

Cheryl A Kerfeld

List of Publications by Year in descending order

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147
papers

14,232
citations

22153

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docs citations

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times ranked

11646
citing authors

#	ARTICLE	IF	CITATIONS
1	UV Excitation of Carotenoid Binding Proteins OCP and HCP: Excited-State Dynamics and Product Formation. <i>ChemPhotoChem</i> , 2022, 6, .	3.0	2
2	Toward a glycol radical enzyme containing synthetic bacterial microcompartment to produce pyruvate from formate and acetate. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	24
3	BMC Caller: a webtool to identify and analyze bacterial microcompartment types in sequence data. <i>Biology Direct</i> , 2022, 17, 9.	4.6	8
4	Atypical Carboxysome Loci: JEEPs or Junk?. <i>Frontiers in Microbiology</i> , 2022, 13, .	3.5	0
5	A genomic catalog of Earth's microbiomes. <i>Nature Biotechnology</i> , 2021, 39, 499-509.	17.5	457
6	Bioenergetics Theory and Components The Shells of Bacterial Microcompartments. , 2021, , 108-122.		0
7	Validation of an insertion-engineered isoprene synthase as a strategy to functionalize terpene synthases. <i>RSC Advances</i> , 2021, 11, 29997-30005.	3.6	1
8	A Survey of Bacterial Microcompartment Distribution in the Human Microbiome. <i>Frontiers in Microbiology</i> , 2021, 12, 669024.	3.5	5
9	Clues to the function of bacterial microcompartments from ancillary genes. <i>Biochemical Society Transactions</i> , 2021, 49, 1085-1098.	3.4	1
10	A catalog of the diversity and ubiquity of bacterial microcompartments. <i>Nature Communications</i> , 2021, 12, 3809.	12.8	55
11	Liposome-based measurement of light-driven chloride transport kinetics of halorhodopsin. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2021, 1863, 183637.	2.6	4
12	Evolutionary relationships among shell proteins of carboxysomes and metabolosomes. <i>Current Opinion in Microbiology</i> , 2021, 63, 1-9.	5.1	27
13	Integrated Structural Studies for Elucidating Carotenoid-Protein Interactions. <i>Advances in Experimental Medicine and Biology</i> , 2021, , .	1.6	1
14	Cyanobacterial carboxysomes contain an unique rubisco-activase-like protein. <i>New Phytologist</i> , 2020, 225, 793-806.	7.3	29
15	Redox Characterization of Electrode-Immobilized Bacterial Microcompartment Shell Proteins Engineered To Bind Metal Centers. <i>ACS Applied Bio Materials</i> , 2020, 3, 685-692.	4.6	9
16	Comparative ultrafast spectroscopy and structural analysis of OCP1 and OCP2 from <i>Tolypothrix</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2020, 1861, 148120.	1.0	22
17	Visualizing in Vivo Dynamics of Designer Nanoscaffolds. <i>Nano Letters</i> , 2020, 20, 208-217.	9.1	9
18	Structural analysis of a new carotenoid-binding protein: the C-terminal domain homolog of the OCP. <i>Scientific Reports</i> , 2020, 10, 15564.	3.3	18

