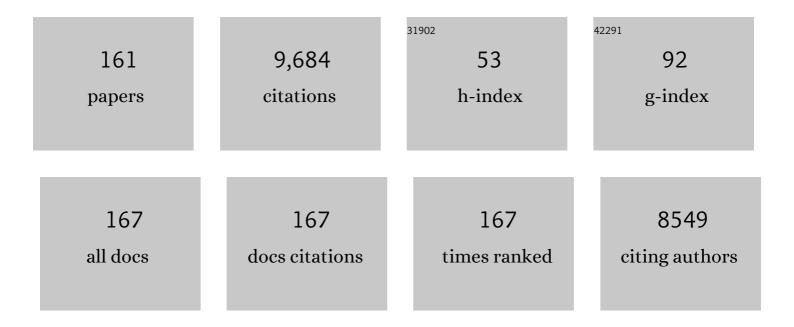
List of Publications by Year in descending order

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RDIAN C FOX

#	Article	IF	CITATIONS
1	Identification and characterization of a set of monocot BAHD monolignol transferases. Plant Physiology, 2022, 189, 37-48.	2.3	10
2	<i>p</i> HBMT1, a BAHD-family monolignol acyltransferase, mediates lignin acylation in poplar. Plant Physiology, 2022, 188, 1014-1027.	2.3	18
3	Multifunctional cellulases are potent, versatile tools for a renewable bioeconomy. Current Opinion in Biotechnology, 2021, 67, 141-148.	3.3	14
4	Mannose- and Mannobiose-Specific Responses of the Insect-Associated Cellulolytic Bacterium <i>Streptomyces</i> sp. Strain SirexAA-E. Applied and Environmental Microbiology, 2021, 87, e0271920.	1.4	6
5	Spectroscopic investigation of iron(III) cysteamine dioxygenase in the presence of substrate (analogs): implications for the nature of substrate-bound reaction intermediates. Journal of Biological Inorganic Chemistry, 2021, 26, 947-955.	1.1	2
6	The Crystal Structure of Cysteamine Dioxygenase Reveals the Origin of the Large Substrate Scope of This Vital Mammalian Enzyme. Biochemistry, 2021, 60, 3728-3737.	1.2	10
7	PreDSLpmo: A neural network-based prediction tool for functional annotation of lytic polysaccharide monooxygenases. Journal of Biotechnology, 2020, 308, 148-155.	1.9	0
8	A structural and kinetic survey of GH5_4 endoglucanases reveals determinants of broad substrate specificity and opportunities for biomass hydrolysis. Journal of Biological Chemistry, 2020, 295, 17752-17769.	1.6	15
9	A bacterial biosynthetic pathway for methylated furan fatty acids. Journal of Biological Chemistry, 2020, 295, 9786-9801.	1.6	18
10	Spectroscopic Investigation of Cysteamine Dioxygenase. Biochemistry, 2020, 59, 2450-2458.	1.2	10
11	Solid-State NMR Studies of Solvent-Mediated, Acid-Catalyzed Woody Biomass Pretreatment for Enzymatic Conversion of Residual Cellulose. ACS Sustainable Chemistry and Engineering, 2020, 8, 6551-6563.	3.2	10
12	Extent and Origins of Functional Diversity in a Subfamily of Glycoside Hydrolases. Journal of Molecular Biology, 2019, 431, 1217-1233.	2.0	13
13	Determination of glycoside hydrolase specificities during hydrolysis of plant cell walls using glycome profiling. Biotechnology for Biofuels, 2017, 10, 31.	6.2	18
14	In-crystal reaction cycle of a toluene-bound diiron hydroxylase. Nature, 2017, 544, 191-195.	13.7	45
15	Reply to Kiser: Dioxygen binding in NOV1 crystal structures. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E6029-E6030.	3.3	4
16	Functional characterization of three GH10 xylanases. FASEB Journal, 2017, 31, 607.1.	0.2	0
17	Structure and mechanism of NOV1, a resveratrol-cleaving dioxygenase. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 14324-14329.	3.3	50
18	Cell-free translation and purification of Arabidopsis thaliana regulator of G signaling 1 protein. Protein Expression and Purification, 2016, 126, 33-41.	0.6	8

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19	Evolution and Ecology of <i>Actinobacteria</i> and Their Bioenergy Applications. Annual Review of Microbiology, 2016, 70, 235-254.	2.9	249
20	Evolution of High Cellulolytic Activity in Symbiotic Streptomyces through Selection of Expanded Gene Content and Coordinated Gene Expression. PLoS Biology, 2016, 14, e1002475.	2.6	68
