2.1	13
30	Dynamics of nitric oxide controlled by protein complex in bacterial system. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 9888-9893.	7.1	35
31	Structures of the Heme Acquisition Protein HasA with Iron(III) $\hat{\pm}$ 5,15 $\hat{\pm}$ Diphenylporphyrin and Derivatives Thereof as an Artificial Prosthetic Group. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 15279-15283.	13.8	15
32	Structures of the Heme Acquisition Protein HasA with Iron(III) $\hat{\pm}$ 5,15 $\hat{\pm}$ Diphenylporphyrin and Derivatives Thereof as an Artificial Prosthetic Group. <i>Angewandte Chemie</i> , 2017, 129, 15481-15485.	2.0	6
33	Manganese(V) Porphycene Complex Responsible for Inert C $\hat{\pm}$ H Bond Hydroxylation in a Myoglobin Matrix. <i>Journal of the American Chemical Society</i> , 2017, 139, 18460-18463.	13.7	60
34	Capturing an initial intermediate during the P450 _{nor} enzymatic reaction using time-resolved XFEL crystallography and caged-substrate. <i>Nature Communications</i> , 2017, 8, 1585.	12.8	74
35	Control of stereoselectivity of benzylic hydroxylation catalysed by wild-type cytochrome P450BM3 using decoy molecules. <i>Catalysis Science and Technology</i> , 2017, 7, 3332-3338.	4.1	30
36	A nearly on-axis spectroscopic system for simultaneously measuring UV $\hat{\pm}$ visible absorption and X-ray diffraction in the SPring-8 structural genomics beamline. <i>Journal of Synchrotron Radiation</i> , 2016, 23, 334-338.	2.4	4

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37	Regulatory Implications of Structural Changes in Tyr201 of the Oxygen Sensor Protein FixL. <i>Biochemistry</i> , 2016, 55, 4027-4035.	2.5	9
38	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
39	A Study of the Dynamics of the Heme Pocket and C-helix in CooA upon CO Dissociation Using Time-Resolved Visible and UV Resonance Raman Spectroscopy. <i>Journal of Physical Chemistry B</i> , 2016, 120, 7836-7843.	2.6	7
40	Crystal structure of bacterial haem importer complex in the inward-facing conformation. <i>Nature Communications</i> , 2016, 7, 13411.	12.8	40
41	CHAPTER 6. Structure and Function of Nitric Oxide Reductases. 2-Oxoglutarate-Dependent Oxygenases, 2016, , 114-140.	0.8	1
42	Activation of Wild-Type Cytochrome P450BM3 by the Next Generation of Decoy Molecules: Enhanced Hydroxylation of Gaseous Alkanes and Crystallographic Evidence. <i>ACS Catalysis</i> , 2015, 5, 150-156.	11.2	73
43	Initial O ₂ Insertion Step of the Tryptophan Dioxygenase Reaction Proposed by a Heme-Modification Study. <i>Biochemistry</i> , 2015, 54, 3604-3616.	2.5	24
44	Structure of the response regulator ChrA in the haem-sensing two-component system of <i>Corynebacterium diphtheriae</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2015, 71, 966-971.	0.8	5
45	Structure-Function Analyses of Cytochrome P450revl Involved in Reveromycin A Biosynthesis and Evaluation of the Biological Activity of Its Substrate, Reveromycin T. <i>Journal of Biological Chemistry</i> , 2014, 289, 32446-32458.	3.4	18
46	Structures of reduced and ligand-bound nitric oxide reductase provide insights into functional differences in respiratory enzymes. <i>Proteins: Structure, Function and Bioinformatics</i> , 2014, 82, 1258-1271.	2.6	29
47	Inhibition of Heme Uptake in <i>Pseudomonas aeruginosa</i> by its Hemophore (HasA _p) Bound to Synthetic Metal Complexes. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 2862-2866.	13.8	34
48	Disulfide bonds regulate binding of exogenous ligand to human cytoglobin. <i>Journal of Inorganic Biochemistry</i> , 2014, 135, 20-27.	3.5	32
49	H ₂ O ₂ -dependent substrate oxidation by an engineered diiron site in a bacterial hemerythrin. <i>Chemical Communications</i> , 2014, 50, 3421-3423.	4.1	9