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19	Engineered bacterial microcompartments: apps for programming metabolism. <i>Current Opinion in Biotechnology</i> , 2020, 65, 225-232.	6.6	20
20	Binding Options for the Small Subunit-Like Domain of Cyanobacteria to Rubisco. <i>Frontiers in Microbiology</i> , 2020, 11, 187.	3.5	2
21	Ubiquity and functional uniformity in CO ₂ concentrating mechanisms in multiple phyla of <i>Bacteria</i> is suggested by a diversity and prevalence of genes encoding candidate dissolved inorganic carbon transporters. <i>FEMS Microbiology Letters</i> , 2020, 367, .	1.8	10
22	Excited-State Properties of Canthaxanthin in Cyanobacterial Carotenoid-Binding Proteins HCP2 and HCP3. <i>Journal of Physical Chemistry B</i> , 2020, 124, 4896-4905.	2.6	14
23	Bacterial microcompartments: catalysis-enhancing metabolic modules for next generation metabolic and biomedical engineering. <i>BMC Biology</i> , 2019, 17, 79.	3.8	32
24	A designed bacterial microcompartment shell with tunable composition and precision cargo loading. <i>Metabolic Engineering</i> , 2019, 54, 286-291.	7.0	42
25	Structural Characterization of a Synthetic Tandem-Domain Bacterial Microcompartment Shell Protein Capable of Forming Icosahedral Shell Assemblies. <i>ACS Synthetic Biology</i> , 2019, 8, 668-674.	3.8	24
26	Structural and spectroscopic characterization of HCP2. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2019, 1860, 414-424.	1.0	21
27	Engineering the orange carotenoid protein for applications in synthetic biology. <i>Current Opinion in Structural Biology</i> , 2019, 57, 110-117.	5.7	14
28	X-ray radiolytic labeling reveals the molecular basis of orange carotenoid protein photoprotection and its interactions with fluorescence recovery protein. <i>Journal of Biological Chemistry</i> , 2019, 294, 8848-8860.	3.4	25
29	The Plasticity of Molecular Interactions Governs Bacterial Microcompartment Shell Assembly. <i>Structure</i> , 2019, 27, 749-763.e4.	3.3	50
30	Glycyl Radical Enzyme-Associated Microcompartments: Redox-Replete Bacterial Organelles. <i>MBio</i> , 2019, 10, .	4.1	30
31	Structure of a Synthetic β -Carboxysome Shell. <i>Plant Physiology</i> , 2019, 181, 1050-1058.	4.8	54
32	Heterohexamers Formed by CcmK3 and CcmK4 Increase the Complexity of Beta Carboxysome Shells. <i>Plant Physiology</i> , 2019, 179, 156-167.	4.8	61
33	Functionalization of Bacterial Microcompartment Shell Proteins With Covalently Attached Heme. <i>Frontiers in Bioengineering and Biotechnology</i> , 2019, 7, 432.	4.1	17
34	Characterization of Novel Homologs to the C-terminal Domain of the Orange Carotenoid Protein. <i>FASEB Journal</i> , 2019, 33, 779.45.	0.5	1
35	Genomes of ubiquitous marine and hypersaline <i>Hydrogenovibrio</i> , <i>Thiomicrothrix</i> and <i>Thiomicrospira</i> spp. encode a diversity of mechanisms to sustain chemolithoautotrophy in heterogeneous environments. <i>Environmental Microbiology</i> , 2018, 20, 2686-2708.	3.8	32
36	Bacterial microcompartments. <i>Nature Reviews Microbiology</i> , 2018, 16, 277-290.	28.6	328