21	Development of a High Throughput Platform for Screening Glycoside Hydrolases Based on Oxime-NIMS. Frontiers in Bioengineering and Biotechnology, 2015, 3, 153.	2.0	14
22	Use of Nanostructure-Initiator Mass Spectrometry to Deduce Selectivity of Reaction in Glycoside Hydrolases. Frontiers in Bioengineering and Biotechnology, 2015, 3, 165.	2.0	6
23	Multifunctional cellulase catalysis targeted by fusion to different carbohydrate-binding modules. Biotechnology for Biofuels, 2015, 8, 220.	6.2	49
24	Spectroscopic and Computational Investigation of the H155A Variant of Cysteine Dioxygenase: Geometric and Electronic Consequences of a Third-Sphere Amino Acid Substitution. Biochemistry, 2015, 54, 2874-2884.	1.2	26
25	X-ray structure of a mammalian stearoyl-CoA desaturase. Nature, 2015, 524, 252-256.	13.7	213
26	Expression platforms for producing eukaryotic proteins: a comparison of E. coli cell-based and wheat germ cell-free synthesis, affinity and solubility tags, and cloning strategies. Journal of Structural and Functional Genomics, 2015, 16, 67-80.	1.2	12
27	Active Site and Laminarin Binding in Glycoside Hydrolase Family 55. Journal of Biological Chemistry, 2015, 290, 11819-11832.	1.6	36
28	Structure of T4moF, the Toluene 4-Monooxygenase Ferredoxin Oxidoreductase. Biochemistry, 2015, 54, 5980-5988.	1.2	8
29	Structureâ€guided analysis of catalytic specificity of the abundantly secreted chitosanase SACTE_5457 from <i>Streptomyces</i> sp. SirexAAâ€E. Proteins: Structure, Function and Bioinformatics, 2014, 82, 1245-1257.	1.5	33
30	Structural basis for biomolecular recognition in overlapping binding sites in a diiron enzyme system. Nature Communications, 2014, 5, 5009.	5.8	24
31	Evolution of substrate specificity in bacterial AA10 lytic polysaccharide monooxygenases. Biotechnology for Biofuels, 2014, 7, 109.	6.2	69
32	The Oligomeric States of the Purified Sigma-1 Receptor Are Stabilized by Ligands. Journal of Biological Chemistry, 2014, 289, 20333-20344.	1.6	92
33	Rapid Kinetic Characterization of Glycosyl Hydrolases Based on Oxime Derivatization and Nanostructure-Initiator Mass Spectrometry (NIMS). ACS Chemical Biology, 2014, 9, 1470-1479.	1.6	36
34	Cell-Free Protein Synthesis for Functional and Structural Studies. Methods in Molecular Biology, 2014, 1091, 161-178.	0.4	19
35	Spectroscopic and Computational Investigation of Iron(III) Cysteine Dioxygenase: Implications for the Nature of the Putative Superoxo-Fe(III) Intermediate. Biochemistry, 2014, 53, 5759-5770.	1.2	26
36	Cellulolytic Streptomyces Strains Associated with Herbivorous Insects Share a Phylogenetically Linked Capacity To Degrade Lignocellulose. Applied and Environmental Microbiology, 2014, 80, 4692-4701.	1.4	70

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37	Functional Evolution of Ribonuclease Inhibitor: Insights from Birds and Reptiles. Journal of Molecular Biology, 2014, 426, 3041-3056.	2.0	56
38	Expression, purification and characterization of a functional carbohydrate-binding module from Streptomyces sp. SirexAA-E. Protein Expression and Purification, 2014, 98, 1-9.	0.6	24
39	Cell-Free Translation of Biofuel Enzymes. Methods in Molecular Biology, 2014, 1118, 71-95.	0.4	24
40	Biochemical Properties and Atomic Resolution Structure of a Proteolytically Processed β-Mannanase from Cellulolytic Streptomyces sp. SirexAA-E. PLoS ONE, 2014, 9, e94166.	1.1	18
