

# Keith Moffat

## List of Publications by Year in descending order

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95  
papers

10,262  
citations

41344

49  
h-index

42399

92  
g-index

99  
all docs

99  
docs citations

99  
times ranked

6199  
citing authors

#	ARTICLE	IF	CITATIONS
1	Light-induced protein structural dynamics in bacteriophytochrome revealed by time-resolved x-ray solution scattering. <i>Science Advances</i> , 2022, 8, .	10.3	10
2	The primary structural photoresponse of phytochrome proteins captured by a femtosecond X-ray laser. <i>ELife</i> , 2020, 9, .	6.0	78
3	Laue diffraction and time-resolved crystallography: a personal history. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2019, 377, 20180243.	3.4	17
4	Structural basis for light control of cell development revealed by crystal structures of a myxobacterial phytochrome. <i>IUCr</i> , 2018, 5, 619-634.	2.2	33
5	Femtosecond structural photobiology. <i>Science</i> , 2018, 361, 127-128.	12.6	2
6	Small crystals, fast dynamics and noisy data are indeed beautiful. <i>IUCr</i> , 2017, 4, 303-305.	2.2	2
7	Femtosecond structural dynamics drives the trans/cis isomerization in photoactive yellow protein. <i>Science</i> , 2016, 352, 725-729.	12.6	348
8	Bacteriophytochrome Photoisomerization Proceeds Homogeneously Despite Heterogeneity in Ground State. <i>Biophysical Journal</i> , 2016, 111, 2125-2134.	0.5	21
9	The room temperature crystal structure of a bacterial phytochrome determined by serial femtosecond crystallography. <i>Scientific Reports</i> , 2016, 6, 35279.	3.3	39
10	Photocycle populations with femtosecond excitation of crystalline photoactive yellow protein. <i>Chemical Physics Letters</i> , 2016, 654, 63-71.	2.6	32
11	Structure of the response regulator RPA3017 involved in red-light signaling in <i>Rhodospseudomonas palustris</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2015, 71, 1215-1222.	0.8	5
12	Light Signaling Mechanism of Two Tandem Bacteriophytochromes. <i>Structure</i> , 2015, 23, 1179-1189.	3.3	42
13	Signal to noise considerations for single crystal femtosecond time resolved crystallography of the Photoactive Yellow Protein. <i>Faraday Discussions</i> , 2014, 171, 439-455.	3.2	19
14	Time-resolved serial crystallography captures high-resolution intermediates of photoactive yellow protein. <i>Science</i> , 2014, 346, 1242-1246.	12.6	418
15	Reply to 'Contradictions in X-ray structures of intermediates in the photocycle of photoactive yellow protein'. <i>Nature Chemistry</i> , 2014, 6, 259-260.	13.6	23
16	Time-resolved crystallography and protein design: signalling photoreceptors and optogenetics. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2014, 369, 20130568.	4.0	36
17	FTIR Spectroscopy Revealing Light-Dependent Refolding of the Conserved Tongue Region of Bacteriophytochrome. <i>Journal of Physical Chemistry Letters</i> , 2014, 5, 2512-2515.	4.6	49
18	Volume-conserving trans $\leftrightarrow$ cis isomerization pathways in photoactive yellow protein visualized by picosecond X-ray crystallography. <i>Nature Chemistry</i> , 2013, 5, 212-220.	13.6	178