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37	Fluorescence and Excited-State Conformational Dynamics of the Orange Carotenoid Protein. <i>Journal of Physical Chemistry B</i> , 2018, 122, 1792-1800.	2.6	14
38	Engineering nanoreactors using bacterial microcompartment architectures. <i>Current Opinion in Biotechnology</i> , 2018, 51, 1-7.	6.6	55
39	In Vitro Assembly of Diverse Bacterial Microcompartment Shell Architectures. <i>Nano Letters</i> , 2018, 18, 7030-7037.	9.1	61
40	Light-Driven Chloride Transport Kinetics of Halorhodopsin. <i>Biophysical Journal</i> , 2018, 115, 353-360.	0.5	9
41	Programmed loading and rapid purification of engineered bacterial microcompartment shells. <i>Nature Communications</i> , 2018, 9, 2881.	12.8	92
42	Structural and functional insights into the unique CBSâ€“CP12 fusion protein family in cyanobacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 7141-7146.	7.1	20
43	Free-electron laser data for multiple-particle fluctuation scattering analysis. <i>Scientific Data</i> , 2018, 5, 180201.	5.3	6
44	Structure and functions of Orange Carotenoid Protein homologs in cyanobacteria. <i>Current Opinion in Plant Biology</i> , 2017, 37, 1-9.	7.1	60
45	Î²-Carboxysome bioinformatics: identification and evolution of new bacterial microcompartment protein gene classes and core locus constraints. <i>Journal of Experimental Botany</i> , 2017, 68, 3841-3855.	4.8	36
46	Synthetic <scp>OCP</scp> heterodimers are photoactive and recapitulate the fusion of two primitive carotenoproteins in the evolution of cyanobacterial photoprotection. <i>Plant Journal</i> , 2017, 91, 646-656.	5.7	33
47	Connecting Earth observation to high-throughput biodiversity data. <i>Nature Ecology and Evolution</i> , 2017, 1, 176.	7.8	156
48	Assembly principles and structure of a 6.5-MDa bacterial microcompartment shell. <i>Science</i> , 2017, 356, 1293-1297.	12.6	187
49	In Vitro Characterization and Concerted Function of Three Core Enzymes of a Glycyl Radical Enzyme - Associated Bacterial Microcompartment. <i>Scientific Reports</i> , 2017, 7, 42757.	3.3	51
50	Structural and Functional Characterization of a Short-Chain Flavodoxin Associated with a Noncanonical 1,2-Propanediol Utilization Bacterial Microcompartment. <i>Biochemistry</i> , 2017, 56, 5679-5690.	2.5	4
51	Structure, function and evolution of the cyanobacterial orange carotenoid protein and its homologs. <i>New Phytologist</i> , 2017, 215, 937-951.	7.3	87
52	Carboxysomes: metabolic modules for CO2 fixation. <i>FEMS Microbiology Letters</i> , 2017, 364, .	1.8	86
53	A bioarchitectonic approach to the modular engineering of metabolism. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160387.	4.0	16
54	Raman Optical Activity Reveals Carotenoid Photoactivation Events in the Orange Carotenoid Protein in Solution. <i>Journal of the American Chemical Society</i> , 2017, 139, 10456-10460.	13.7	38

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55	Additional families of orange carotenoid proteins in the photoprotective system of cyanobacteria. <i>Nature Plants</i> , 2017, 3, 17089.	9.3	70
56	David W. Krogmann, 1931–2016. <i>Photosynthesis Research</i> , 2017, 132, 1-12.	2.9	7
57	Engineering the Bacterial Microcompartment Domain for Molecular Scaffolding Applications. <i>Frontiers in Microbiology</i> , 2017, 8, 1441.	3.5	57
58	Structure, Diversity, and Evolution of a New Family of Soluble Carotenoid-Binding Proteins in Cyanobacteria. <i>Molecular Plant</i> , 2016, 9, 1379-1394.	8.3	83
59	Cyanobacterial photoprotection by the orange carotenoid protein. <i>Nature Plants</i> , 2016, 2, 16180.	9.3	166
60	Different Functions of the Paralogs to the N-Terminal Domain of the Orange Carotenoid Protein in the Cyanobacterium <i>Anabaena</i> sp. PCC 7120. <i>Plant Physiology</i> , 2016, 171, 1852-1866.	4.8	76
61	Assembly, function and evolution of cyanobacterial carboxysomes. <i>Current Opinion in Plant Biology</i> , 2016, 31, 66-75.	7.1	197
62	Interrelated modules in cyanobacterial photosynthesis: the carbon-concentrating mechanism, photorespiration, and light perception. <i>Journal of Experimental Botany</i> , 2016, 67, 2931-2940.	4.8	19
63	Rewiring <i>Escherichia coli</i> for carbon-dioxide fixation. <i>Nature Biotechnology</i> , 2016, 34, 1035-1036.	17.5	12
64	Protein Nanotubes: Purification and Characterization of Protein Nanotubes Assembled from a Single Bacterial Microcompartment Shell Subunit (<i>Adv. Mater. Interfaces</i> 1/2016). <i>Advanced Materials Interfaces</i> , 2016, 3, .	3.7	0
65	Biochemical characterization of predicted Precambrian RuBisCO. <i>Nature Communications</i> , 2016, 7, 10382.	12.8	112
66	Bacterial microcompartments as metabolic modules for plant synthetic biology. <i>Plant Journal</i> , 2016, 87, 66-75.	5.7	32
67	Cyanobacterial ultrastructure in light of genomic sequence data. <i>Photosynthesis Research</i> , 2016, 129, 147-157.	2.9	42
68	Visualization of Bacterial Microcompartment Facet Assembly Using High-Speed Atomic Force Microscopy. <i>Nano Letters</i> , 2016, 16, 1590-1595.	9.1	106
69	Purification and Characterization of Protein Nanotubes Assembled from a Single Bacterial Microcompartment Shell Subunit. <i>Advanced Materials Interfaces</i> , 2016, 3, 1500295.	3.7	38
70	Structure and Function of a Bacterial Microcompartment Shell Protein Engineered to Bind a [4Fe-4S] Cluster. <i>Journal of the American Chemical Society</i> , 2016, 138, 5262-5270.	13.7	58
71	Production and Characterization of Synthetic Carboxysome Shells with Incorporated Luminal Proteins. <i>Plant Physiology</i> , 2016, 170, 1868-77.	4.8	64
72	The Structural Basis of Coenzyme A Recycling in a Bacterial Organelle. <i>PLoS Biology</i> , 2016, 14, e1002399.	5.6	40