41	Coordinating the impact of structural genomics on the human α-helical transmembrane proteome. Nature Structural and Molecular Biology, 2013, 20, 135-138.	3.6	64
42	Improved expression and purification of sigma 1 receptor fused to maltose binding protein by alteration of linker sequence. Protein Expression and Purification, 2013, 89, 203-209.	0.6	11
43	Aerobic deconstruction of cellulosic biomass by an insect-associated Streptomyces. Scientific Reports, 2013, 3, 1030.	1.6	107
44	Spectroscopic and Computational Characterization of the NO Adduct of Substrate-Bound Fe(II) Cysteine Dioxygenase: Insights into the Mechanism of O ₂ Activation. Biochemistry, 2013, 52, 6040-6051.	1.2	32
45	Mutations in FLS2 Ser-938 Dissect Signaling Activation in FLS2-Mediated Arabidopsis Immunity. PLoS Pathogens, 2013, 9, e1003313.	2.1	57
46	Cellâ€free production of integral membrane aspartic acid proteases reveals zincâ€dependent methyltransferase activity of the <i><scp>P</scp>seudomonas aeruginosa</i> prepilin peptidase PilD. MicrobiologyOpen, 2013, 2, 94-104.	1.2	21
47	Fusion of Dioxygenase and Lignin-binding Domains in a Novel Secreted Enzyme from Cellulolytic Streptomyces sp. SirexAA-E. Journal of Biological Chemistry, 2013, 288, 18574-18587.	1.6	40
48	Function of Shaker potassium channels produced by cell-free translation upon injection into Xenopus oocytes. Scientific Reports, 2013, 3, 1040.	1.6	22
49	Crystallographic Analysis of Active Site Contributions to Regiospecificity in the Diiron Enzyme Toluene 4-Monooxygenase. Biochemistry, 2012, 51, 1101-1113.	1.2	19
50	Amino Acid Determinants of Substrate Selectivity in the <i>Trypanosoma brucei</i> Sphingolipid Synthase Family. Biochemistry, 2011, 50, 8853-8861.	1.2	8
51	Robotic large-scale application of wheat cell-free translation to structural studies including membrane proteins. New Biotechnology, 2011, 28, 239-249.	2.4	24
52	Structure of cellobiose phosphorylase from <i>Clostridium thermocellum</i> in complex with phosphate. Acta Crystallographica Section F: Structural Biology Communications, 2011, 67, 1345-1349.	0.7	25
53	Structural architecture of <i>Galdieria sulphuraria</i> DCN1L. Proteins: Structure, Function and Bioinformatics, 2011, 79, 1329-1336.	1.5	4
54	Global Gene Expression Patterns in <i>Clostridium thermocellum</i> as Determined by Microarray Analysis of Chemostat Cultures on Cellulose or Cellobiose. Applied and Environmental Microbiology, 2011, 77, 1243-1253.	1.4	75

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55	Maltose–neopentyl glycol (MNG) amphiphiles for solubilization, stabilization and crystallization of membrane proteins. Nature Methods, 2010, 7, 1003-1008.	9.0	397
56	Cell-free Synthesis and Functional Characterization of Sphingolipid Synthases from Parasitic Trypanosomatid Protozoa. Journal of Biological Chemistry, 2010, 285, 20580-20587.	1.6	37
57	Spectroscopic and Computational Characterization of Substrate-Bound Mouse Cysteine Dioxygenase: Nature of the Ferrous and Ferric Cysteine Adducts and Mechanistic Implications. Biochemistry, 2010, 49, 6033-6041.	1.2	63
58	Cell-Free Protein Synthesis Technology in NMR High-Throughput Structure Determination. Methods in Molecular Biology, 2010, 607, 127-147.	0.4	30
59	The Center for Eukaryotic Structural Genomics. Journal of Structural and Functional Genomics, 2009, 10, 165-179.	1.2	33