50	Constructing new proton pathways in nitric oxide reductases. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2014, 1837, e102.	1.0	0
51	Palladium-Nanoparticle-Catalyzed 1,7-Palladium Migration Involving C-H Activation, Followed by Intramolecular Amination: Regioselective Synthesis of N1-Arylbenzotriazoles and an Evaluation of Their Inhibitory Activity toward Indoleamine 2,3-Dioxygenase. <i>Journal of Organic Chemistry</i> , 2014, 79, 6366-6371.	3.2	43
52	Characterization of quinol-dependent nitric oxide reductase from <i>Geobacillus stearothermophilus</i> : Enzymatic activity and active site structure. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2014, 1837, 1019-1026.	1.0	12
53	Crystal Structure, Exogenous Ligand Binding, and Redox Properties of an Engineered Diiron Active Site in a Bacterial Hemerythrin. <i>Inorganic Chemistry</i> , 2013, 52, 13014-13020.	4.0	10
54	Resonance Raman study on indoleamine 2,3-dioxygenase: Control of reactivity by substrate-binding. <i>Chemical Physics</i> , 2013, 419, 178-183.	1.9	1

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55	Crystal structures of nitric oxide reductases provide key insights into functional conversion of respiratory enzymes. <i>IUBMB Life</i> , 2013, 65, 217-226.	3.4	33
56	Ultraviolet Resonance Raman Observations of the Structural Dynamics of Rhizobial Oxygen Sensor FixL on Ligand Recognition. <i>Journal of Physical Chemistry B</i> , 2013, 117, 15786-15791.	2.6	13
57	Structural basis for nitrous oxide generation by bacterial nitric oxide reductases. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2012, 367, 1195-1203.	4.0	47
58	Structural Basis for the Transcriptional Regulation of Heme Homeostasis in <i>Lactococcus lactis</i> . <i>Journal of Biological Chemistry</i> , 2012, 287, 30755-30768.	3.4	55
59	Diversity and Substrate Specificity in the Structures of Steroidogenic Cytochrome P450 Enzymes. <i>Biological and Pharmaceutical Bulletin</i> , 2012, 35, 818-823.	1.4	39
60	Coupling Reaction of Indolepyruvic Acid by StaD and Its Product: Implications for Biosynthesis of Indolocarbazole and Violacein. <i>ChemBioChem</i> , 2012, 13, 2495-2500.	2.6	14
61	Interactions of Soluble Guanylate Cyclase with a P-Site Inhibitor: Effects of Gaseous Heme Ligands, Azide, and Allosteric Activators on the Binding of 2 α -Deoxy-3 β -GMP. <i>Biochemistry</i> , 2012, 51, 9277-9289.	2.5	7
62	Structure and function of bacterial nitric oxide reductases. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 1907-1913.	1.0	85
63	Proton transfer in the quinol-dependent nitric oxide reductase from <i>Geobacillus stearothermophilus</i> during reduction of oxygen. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 1914-1920.	1.0	11
64	Crystal structure of quinol-dependent nitric oxide reductase from <i>Geobacillus stearothermophilus</i> . <i>Nature Structural and Molecular Biology</i> , 2012, 19, 238-245.	8.2	106
65	Structural basis for oxygen sensing and signal transduction of the heme-based sensor protein Aer2 from <i>Pseudomonas aeruginosa</i> . <i>Chemical Communications</i> , 2012, 48, 6523.	4.1	29
66	Chiral α -Substrate α -Assisted Stereoselective Epoxidation Catalyzed by H ₂ O ₂ -Dependent Cytochrome P450 _{SP1} . <i>Chemistry - an Asian Journal</i> , 2012, 7, 2286-2293.	3.3	26
67	Molecular structure and function of bacterial nitric oxide reductase. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 680-687.	1.0	52
68	Molecular Dynamics Simulations Reveal Proton Transfer Pathways in Cytochrome C-Dependent Nitric Oxide Reductase. <i>PLoS Computational Biology</i> , 2012, 8, e1002674.	3.2	27
69	Resonance Raman study on the oxygenated and the ferryl-oxo species of indoleamine 2,3-dioxygenase during catalytic turnover. <i>Faraday Discussions</i> , 2011, 148, 239-247.	3.2	17