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73	Structural Characterization of a Newly Identified Component of H^+ -Carboxysomes: The AAA+ Domain Protein CsoCbbQ. <i>Scientific Reports</i> , 2015, 5, 16243.	3.3	45
74	Plug-and-play for improving primary productivity. <i>American Journal of Botany</i> , 2015, 102, 1949-1950.	1.7	8
75	Advances in Understanding Carboxysome Assembly in <i>Prochlorococcus</i> and <i>Synechococcus</i> Implicate CsoS2 as a Critical Component. <i>Life</i> , 2015, 5, 1141-1171.	2.4	82
76	Bioinformatic analysis of the distribution of inorganic carbon transporters and prospective targets for bioengineering to increase C_i uptake by cyanobacteria. <i>Photosynthesis Research</i> , 2015, 126, 99-109.	2.9	18
77	A 12 Å... carotenoid translocation in a photoswitch associated with cyanobacterial photoprotection. <i>Science</i> , 2015, 348, 1463-1466.	12.6	192
78	Operational properties of fluctuation X-ray scattering data. <i>IUCr</i> , 2015, 2, 309-316.	2.2	21
79	Bioinformatic Characterization of Glycyl Radical Enzyme-Associated Bacterial Microcompartments. <i>Applied and Environmental Microbiology</i> , 2015, 81, 8315-8329.	3.1	59
80	Bacterial microcompartment assembly: The key role of encapsulation peptides. <i>Communicative and Integrative Biology</i> , 2015, 8, e1039755.	1.4	77
81	Streamlined Construction of the Cyanobacterial CO_2 -Fixing Organelle via Protein Domain Fusions for Use in Plant Synthetic Biology. <i>Plant Cell</i> , 2015, 27, 2637-2644.	6.6	49
82	Local and global structural drivers for the photoactivation of the orange carotenoid protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E5567-74.	7.1	121
83	Bacterial microcompartments and the modular construction of microbial metabolism. <i>Trends in Microbiology</i> , 2015, 23, 22-34.	7.7	165
84	Engineering Bacterial Microcompartment Shells: Chimeric Shell Proteins and Chimeric Carboxysome Shells. <i>ACS Synthetic Biology</i> , 2015, 4, 444-453.	3.8	88
85	Bayesian Analysis of Congruence of Core Genes in <i>Prochlorococcus</i> and <i>Synechococcus</i> and Implications on Horizontal Gene Transfer. <i>PLoS ONE</i> , 2014, 9, e85103.	2.5	12
86	Phylum-wide comparative genomics unravel the diversity of secondary metabolism in Cyanobacteria. <i>BMC Genomics</i> , 2014, 15, 977.	2.8	175
87	A Taxonomy of Bacterial Microcompartment Loci Constructed by a Novel Scoring Method. <i>PLoS Computational Biology</i> , 2014, 10, e1003898.	3.2	227
88	Introduction of a Synthetic CO_2 -fixing Photorespiratory Bypass into a Cyanobacterium. <i>Journal of Biological Chemistry</i> , 2014, 289, 9493-9500.	3.4	87
89	Structural and Functional Modularity of the Orange Carotenoid Protein: Distinct Roles for the N- and C-Terminal Domains in Cyanobacterial Photoprotection. <i>Plant Cell</i> , 2014, 26, 426-437.	6.6	114
90	Specificity of the Cyanobacterial Orange Carotenoid Protein: Influences of Orange Carotenoid Protein and Phycobilisome Structures. <i>Plant Physiology</i> , 2014, 164, 790-804.	4.8	30