60	Transformation of RDX and other energetic compounds by xenobiotic reductases XenA and XenB. Applied Microbiology and Biotechnology, 2009, 84, 535-544.	1.7	65
61	Xâ€ray structure of ILL2, an auxinâ€conjugate amidohydrolase from <i>Arabidopsis thaliana</i> . Proteins: Structure, Function and Bioinformatics, 2009, 74, 61-71.	1.5	42
62	Discovery of sarcosine dimethylglycine methyltransferase from <i>Galdieria sulphuraria</i> . Proteins: Structure, Function and Bioinformatics, 2009, 74, 368-377.	1.5	11
63	Xâ€ray structure of <i>Danio rerio</i> secretagogin: A hexaâ€EFâ€hand calcium sensor. Proteins: Structure, Function and Bioinformatics, 2009, 76, 477-483.	1.5	22
64	X-ray structure of putative acyl-ACP desaturase DesA2 from Mycobacterium tuberculosis H37Rv. Protein Science, 2009, 14, 1508-1517.	3.1	34
65	Role for Threonine 201 in the Catalytic Cycle of the Soluble Diiron Hydroxylase Toluene 4-Monooxygenase,. Biochemistry, 2009, 48, 3838-3846.	1.2	25
66	Crystallographic and Catalytic Studies of the Peroxide-Shunt Reaction in a Diiron Hydroxylase. Biochemistry, 2009, 48, 8932-8939.	1.2	50
67	Flexi Vector Cloning. Methods in Molecular Biology, 2009, 498, 55-73.	0.4	28
68	Autoinduction of Protein Expression. Current Protocols in Protein Science, 2009, 56, Unit 5.23.	2.8	61
69	Chapter 37 Cell-Free Translation of Integral Membrane Proteins into Unilamelar Liposomes. Methods in Enzymology, 2009, 463, 647-673.	0.4	43
70	X-ray structure of a soluble Rieske-type ferredoxin fromMus musculus. Acta Crystallographica Section D: Biological Crystallography, 2008, 64, 933-940.	2.5	5
71	Structural and functional characterization of a novel phosphatase from the <i>Arabidopsis thaliana</i> gene locus At1g05000. Proteins: Structure, Function and Bioinformatics, 2008, 73, 241-253.	1.5	17
72	Structural genomics: from genes to structures with valuable materials and many questions in between. Nature Methods, 2008, 5, 129-132.	9.0	45

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73	Enhanced Bacterial Protein Expression During Auto-Induction Obtained by Alteration of Lac Repressor Dosage and Medium Composition. Biotechnology Progress, 2008, 23, 585-598.	1.3	141
74	Structure of Human J-type Co-chaperone HscB Reveals a Tetracysteine Metal-binding Domain. Journal of Biological Chemistry, 2008, 283, 30184-30192.	1.6	38
75	Soluble expression and purification of the oxidoreductase component of toluene 4-monooxygenase. Protein Expression and Purification, 2008, 57, 9-16.	0.6	17
76	A Protein Structure Initiative approach to expression, purification, and in situ delivery of human cytochrome b5 to membrane vesicles. Protein Expression and Purification, 2008, 58, 229-241.	0.6	23
77	Wheat germ cell-free translation, purification, and assembly of a functional human stearoyl-CoA desaturase complex. Protein Expression and Purification, 2008, 62, 171-178.	0.6	82
78	Geometric and Electronic Structure Studies of the Binuclear Nonheme Ferrous Active Site of Toluene-4-monooxygenase: Parallels with Methane Monooxygenase and Insight into the Role of the Effector Proteins in O ₂ Activation. Journal of the American Chemical Society, 2008, 130, 7098-7109.	6.6	41
79	In Vivo Inactivation of the Mycobacterial Integral Membrane Stearoyl Coenzyme A Desaturase DesA3 by a C-Terminus-Specific Degradation Process. Journal of Bacteriology, 2008, 190, 6686-6696.	1.0	9