70	Crystal Structure and Spectroscopic Studies of a Stable Mixed-Valent State of the Hemerythrin-like Domain of a Bacterial Chemotaxis Protein. <i>Inorganic Chemistry</i> , 2011, 50, 4892-4899.	4.0	20
71	Crystal structure of the carbon monoxide complex of human cytoglobin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2011, 79, 1143-1153.	2.6	22
72	Bioconversion of vitamin D to its active form by bacterial or mammalian cytochrome P450. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2011, 1814, 249-256.	2.3	30

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73	Application of micro-reactor chip technique for millisecond quenching of deuterium incorporation into 70S ribosomal protein complex. <i>International Journal of Mass Spectrometry</i> , 2011, 302, 132-138.	1.5	13
74	Characterization and Functional Modification of StaC and RebC, Which Are Involved in the Pyrrole Oxidation of Indolocarbazole Biosynthesis. <i>Bioscience, Biotechnology and Biochemistry</i> , 2011, 75, 2184-2193.	1.3	18
75	Crystal Structure of H ₂ O ₂ -dependent Cytochrome P450SP1± with Its Bound Fatty Acid Substrate. <i>Journal of Biological Chemistry</i> , 2011, 286, 29941-29950.	3.4	103
76	Identification of the Fe=O and the Fe=O Heme Species for Indoleamine 2,3-Dioxygenase during Catalytic Turnover. <i>Chemistry Letters</i> , 2010, 39, 36-37.	1.3	29
77	Unique Properties and Reactivity of High-Valent Manganese ^{IV} Oxo versus Manganese ^{III} Hydroxo in the Salen Platform. <i>Inorganic Chemistry</i> , 2010, 49, 6664-6672.	4.0	67
78	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
79	Three-step hydroxylation of vitamin D ₃ by a genetically engineered CYP105A1. <i>FEBS Journal</i> , 2010, 277, 3999-4009.	4.7	33
80	A Novel Glycerophosphodiester Phosphodiesterase, GDE5, Controls Skeletal Muscle Development via a Non-enzymatic Mechanism. <i>Journal of Biological Chemistry</i> , 2010, 285, 27652-27663.	3.4	49
81	Structural Basis of Biological N ₂ O Generation by Bacterial Nitric Oxide Reductase. <i>Science</i> , 2010, 330, 1666-1670.	12.6	292
82	Mg ²⁺ Dependence of 70 S Ribosomal Protein Flexibility Revealed by Hydrogen/Deuterium Exchange and Mass Spectrometry. <i>Journal of Biological Chemistry</i> , 2010, 285, 5646-5652.	3.4	18
83	Hydrophobic Residues Regulate Distal Histidine Coordinations in Human Cgb and Ngb. , 2010, , .		0
84	Excited States of Fluorescent Proteins, mKO and DsRed: Chromophore ⁺ Protein Electrostatic Interaction Behind the Color Variations. <i>Journal of Physical Chemistry B</i> , 2010, 114, 2971-2979.	2.6	43
85	A Specific Interaction of Tryptophan with CO of CO-Bound Indoleamine 2,3-Dioxygenase Identified by Resonance Raman Spectroscopy. <i>Biochemistry</i> , 2010, 49, 10081-10088.	2.5	6
86	ONIOM Study on a Missing Piece in Our Understanding of Heme Chemistry: Bacterial Tryptophan 2,3-Dioxygenase with Dual Oxidants. <i>Journal of the American Chemical Society</i> , 2010, 132, 11993-12005.	13.7	74
87	X-ray Crystal Structure of Michaelis Complex of Aldoxime Dehydratase. <i>Journal of Biological Chemistry</i> , 2009, 284, 32089-32096.	3.4	55
88	Cooperative Binding of L-Trp to Human Tryptophan 2,3-Dioxygenase: Resonance Raman Spectroscopic Analysis. <i>Journal of Biochemistry</i> , 2009, 145, 505-515.	1.7	18
89	Structure of PAS-Linked Histidine Kinase and the Response Regulator Complex. <i>Structure</i> , 2009, 17, 1333-1344.	3.3	93
90	Heme-dependent autophosphorylation of a heme sensor kinase, ChrS, from <i>Corynebacterium diphtheriae</i> reconstituted in proteoliposomes. <i>FEBS Letters</i> , 2009, 583, 2244-2248.	2.8	27

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91	Theoretical and Experimental Studies of the Conversion of Chromopyrrolic Acid to an Antitumor Derivative by Cytochrome P450 StaP: The Catalytic Role of Water Molecules. <i>Journal of the American Chemical Society</i> , 2009, 131, 6748-6762.	13.7	64