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19	Resolution of structural heterogeneity in dynamic crystallography. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 946-959.	2.5	32
20	Pigment-Protein Interactions in Phytochromes Probed by Fluorescence Line Narrowing Spectroscopy. <i>Journal of Physical Chemistry B</i> , 2013, 117, 14940-14950.	2.6	7
21	Coiled-coil dimerization of the LOV2 domain of the blue-light photoreceptor phototropin 1 from <i>Arabidopsis thaliana</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013, 69, 1316-1321.	0.7	33
22	From Dusk till Dawn: One-Plasmid Systems for Light-Regulated Gene Expression. <i>Journal of Molecular Biology</i> , 2012, 416, 534-542.	4.2	207
23	Time-resolved structural studies at synchrotrons and X-ray free electron lasers: opportunities and challenges. <i>Current Opinion in Structural Biology</i> , 2012, 22, 651-659.	5.7	144
24	Crystal Structures of Aureochrome1 LOV Suggest New Design Strategies for Optogenetics. <i>Structure</i> , 2012, 20, 698-706.	3.3	67
25	Fluorescence quantum yield and photochemistry of bacteriophytochrome constructs. <i>Physical Chemistry Chemical Physics</i> , 2011, 13, 11985.	2.8	70
26	Primary Reactions of Bacteriophytochrome Observed with Ultrafast Mid-Infrared Spectroscopy. <i>Journal of Physical Chemistry A</i> , 2011, 115, 3778-3786.	2.5	43
27	Temperature-scan cryocrystallography reveals reaction intermediates in bacteriophytochrome. <i>Nature</i> , 2011, 479, 428-432.	27.8	155
28	Cluster Analysis of Time-Dependent Crystallographic Data: Direct Identification of Time-Independent Structural Intermediates. <i>Biophysical Journal</i> , 2011, 100, 440-449.	0.5	19
29	The Primary Photophysics of the <i>Avena sativa</i> Phototropin 1 LOV2 Domain Observed with Time-resolved Emission Spectroscopy. <i>Photochemistry and Photobiology</i> , 2011, 87, 534-541.	2.5	18
30	Structure and Function of Plant Photoreceptors. <i>Annual Review of Plant Biology</i> , 2010, 61, 21-47.	18.7	436
31	Picosecond Structural Dynamics at the Advanced Photon Source. <i>Synchrotron Radiation News</i> , 2010, 23, 18-25.	0.8	0
32	Proton-transfer and hydrogen-bond interactions determine fluorescence quantum yield and photochemical efficiency of bacteriophytochrome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 9170-9175.	7.1	132
33	Addition at the Molecular Level: Signal Integration in Designed Per-ARNT-Sim Receptor Proteins. <i>Journal of Molecular Biology</i> , 2010, 400, 477-486.	4.2	73
34	Engineered photoreceptors as novel optogenetic tools. <i>Photochemical and Photobiological Sciences</i> , 2010, 9, 1286-1300.	2.9	195
35	Conformational differences between the Pfr and Pr states in <i>Pseudomonas aeruginosa</i> bacteriophytochrome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 15639-15644.	7.1	133
36	Structure and Signaling Mechanism of Per-ARNT-Sim Domains. <i>Structure</i> , 2009, 17, 1282-1294.	3.3	457

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37	Design and Signaling Mechanism of Light-Regulated Histidine Kinases. <i>Journal of Molecular Biology</i> , 2009, 385, 1433-1444.	4.2	316
38	Changes in Quaternary Structure in the Signaling Mechanisms of PAS Domains. <i>Biochemistry</i> , 2008, 47, 12078-12086.	2.5	45
39	Crystal structure of <i>Pseudomonas aeruginosa</i> bacteriophytochrome: Photoconversion and signal transduction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 14715-14720.	7.1	293
40	Light-activated DNA binding in a designed allosteric protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 10709-10714.	7.1	273
41	Crystal structure of the chromophore binding domain of an unusual bacteriophytochrome, RpBphP3, reveals residues that modulate photoconversion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 12571-12576.	7.1	167
42	Structural Basis for Light-dependent Signaling in the Dimeric LOV Domain of the Photosensor YtvA. <i>Journal of Molecular Biology</i> , 2007, 373, 112-126.	4.2	211
43	Structure of the Redox Sensor Domain of <i>Azotobacter vinelandii</i> NifL at Atomic Resolution: Signaling, Dimerization, and Mechanism. <i>Biochemistry</i> , 2007, 46, 3614-3623.	2.5	103
44	N- and C-Terminal Flanking Regions Modulate Light-Induced Signal Transduction in the LOV2 Domain of the Blue Light Sensor Phototropin 1 from <i>Avena sativa</i> . <i>Biochemistry</i> , 2007, 46, 14001-14009.	2.5	283
45	Time-Resolved Crystallographic Studies of the Heme Domain of the Oxygen Sensor FixL: Structural Dynamics of Ligand Rebinding and Their Relation to Signal Transduction. <i>Biochemistry</i> , 2007, 46, 4706-4715.	2.5	45
46	Influence of the Crystalline State on Photoinduced Dynamics of Photoactive Yellow Protein Studied by Ultraviolet-Visible Transient Absorption Spectroscopy. <i>Biophysical Journal</i> , 2006, 90, 4224-4235.	0.5	52
47	Crystal Structures of the <i>Synechocystis</i> Photoreceptor Slr1694 Reveal Distinct Structural States Related to Signaling. <i>Biochemistry</i> , 2006, 45, 12687-12694.	2.5	140
48	A Structural Pathway for Signaling in the E46Q Mutant of Photoactive Yellow Protein. <i>Structure</i> , 2005, 13, 55-63.	3.3	73
49	From The Cover: Visualizing reaction pathways in photoactive yellow protein from nanoseconds to seconds. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 7145-7150.	7.1	256
50	Structure of a Novel Photoreceptor, the BLUF Domain of AppA from <i>Rhodobacter sphaeroides</i> . <i>Biochemistry</i> , 2005, 44, 7998-8005.	2.5	217
51	Crystal Structures of Deoxy and CO-Bound FixLH Reveal Details of Ligand Recognition and Signaling. <i>Biochemistry</i> , 2005, 44, 4627-4635.	2.5	78
52	Structural Heterogeneity of Cryotrapped Intermediates in the Bacterial Blue Light Photoreceptor, Photoactive Yellow Protein. <i>Photochemistry and Photobiology</i> , 2004, 80, 7.	2.5	26
53	Time-resolved crystallographic studies of light-induced structural changes in the photosynthetic reaction center. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 5982-5987.	7.1	65
54	Chromophore Conformation and the Evolution of Tertiary Structural Changes in Photoactive Yellow Protein. <i>Structure</i> , 2004, 12, 1039-1045.	3.3	65