80	Structural consequences of effector protein complex formation in a diiron hydroxylase. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 19194-19198.	3.3	92
81	A combined approach to improving large-scale production of tobacco etch virus protease. Protein Expression and Purification, 2007, 55, 53-68.	0.6	240
82	Characterization of the Nitrosyl Adduct of Substrate-Bound Mouse Cysteine Dioxygenase by Electron Paramagnetic Resonance:  Electronic Structure of the Active Site and Mechanistic Implications. Biochemistry, 2007, 46, 8569-8578.	1.2	99
83	Structures of proteins of biomedical interest from the Center for Eukaryotic Structural Genomics. Journal of Structural and Functional Genomics, 2007, 8, 73-84.	1.2	10
84	Small-scale, semi-automated purification of eukaryotic proteins for structure determination. Journal of Structural and Functional Genomics, 2007, 8, 153-166.	1.2	27
85	Component Interactions and Implications for Complex Formation in the Multicomponent Toluene 4-Monooxygenase. Biochemistry, 2006, 45, 5478-5485.	1.2	11
86	Identification of Rv3230c as the NADPH Oxidoreductase of a Two-Protein DesA3 Acyl-CoA Desaturase in Mycobacterium tuberculosis H37Rv. Biochemistry, 2006, 45, 13476-13486.	1.2	25
87	Identification of the Binding Region of the [2Fe-2S] Ferredoxin in Stearoyl-Acyl Carrier Protein Desaturase:  Insight into the Catalytic Complex and Mechanism of Action. Biochemistry, 2006, 45, 4848-4858.	1.2	31
88	Solution Structures of Spinach Acyl Carrier Protein with Decanoate and Stearateâ€,‡. Biochemistry, 2006, 45, 5217-5227.	1.2	86
89	Preparation of isotopically labeled spinach acyl–acyl carrier protein for NMR structural studies. Protein Expression and Purification, 2006, 46, 446-455.	0.6	10
90	High efficiency single step production of expression plasmids from cDNA clones using the Flexi Vector cloning system. Protein Expression and Purification, 2006, 47, 562-570.	0.6	58

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91	Structure and mechanism of mouse cysteine dioxygenase. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 3084-3089.	3.3	174
92	Fluorescence anisotropy assay for proteolysis of specifically labeled fusion proteins. Analytical Biochemistry, 2005, 336, 75-86.	1.1	23
93	Reaction mechanisms of non-heme diiron hydroxylases characterized in whole cells. Journal of Inorganic Biochemistry, 2005, 99, 1998-2006.	1.5	49
94	High-throughput Purification and Quality Assurance of Arabidopsis thaliana Proteins for Eukaryotic Structural Genomics. Journal of Structural and Functional Genomics, 2005, 6, 143-147.	1.2	67
95	The structure at 2.4â€Ã resolution of the protein from gene locus At3g21360, a putative Fell/2-oxoglutarate-dependent enzyme fromArabidopsis thaliana. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 469-472.	0.7	4
96	Comparison of cell-based and cell-free protocols for producing target proteins from the Arabidopsis thaliana genome for structural studies. Proteins: Structure, Function and Bioinformatics, 2005, 59, 633-643.	1.5	56
97	Identification of transcribed sequences in Arabidopsis thaliana by using high-resolution genome tiling arrays. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 4453-4458.	3.3	147
98	Crystal Structures and Functional Studies of T4moD, the Toluene 4-Monooxygenase Catalytic Effector Proteinâ€,‡. Biochemistry, 2005, 44, 7131-7142.	1.2	24