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91	Characterization of a Planctomycetal Organelle: a Novel Bacterial Microcompartment for the Aerobic Degradation of Plant Saccharides. <i>Applied and Environmental Microbiology</i> , 2014, 80, 2193-2205.	3.1	124
92	Assembly of Robust Bacterial Microcompartment Shells Using Building Blocks from an Organelle of Unknown Function. <i>Journal of Molecular Biology</i> , 2014, 426, 2217-2228.	4.2	102
93	Dynamic cyanobacterial response to hydration and dehydration in a desert biological soil crust. <i>ISME Journal</i> , 2013, 7, 2178-2191.	9.8	217
94	Two new high-resolution crystal structures of carboxysome pentamer proteins reveal high structural conservation of CcmL orthologs among distantly related cyanobacterial species. <i>Photosynthesis Research</i> , 2013, 118, 9-16.	2.9	42
95	Biogenesis of a Bacterial Organelle: The Carboxysome Assembly Pathway. <i>Cell</i> , 2013, 155, 1131-1140.	28.9	274
96	Carotenoid-protein interaction alters the S1 energy of hydroxyechinenone in the Orange Carotenoid Protein. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2013, 1827, 248-254.	1.0	57
97	Structural, Mechanistic and Genomic Insights into OCP-Mediated Photoprotection. <i>Advances in Botanical Research</i> , 2013, 65, 1-26.	1.1	14
98	Cyanobacterial-based approaches to improving photosynthesis in plants. <i>Journal of Experimental Botany</i> , 2013, 64, 787-798.	4.8	121
99	Genome Mining Expands the Chemical Diversity of the Cyanobactin Family to Include Highly Modified Linear Peptides. <i>Chemistry and Biology</i> , 2013, 20, 1033-1043.	6.0	90
100	The Orange Carotenoid Protein: a blue-green light photoactive protein. <i>Photochemical and Photobiological Sciences</i> , 2013, 12, 1135-1143.	2.9	162
101	Comparative Analysis of 126 Cyanobacterial Genomes Reveals Evidence of Functional Diversity Among Homologs of the Redox-Regulated CP12 Protein. <i>Plant Physiology</i> , 2013, 161, 824-835.	4.8	47
102	Evidence for the widespread distribution of CRISPR-Cas system in the Phylum <i>Cyanobacteria</i> . <i>RNA Biology</i> , 2013, 10, 687-693.	3.1	86
103	Frontiers, Opportunities, and Challenges in Biochemical and Chemical Catalysis of CO ₂ Fixation. <i>Chemical Reviews</i> , 2013, 113, 6621-6658.	47.7	1,786
104	Improving the coverage of the cyanobacterial phylum using diversity-driven genome sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 1053-1058.	7.1	769
105	The crystal structures of the tri-functional <i>Chloroflexus aurantiacus</i> and bi-functional <i>Rhodobacter sphaeroides</i> malyl-CoA lyases and comparison with CitE-like superfamily enzymes and malate synthases. <i>BMC Structural Biology</i> , 2013, 13, 28.	2.3	11
106	The Structure of CcmP, a Tandem Bacterial Microcompartment Domain Protein from the $\hat{1}^2$ -Carboxysome, Forms a Subcompartment Within a Microcompartment. <i>Journal of Biological Chemistry</i> , 2013, 288, 16055-16063.	3.4	104
107	Crystal structure of the FRP and identification of the active site for modulation of OCP-mediated photoprotection in cyanobacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 10022-10027.	7.1	102
108	Isolation and Characterization of the <i>Prochlorococcus</i> Carboxysome Reveal the Presence of the Novel Shell Protein CsoS1D. <i>Journal of Bacteriology</i> , 2012, 194, 787-795.	2.2	67