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55	Analysis of experimental time-resolved crystallographic data by singular value decomposition. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004, 60, 860-871.	2.5	50
56	Short hydrogen bonds in photoactive yellow protein. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2004, 60, 1008-1016.	2.5	97
57	Protein kinetics: Structures of intermediates and reaction mechanism from time-resolved x-ray data. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 4799-4804.	7.1	88
58	The LOV2 Domain of Phototropin: A Reversible Photochromic Switch. <i>Journal of the American Chemical Society</i> , 2004, 126, 4512-4513.	13.7	102
59	Analytical trapping: extraction of time-independent structures from time-dependent crystallographic data. <i>Journal of Structural Biology</i> , 2004, 147, 211-222.	2.8	20
60	Structural Heterogeneity of Cryotrapped Intermediates in the Bacterial Blue Light Photoreceptor, Photoactive Yellow Protein. <i>Photochemistry and Photobiology</i> , 2004, 80, 7-14.	2.5	3
61	Purification and Initial Characterization of a Putative Blue Light Regulated Phosphodiesterase from <i>Escherichia coli</i> . <i>Photochemistry and Photobiology</i> , 2004, 80, 542-7.	2.5	23
62	The LOV Domain Family: Photoresponsive Signaling Modules Coupled to Diverse Output Domains. <i>Biochemistry</i> , 2003, 42, 2-10.	2.5	387
63	Application of Singular Value Decomposition to the Analysis of Time-Resolved Macromolecular X-Ray Data. <i>Biophysical Journal</i> , 2003, 84, 2112-2129.	0.5	146
64	Primary Reactions of the LOV2 Domain of Phototropin, a Plant Blue-Light Photoreceptor. <i>Biochemistry</i> , 2003, 42, 3385-3392.	2.5	214
65	The frontiers of time-resolved macromolecular crystallography: movies and chirped X-ray pulses. <i>Faraday Discussions</i> , 2003, 122, 65-77.	3.2	51
66	Crystal structure of a photoactive yellow protein from a sensor histidine kinase: Conformational variability and signal transduction. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 1649-1654.	7.1	39
67	Photoexcited Structure of a Plant Photoreceptor Domain Reveals a Light-Driven Molecular Switch. <i>Plant Cell</i> , 2002, 14, 1067-1075.	6.6	358
68	Protein Conformational Relaxation and Ligand Migration in Myoglobin: A Nanosecond to Millisecond Molecular Movie from Time-Resolved Laue X-ray Diffraction. <i>Biochemistry</i> , 2001, 40, 13802-13815.	2.5	329
69	A Molecular Movie at 1.8 Å... Resolution Displays the Photocycle of Photoactive Yellow Protein, a Eubacterial Blue-Light Receptor, from Nanoseconds to Seconds. <i>Biochemistry</i> , 2001, 40, 13788-13801.	2.5	190
70	Time-Resolved Biochemical Crystallography: A Mechanistic Perspective. <i>Chemical Reviews</i> , 2001, 101, 1569-1582.	47.7	180
71	Extraction of accurate structure-factor amplitudes from Laue data: wavelength normalization with wiggler and undulator X-ray sources. <i>Journal of Synchrotron Radiation</i> , 2000, 7, 236-244.	2.4	30
72	TIME-RESOLVED MACROMOLECULAR CRYSTALLOGRAPHY. , 2000, , .		0