99	Oxygen-18 tracer studies of enzyme reactions with radical/cation diagnostic probes. Biochemical and Biophysical Research Communications, 2005, 338, 240-249.	1.0	5
100	Expression, purification, and physical characterization of Escherichia coli lipoyl(octanoyl)transferase. Protein Expression and Purification, 2005, 39, 269-282.	0.6	35
101	Protocols for production of selenomethionine-labeled proteins in 2-L polyethylene terephthalate bottles using auto-induction medium. Protein Expression and Purification, 2005, 40, 256-267.	0.6	104
102	Auto-induction medium for the production of [U-15N]- and [U-13C, U-15N]-labeled proteins for NMR screening and structure determination. Protein Expression and Purification, 2005, 40, 268-278.	0.6	91
103	Production in two-liter beverage bottles of proteins for NMR structure determination labeled with either15N- or13C-15N. Journal of Structural and Functional Genomics, 2004, 5, 87-93.	1.2	22
104	Solution structure of T4moC, the Rieske ferredoxin component of the toluene 4-monooxygenase complex. Journal of Biological Inorganic Chemistry, 2004, 9, 945-953.	1.1	11
105	Results from high-throughput DNA cloning of Arabidopsis thaliana target genes using site-specific recombination. Journal of Structural and Functional Genomics, 2004, 5, 267-276.	1.2	74
106	Crystal structure of the protein from gene At3g17210 of Arabidopsis thaliana. Proteins: Structure, Function and Bioinformatics, 2004, 57, 218-220.	1.5	7
107	Crystal structure of At2g03760, a putative steroid sulfotransferase fromArabidopsis thaliana. Proteins: Structure, Function and Bioinformatics, 2004, 57, 854-857.	1.5	14
108	Crystallization and preliminary analysis of xenobiotic reductase A and ligand complexes fromPseudomonas putidall-B. Acta Crystallographica Section D: Biological Crystallography, 2004, 60, 957-961.	2.5	7

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109	Crystallization and preliminary analysis of xenobiotic reductase B fromPseudomonas fluorescensl-C. Acta Crystallographica Section D: Biological Crystallography, 2004, 60, 1289-1291.	2.5	11
110	Reactions of the Diiron Enzyme Stearoyl-Acyl Carrier Protein Desaturase. ChemInform, 2004, 35, no.	0.1	0
111	Remarkable Aliphatic Hydroxylation by the Diiron Enzyme Toluene 4-Monooxygenase in Reactions with Radical or Cation Diagnostic Probes Norcarane, 1,1-Dimethylcyclopropane, and 1,1-Diethylcyclopropaneâ€. Biochemistry, 2004, 43, 15688-15701.	1.2	35
112	Reactions of the Diiron Enzyme Stearoyl-Acyl Carrier Protein Desaturase. Accounts of Chemical Research, 2004, 37, 421-429.	7.6	157
113	Crystallization and preliminary analysis of native and N-terminal truncated isoforms of toluene-4-monooxygenase catalytic effector protein. Acta Crystallographica Section D: Biological Crystallography, 2003, 59, 572-575.	2.5	3
114	Rapid-Mix and Chemical Quench Studies of Ferredoxin-Reduced Stearoyl-Acyl Carrier Protein Desaturaseâ€. Biochemistry, 2003, 42, 5857-5866.	1.2	21
115	Insight into the mechanism of aromatic hydroxylation by toluene 4-monooxygenase by use of specifically deuterated toluene and p-xylene. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 3784-3789.	3.3	98
116	Chain Cleavage and Sulfoxidation of Thiastearoyl-ACP upon Reaction with Stearoyl-ACP Desaturaseâ€. Biochemistry, 2003, 42, 7828-7835.	1.2	15