92	Crystal Structure of a New Cyan Fluorescent Protein and Its Hue-Shifted Variants. <i>Biochemistry</i> , 2009, 48, 5276-5283.	2.5	4
93	Ligand Energy Controls the Heme-Fe Valence in Aqueous Myoglobins. <i>Journal of the Physical Society of Japan</i> , 2009, 78, 044802.	1.6	22
94	Observation of a calcium-binding site in the \hat{I}^3 -class carbonic anhydrase from <i>Pyrococcus horikoshii</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2008, 64, 1012-1019.	2.5	48
95	Separation of phosphoprotein isotypes having the same number of phosphate groups using phosphate-affinity SDS-PAGE. <i>Proteomics</i> , 2008, 8, 2994-3003.	2.2	81
96	Molecular Design of Heteroprotein Assemblies Providing a Bionanocup as a Chemical Reactor. <i>Small</i> , 2008, 4, 50-54.	10.0	25
97	Synthesis and biological activity of 1-methyl-tryptophan-tirapazamine hybrids as hypoxia-targeting indoleamine 2,3-dioxygenase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2008, 16, 8661-8669.	3.0	24
98	Structural Basis of the Signal Transduction in the Two-Component System. <i>Advances in Experimental Medicine and Biology</i> , 2008, 631, 22-39.	1.6	22
99	Purification and functional characterization of human $11\hat{I}^2$ hydroxylase expressed in <i>Escherichia coli</i> . <i>FEBS Journal</i> , 2008, 275, 799-810.	4.7	44
100	Transient Intermediates from Mn(salen) with Sterically Hindered Mesityl Groups: Interconversion between Mn ^{IV} -Phenolate and Mn ^{III} -Phenoxy Radicals as an Origin for Unique Reactivity. <i>Inorganic Chemistry</i> , 2008, 47, 1674-1686.	4.0	77
101	Structural Characterization of a Thiazoline-Containing Chromophore in an Orange Fluorescent Protein, Monomeric Kusabira Orange. <i>Biochemistry</i> , 2008, 47, 11573-11580.	2.5	53
102	Density Functional Theory Study on a Missing Piece in Understanding of Heme Chemistry: The Reaction Mechanism for Indoleamine 2,3-Dioxygenase and Tryptophan 2,3-Dioxygenase. <i>Journal of the American Chemical Society</i> , 2008, 130, 12299-12309.	13.7	80
103	Structure-Based Design of a Highly Active Vitamin D Hydroxylase from <i>Streptomyces griseolus</i> CYP105A1. <i>Biochemistry</i> , 2008, 47, 11964-11972.	2.5	46
104	Crystal Structure of CYP105A1 (P450SU-1) in Complex with $1\hat{I}^{\pm,25}$ -Dihydroxyvitamin D ₃ . <i>Biochemistry</i> , 2008, 47, 4017-4027.	2.5	78
105	Crystal Structure of VioE, a Key Player in the Construction of the Molecular Skeleton of Violacein. <i>Journal of Biological Chemistry</i> , 2008, 283, 6459-6466.	3.4	38
106	<i>Escherichia coli</i> Cytosolic Glycerophosphodiester Phosphodiesterase (UgpQ) Requires Mg ²⁺ , Co ²⁺ , or Mn ²⁺ for Its Enzyme Activity. <i>Journal of Bacteriology</i> , 2008, 190, 1219-1223.	2.2	54
107	Light-dependent regulation of structural flexibility in a photochromic fluorescent protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 9227-9232.	7.1	150
108	Crystal structures and catalytic mechanism of cytochrome P450 StaP that produces the indolocarbazole skeleton. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 11591-11596.	7.1	108

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109	Mechanistic studies on the intramolecular one-electron transfer between the two flavins in the human endothelial NOS reductase domain. <i>Archives of Biochemistry and Biophysics</i> , 2007, 465, 254-265.	3.0	8
110	Structure and Ligand Binding Properties of Myoglobins Reconstituted with Monodepropionated Heme: Functional Role of Each Heme Propionate Side Chain. <i>Biochemistry</i> , 2007, 46, 9406-9416.	2.5	42
111	Resonance Raman Observation of the Structural Dynamics of FixL on Signal Transduction and Ligand Discrimination. <i>Biochemistry</i> , 2007, 46, 6086-6096.	2.5	23