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73	Laue crystallography: coming of age. <i>Journal of Synchrotron Radiation</i> , 1999, 6, 891-917.	2.4	122
74	Ultrafast time-resolved crystallography. <i>Nature Structural Biology</i> , 1998, 5, 641-643.	9.7	29
75	Structure Refinement Against Synchrotron Laue Data: Strategies for Data Collection and Reduction. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1998, 54, 367-377.	2.5	4
76	Energy Transduction on the Nanosecond Time Scale: Early Structural Events in a Xanthopsin Photocycle. <i>Science</i> , 1998, 279, 1946-1950.	12.6	302
77	[22] Laue diffraction. <i>Methods in Enzymology</i> , 1997, 277, 433-447.	1.0	35
78	Structure of a Protein Photocycle Intermediate by Millisecond Time-Resolved Crystallography. <i>Science</i> , 1997, 275, 1471-1475.	12.6	445
79	Synchrotron radiation applications to macromolecular crystallography. <i>Current Opinion in Structural Biology</i> , 1997, 7, 689-696.	5.7	20
80	Time-resolved structures of macromolecules at the ESRF: Single-pulse Laue diffraction, stroboscopic data collection and femtosecond flash photolysis. <i>Nuclear Instruments and Methods in Physics Research, Section A: Accelerators, Spectrometers, Detectors and Associated Equipment</i> , 1997, 398, 69-84.	1.6	85
81	Insights into specificity of cleavage and mechanism of cell entry from the crystal structure of the highly specific <i>Aspergillus</i> ribotoxin, restrictocin. <i>Structure</i> , 1996, 4, 837-852.	3.3	89
82	Optical Studies of a Bacterial Photoreceptor Protein, Photoactive Yellow Protein, in Single Crystals. <i>Biochemistry</i> , 1995, 34, 879-890.	2.5	75
83	Freeze trapping of reaction intermediates. <i>Current Opinion in Structural Biology</i> , 1995, 5, 656-663.	5.7	76
84	Optical monitoring of protein crystals in time-resolved X-ray experiments: Microspectrophotometer design and performance. <i>Review of Scientific Instruments</i> , 1994, 65, 1506-1511.	1.3	29
85	Structure of isothiocyanate methemoglobin. <i>Journal of Molecular Biology</i> , 1981, 145, 815-824.	4.2	19
86	Structure of imidazole methemoglobin. <i>Journal of Molecular Biology</i> , 1981, 147, 325-335.	4.2	19
87	Structure of nitric oxide hemoglobin. <i>Journal of Molecular Biology</i> , 1979, 134, 401-417.	4.2	75
88	Structure of azide methemoglobin. <i>Journal of Molecular Biology</i> , 1979, 134, 419-429.	4.2	24
89	Structure of hemoglobin reconstituted with mesoheme. <i>Journal of Molecular Biology</i> , 1977, 113, 419-430.	4.2	19
90	The structure of metmanganoglobin. <i>Journal of Molecular Biology</i> , 1976, 104, 669-685.	4.2	28

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91	Structure of cyanide methemoglobin. Journal of Molecular Biology, 1976, 104, 687-706.	4.2	103
92	Structure of fluoride methemoglobin. Journal of Molecular Biology, 1976, 104, 723-728.	4.2	41
93	The structure of hemoglobin reconstituted with deuteroheme. Journal of Molecular Biology, 1976, 106, 895-902.	4.2	18
94	Crystallographic studies on manganese hemoglobin. Journal of the American Chemical Society, 1974, 96, 5259-5261.	13.7	14
95	Femtosecond Studies of the Initial Events in the Photocycle of Photoactive Yellow Protein (PYP). , 0, , 381-390.		3