117	Effects of sterculic acid on stearoyl-CoA desaturase in differentiating 3T3-L1 adipocytes. Biochemical and Biophysical Research Communications, 2003, 300, 316-326.	1.0	67
118	N-Isotope effects on the Raman spectra of Fe2S2 ferredoxin and Rieske ferredoxin: evidence for structural rigidity of metal sites. Journal of Biological Inorganic Chemistry, 2003, 8, 318-326.	1.1	36
119	Fluorescence Anisotropy Studies of Enzymeâ^'Substrate Complex Formation in Stearoyl-ACP Desaturase. Biochemistry, 2002, 41, 14472-14481.	1.2	22
120	Molecular Differences Caused by Differentiation of 3T3-L1 Preadipocytes in the Presence of either Dehydroepiandrosterone (DHEA) or 7-Oxo-DHEA. Biochemistry, 2002, 41, 5473-5482.	1.2	33
121	Desaturation, Chain Scission, and Register-Shift of Oxygen-Substituted Fatty Acids during Reaction with Stearoyl-ACP Desaturaseâ€. Biochemistry, 2002, 41, 10141-10148.	1.2	24
122	Combined Participation of Hydroxylase Active Site Residues and Effector Protein Binding in a Para to Ortho Modulation of Toluene 4-Monooxygenase Regiospecificity. Biochemistry, 2002, 41, 3176-3188.	1.2	88
123	Mössbauer and EPR Studies of the Photoactivation of Nitrile Hydratase. Biochemistry, 2001, 40, 7984-7991.	1.2	57
124	Solution Structure of the Toluene 4-Monooxygenase Effector Protein (T4moD)â€,‡. Biochemistry, 2001, 40, 3512-3524.	1.2	28
125	Assignment of 1H, 13C and 15N NMR signals from toluene 4-monooxygenase Rieske ferredoxin in its oxidized state. Journal of Biomolecular NMR, 2001, 21, 73-74.	1.6	0
126	Aromatic Hydroxylation Catalyzed by Toluene 4-Monooxygenase in Organic Solvent/Aqueous Buffer Mixtures. Applied Biochemistry and Biotechnology, 2001, 90, 187-198.	1.4	16

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127	Desaturation of trans-octadecenoyl-acyl carrier protein by stearoyl-acyl carrier protein Δ9 desaturase. Journal of Inorganic Biochemistry, 2000, 78, 7-14.	1.5	17
128	Assignment of 1H, 13C and 15N NMR signals in the toluene 4-monooxygenase effector protein. Journal of Biomolecular NMR, 2000, 16, 359-360.	1.6	1
129	Toluene Monooxygenase-Catalyzed Epoxidation of Alkenes. Applied and Environmental Microbiology, 2000, 66, 1877-1882.	1.4	68
130	Transformation of 2,4,6-Trinitrotoluene by Purified Xenobiotic Reductase B from Pseudomonas fluorescens I-C. Applied and Environmental Microbiology, 2000, 66, 4742-4750.	1.4	148
131	Optimized Expression and Purification of Toluene 4-Monooxygenase Hydroxylase. Protein Expression and Purification, 2000, 20, 58-65.	0.6	32
132	Chemical and Posttranslational Modification of Escherichia coli Acyl Carrier Protein for Preparation of Dansyl-Acyl Carrier Proteins. Protein Expression and Purification, 2000, 20, 274-284.	0.6	16
133	Threonine 201 in the Diiron Enzyme Toluene 4-Monooxygenase Is Not Required for Catalysis. Biochemistry, 2000, 39, 791-799.	1.2	53
134	Resonance Raman Studies of the Stoichiometric Catalytic Turnover of a Substrateâ^'Stearoyl-Acyl Carrier Protein Δ9Desaturase Complexâ€. Biochemistry, 2000, 39, 10507-10513.	1.2	14
135	Differential regulation of the stearoyl-CoA desaturase genes by thiazolidinediones in 3T3-L1 adipocytes. Journal of Lipid Research, 2000, 41, 1310-1316.	2.0	67
136	Role of Hydrophobic Partitioning in Substrate Selectivity and Turnover of theRicinuscommunisStearoyl Acyl Carrier Protein Δ9Desaturaseâ€. Biochemistry, 1999, 38, 12833-12840.	1.2	33