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109	Elucidating Essential Role of Conserved Carboxysomal Protein CcmN Reveals Common Feature of Bacterial Microcompartment Assembly. <i>Journal of Biological Chemistry</i> , 2012, 287, 17729-17736.	3.4	140
110	The Essential Role of the N-Terminal Domain of the Orange Carotenoid Protein in Cyanobacterial Photoprotection: Importance of a Positive Charge for Phycobilisome Binding. <i>Plant Cell</i> , 2012, 24, 1972-1983.	6.6	82
111	The orange carotenoid protein in photoprotection of photosystem II in cyanobacteria. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 158-166.	1.0	171
112	Bioinformatic Identification and Structural Characterization of a New Carboxysome Shell Protein. <i>Advances in Photosynthesis and Respiration</i> , 2012, , 345-356.	1.0	6
113	Photoprotection in Cyanobacteria: The Orange Carotenoid Protein and Energy Dissipation. , 2011, , 395-421.		4
114	Excited-state properties of the 16 kDa red carotenoid protein from <i>Arthrospira maxima</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2011, 1807, 30-35.	1.0	36
115	Comparative analysis of carboxysome shell proteins. <i>Photosynthesis Research</i> , 2011, 109, 21-32.	2.9	112
116	Using BLAST to Teach "E-value-tionary" Concepts. <i>PLoS Biology</i> , 2011, 9, e1001014.	5.6	58
117	Carboxysomal carbonic anhydrases: Structure and role in microbial CO ₂ fixation. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2010, 1804, 382-392.	2.3	109
118	Incorporating Genomics and Bioinformatics across the Life Sciences Curriculum. <i>PLoS Biology</i> , 2010, 8, e1000448.	5.6	54
119	Bacterial Microcompartments. <i>Annual Review of Microbiology</i> , 2010, 64, 391-408.	7.3	299
120	Structural Determinants Underlying Photoprotection in the Photoactive Orange Carotenoid Protein of Cyanobacteria. <i>Journal of Biological Chemistry</i> , 2010, 285, 18364-18375.	3.4	152
121	Identification and Structural Analysis of a Novel Carboxysome Shell Protein with Implications for Metabolite Transport. <i>Journal of Molecular Biology</i> , 2009, 392, 319-333.	4.2	193
122	Protein-based organelles in bacteria: carboxysomes and related microcompartments. <i>Nature Reviews Microbiology</i> , 2008, 6, 681-691.	28.6	421
123	Atomic-Level Models of the Bacterial Carboxysome Shell. <i>Science</i> , 2008, 319, 1083-1086.	12.6	367
124	A photoactive carotenoid protein acting as light intensity sensor. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 12075-12080.	7.1	324
125	Structural Analysis of CsoS1A and the Protein Shell of the <i>Halothiobacillus neapolitanus</i> Carboxysome. <i>PLoS Biology</i> , 2007, 5, e144.	5.6	145
126	The Undergraduate Genomics Research Initiative. <i>PLoS Biology</i> , 2007, 5, e141.	5.6	23