112	X-ray structure and reaction mechanism of human indoleamine 2,3-dioxygenase. <i>International Congress Series</i> , 2007, 1304, 85-97.	0.2	0
113	Hydrogen Peroxide Dependent Monooxygenations by Tricking the Substrate Recognition of Cytochrome P450BSI ² . <i>Angewandte Chemie - International Edition</i> , 2007, 46, 3656-3659.	13.8	132
114	Separation of a phosphorylated histidine protein using phosphate affinity polyacrylamide gel electrophoresis. <i>Analytical Biochemistry</i> , 2007, 360, 160-162.	2.4	50
115	Crystallization and preliminary crystallographic analysis of molybdenum-cofactor biosynthesis protein C from <i>Thermus thermophilus</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007, 63, 27-29.	0.7	4
116	Cloning, expression, purification, crystallization and preliminary X-ray crystallographic study of DHNA synthetase from <i>Geobacillus kaustophilus</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007, 63, 103-105.	0.7	4
117	Cloning, expression, purification, crystallization and preliminary X-ray crystallographic study of molybdopterin synthase from <i>Thermus thermophilus</i> HB8. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007, 63, 324-326.	0.7	2
118	Preliminary X-ray crystallographic study of glucose dehydrogenase from <i>Thermus thermophilus</i> HB8. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007, 63, 446-448.	0.7	1
119	Crystal Structure and Peroxidase Activity of Myoglobin Reconstituted with Iron Porphycene. <i>Inorganic Chemistry</i> , 2006, 45, 10530-10536.	4.0	89
120	Roles of the Heme Distal Residues of FixL in O ₂ Sensing: A Single Convergent Structure of the Heme Moiety Is Relevant to the Downregulation of Kinase Activity. <i>Biochemistry</i> , 2006, 45, 2515-2523.	2.5	31
121	1.25 Å Resolution Crystal Structures of Human Haemoglobin in the Oxy, Deoxy and Carbonmonoxy Forms. <i>Journal of Molecular Biology</i> , 2006, 360, 690-701.	4.2	261
122	The Signaling Pathway in Histidine Kinase and the Response Regulator Complex Revealed by X-ray Crystallography and Solution Scattering. <i>Journal of Molecular Biology</i> , 2006, 362, 123-139.	4.2	27
123	High-resolution structure of human cytoglobin: identification of extra N- and C-termini and a new dimerization mode. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2006, 62, 671-677.	2.5	28
124	Crystallization and preliminary crystallographic studies of human indoleamine 2,3-dioxygenase. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006, 62, 221-223.	0.7	8
125	Design and Synthesis of De Novo Peptide for Manganese Binding. <i>International Journal of Peptide Research and Therapeutics</i> , 2006, 12, 379-385.	1.9	10
126	Crystal structure of human indoleamine 2,3-dioxygenase: Catalytic mechanism of O ₂ incorporation by a heme-containing dioxygenase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 2611-2616.	7.1	389

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127	Tolerance of the Rieske-type [2Fe-2S] cluster in recombinant ferredoxin BphA3 from <i>Pseudomonas</i> sp. KKS102 to histidine ligand mutations. <i>Biochemical Journal</i> , 2005, 388, 869-878.	3.7	22
128	Solution Structure of the C-Terminal Transcriptional Activator Domain of FixJ from <i>Sinorhizobium meliloti</i> and Its Recognition of the fixK Promoter. <i>Biochemistry</i> , 2005, 44, 14835-14844.	2.5	20
129	Structural Characterization of the Proximal and Distal Histidine Environment of Cytoglobin and Neuroglobin. <i>Biochemistry</i> , 2005, 44, 13257-13265.	2.5	62
130	Design of a Cro Fold: Solution Structure of a Monomeric Variant of the De Novo Protein. <i>Journal of Molecular Biology</i> , 2005, 354, 801-814.	4.2	14
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