137	Mössbauer Studies of the Formation and Reactivity of a Quasi-Stable Peroxo Intermediate of Stearoyl-Acyl Carrier Protein Δ9-Desaturaseâ€. Biochemistry, 1999, 38, 12197-12204.	1.2	106
138	Circular Dichroism and Magnetic Circular Dichroism Studies of the Reduced Binuclear Non-Heme Iron Site of Stearoyl-ACP Δ9-Desaturase:  Substrate Binding and Comparison to Ribonucleotide Reductase. Journal of the American Chemical Society, 1999, 121, 2770-2783.	6.6	72
139	Spinach Holo-Acyl Carrier Protein: Overproduction and Phosphopantetheinylation inEscherichia coliBL21(DE3),in VitroAcylation, and Enzymatic Desaturation of Histidine-Tagged Isoform I. Protein Expression and Purification, 1999, 15, 314-326.	0.6	26
140	Application of Fed-Batch Fermentation to the Preparation of Isotopically Labeled or Selenomethionyl-Labeled Proteins. Protein Expression and Purification, 1999, 16, 109-119.	0.6	40
141	Cloning and Sequence Analysis of Two <i>Pseudomonas</i> Flavoprotein Xenobiotic Reductases. Journal of Bacteriology, 1999, 181, 6254-6263.	1.0	141
142	Designing ligands to achieve robust oxidation catalysts. Iron based systems. Coordination Chemistry Reviews, 1998, 174, 361-390.	9.5	71
143	The fundamental, versatile role of diiron enzymes in lipid metabolism. Lipid - Fett, 1998, 100, 103-113.	0.6	33
144	EXAFS and Mössbauer characterization of the Diiron(III) site in stearoyl-acyl carrier protein Δ9– desaturase. Journal of Biological Inorganic Chemistry, 1998, 3, 392-400.	1.1	25

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145	Peroxodiferric Intermediate of Stearoyl-Acyl Carrier Protein Δ9Desaturase: Oxidase Reactivity during Single Turnover and Implications for the Mechanism of Desaturationâ€. Biochemistry, 1998, 37, 14664-14671.	1.2	218
146	Changes in the Regiospecificity of Aromatic Hydroxylation Produced by Active Site Engineering in the Diiron Enzyme Toluene 4-Monooxygenase. Biochemistry, 1997, 36, 9283-9289.	1.2	98
147	Recombinant Toluene-4-monooxygenase:  Catalytic and Mössbauer Studies of the Purified Diiron and Rieske Components of a Four-Protein Complex. Biochemistry, 1996, 35, 9106-9119.	1.2	180
148	Resonance Raman Evidence for an Fe-O-Fe Center in Stearoyl-ACP Desaturase. Primary Sequence Identity with Other Diiron-Oxo Proteins. Biochemistry, 1994, 33, 12776-12786.	1.2	206
149	Oxidation-reduction potentials of the methane monooxygenase hydroxylase component from Methylosinus trichosporium OB3b. Biochemistry, 1994, 33, 713-722.	1.2	119
150	Eight Histidine Residues Are Catalytically Essential in a Membrane-Associated Iron Enzyme, Stearoyl-CoA Desaturase, and Are Conserved in Alkane Hydroxylase and Xylene Monooxygenase. Biochemistry, 1994, 33, 12787-12794.	1.2	723
151	Spectroscopic studies of the coupled binuclear non-heme iron active site in the fully reduced hydroxylase component of methane monooxygenase: comparison to deoxy and deoxy-azide hemerythrin. Journal of the American Chemical Society, 1993, 115, 12409-12422.	6.6	96
152	Moessbauer, EPR, and ENDOR studies of the hydroxylase and reductase components of methane monooxygenase from Methylosinus trichosporium OB3b. Journal of the American Chemical Society, 1993, 115, 3688-3701.	6.6	185
153	A transient intermediate of the methane monooxygenase catalytic cycle containing an FelVFelV cluster. Journal of the American Chemical Society, 1993, 115, 6450-6451.	6.6	337
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