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127	Light-Induced Energy Dissipation in Iron-Starved Cyanobacteria: Roles of OCP and IsiA Proteins. <i>Plant Cell</i> , 2007, 19, 656-672.	6.6	134
128	Structure of the RuBisCO chaperone RbcX from <i>Synechocystis</i> sp. PCC6803. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2007, 63, 1109-1112.	2.5	16
129	A Soluble Carotenoid Protein Involved in Phycobilisome-Related Energy Dissipation in Cyanobacteria. <i>Plant Cell</i> , 2006, 18, 992-1007.	6.6	396
130	The Genome of Deep-Sea Vent Chemolithoautotroph <i>Thiomicrospira crunogena</i> XCL-2. <i>PLoS Biology</i> , 2006, 4, e383.	5.6	144
131	The Structure of \hat{I}^2 -Carbonic Anhydrase from the Carboxysomal Shell Reveals a Distinct Subclass with One Active Site for the Price of Two. <i>Journal of Biological Chemistry</i> , 2006, 281, 7546-7555.	3.4	159
132	Protein Structures Forming the Shell of Primitive Bacterial Organelles. <i>Science</i> , 2005, 309, 936-938.	12.6	420
133	Spectroscopic Properties of the Carotenoid $3\hat{\epsilon}$ -Hydroxyechinenone in the Orange Carotenoid Protein from the Cyanobacterium <i>Arthrospira maxima</i> . <i>Biochemistry</i> , 2005, 44, 3994-4003.	2.5	124
134	Structure and Function of the Water-Soluble Carotenoid-Binding Proteins of Cyanobacteria. <i>Photosynthesis Research</i> , 2004, 81, 215-225.	2.9	73
135	Water-soluble carotenoid proteins of cyanobacteria. <i>Archives of Biochemistry and Biophysics</i> , 2004, 430, 2-9.	3.0	71
136	The 1.6Å resolution structure of Fe-superoxide dismutase from the thermophilic cyanobacterium <i>Thermosynechococcus elongatus</i> . <i>Journal of Biological Inorganic Chemistry</i> , 2003, 8, 707-714.	2.6	25
137	The Crystal Structure of a Cyanobacterial Water-Soluble Carotenoid Binding Protein. <i>Structure</i> , 2003, 11, 55-65.	3.3	267
138	Structural and EPR Characterization of the Soluble Form of Cytochrome c-550 and of the psbV2 Gene Product from the Cyanobacterium <i>Thermosynechococcus elongatus</i> . <i>Plant and Cell Physiology</i> , 2003, 44, 697-706.	3.1	39
139	Structure of cytochromec6 from <i>Arthrospira maxima</i> : an assembly of 24 subunits in a nearly symmetric shell. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2002, 58, 1104-1110.	2.5	4
140	Structures of Cytochrome c-549 and Cytochrome c6 from the Cyanobacterium <i>Arthrospira maxima</i> . <i>Biochemistry</i> , 2001, 40, 9215-9225.	2.5	65
141	Crystal Structure and Possible Dimerization of the High-Potential Iron-Sulfur Protein from <i>Chromatium purpuratum</i> . <i>Biochemistry</i> , 1998, 37, 13911-13917.	2.5	32
142	PHOTOSYNTHETIC CYTOCHROMESc IN CYANOBACTERIA, ALGAE, AND PLANTS. <i>Annual Review of Plant Biology</i> , 1998, 49, 397-425.	14.3	93
143	Structural comparison of cytochrome c2 and cytochrome c6. <i>Photosynthesis Research</i> , 1997, 54, 81-98.	2.9	8
144	Crystals of the Carotenoid Protein from <i>Arthrospira maxima</i> Containing Uniformly Oriented Pigment Molecules. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1997, 53, 720-723.	2.5	15

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145	Crystallization of two integral membrane pigment-protein complexes from the purple sulfur bacterium <i>Chromatium purpuratum</i> . <i>Protein Science</i> , 1993, 2, 1352-1355.	7.6	4
146	Light harvesting in photosystems I and II. <i>Biochemical Society Transactions</i> , 1993, 21, 15-18.	3.4	42
147	The use of non-denaturing Deriphat-polyacrylamide gel electrophoresis to fractionate pigment-protein complexes of purple bacteria. <i>Photosynthesis Research</i> , 1991, 30, 139-143.	2.9	8