Andreas Winkler

List of Publications by Year in descending order

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36 papers

1,632 citations

331670 21 h-index 36 g-index

38 all docs 38 docs citations

38 times ranked

1944 citing authors

#	Article	IF	CITATIONS
1	Characterisation of sequence–structure–function space in sensor–effector integrators of phytochrome-regulated diguanylate cyclases. Photochemical and Photobiological Sciences, 2022, 21, 1761-1779.	2.9	3
2	The Trimeric Major Capsid Protein of Mavirus is stabilized by its Interlocked N-termini Enabling Core Flexibility for Capsid Assembly. Journal of Molecular Biology, 2021, 433, 166859.	4.2	5
3	The PHY Domain Dimer Interface of Bacteriophytochromes Mediates Cross-talk between Photosensory Modules and Output Domains. Journal of Molecular Biology, 2021, 433, 167092.	4.2	3
4	The scope of flavin-dependent reactions and processes in the model plant Arabidopsis thaliana. Phytochemistry, 2021, 189, 112822.	2.9	18
5	A small molecule chaperone rescues the stability and activity of a cancerâ€nssociated variant of NAD(P)H:quinone oxidoreductase 1 <i>inÂvitro</i> . FEBS Letters, 2020, 594, 424-438.	2.8	7
6	Distinct chromophore–protein environments enable asymmetric activation of a bacteriophytochrome-activated diguanylate cyclase. Journal of Biological Chemistry, 2020, 295, 539-551.	3.4	14
7	Influence of the N-terminal segment and the PHY-tongue element on light-regulation in bacteriophytochromes. Journal of Biological Chemistry, 2019, 294, 4498-4510.	3.4	15
8	Bacteriophytochromes – from informative model systems of phytochrome function to powerful tools in cell biology. Current Opinion in Structural Biology, 2019, 57, 72-83.	5.7	44
9	The single berberine bridge enzyme homolog of <i>PhyscomitrellaÂpatens</i> is a cellobiose oxidase. FEBS Journal, 2018, 285, 1923-1943.	4.7	17
10	Oxidation of the FAD cofactor to the 8-formyl-derivative in human electron-transferring flavoprotein. Journal of Biological Chemistry, 2018, 293, 2829-2840.	3.4	18
11	Structure-guided design and functional characterization of an artificial red light–regulated guanylate/adenylate cyclase for optogenetic applications. Journal of Biological Chemistry, 2018, 293, 9078-9089.	3.4	45
12	Molecular Mechanisms of Bacterial Bioluminescence. Computational and Structural Biotechnology Journal, 2018, 16, 551-564.	4.1	108
13	Asymmetric activation mechanism of a homodimeric red light-regulated photoreceptor. ELife, 2018, 7, .	6.0	46
14	Long-range allosteric signaling in red light–regulated diguanylyl cyclases. Science Advances, 2017, 3, e1602498.	10.3	87
15	Photoactivation Mechanism of a Bacterial Light-Regulated Adenylyl Cyclase. Journal of Molecular Biology, 2017, 429, 1336-1351.	4.2	45
16	The family of berberine bridge enzyme-like enzymes: A treasure-trove of oxidative reactions. Archives of Biochemistry and Biophysics, 2017, 632, 88-103.	3.0	100
17	LOVTRAP: an optogenetic system for photoinduced protein dissociation. Nature Methods, 2016, 13, 755-758.	19.0	267
18	Applications of hydrogen deuterium exchange (HDX) for the characterization of conformational dynamics in light-activated photoreceptors. Frontiers in Molecular Biosciences, 2015, 2, 33.	3.5	10

#	Article	IF	CITATIONS
19	Structural and biochemical properties of LuxF from Photobacterium leiognathi. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2015, 1854, 1466-1475.	2.3	13
20	Rapid access to glycopeptide antibiotic precursor peptides coupled with cytochrome P450-mediated catalysis: towards a biomimetic synthesis of glycopeptide antibiotics. Organic and Biomolecular Chemistry, 2015, 13, 2012-2021.	2.8	30
21	Structural Details of Light Activation of the LOV2-based Photoswitch PA-Rac1. ACS Chemical Biology, 2015, 10, 502-509.	3.4	20
22	Multi-PAS domain-mediated protein oligomerization of PpsR from <i>Rhodobacter sphaeroides</i> Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 863-876.	2.5	11
23	Rescuing DNA repair activity by rewiring the H-atom transfer pathway in the radical SAM enzyme, spore photoproduct lyase. Chemical Communications, 2014, 50, 14201-14204.	4.1	16
24	Hexicon 2: Automated Processing of Hydrogen-Deuterium Exchange Mass Spectrometry Data with Improved Deuteration Distribution Estimation. Journal of the American Society for Mass Spectrometry, 2014, 25, 1018-1028.	2.8	43
25	Characterization of Elements Involved in Allosteric Light Regulation of Phosphodiesterase Activity by Comparison of Different Functional BlrP1 States. Journal of Molecular Biology, 2014, 426, 853-868.	4.2	28
26	A ternary AppA–PpsR–DNA complex mediates light regulation of photosynthesis-related gene expression. Nature Structural and Molecular Biology, 2013, 20, 859-867.	8.2	62
27	Catalytic and Structural Role of a Conserved Active Site Histidine in Berberine Bridge Enzyme. Biochemistry, 2012, 51, 6139-6147.	2.5	14
28	Old Yellow Enzymeâ€Catalyzed Dehydrogenation of Saturated Ketones. Advanced Synthesis and Catalysis, 2011, 353, 268-274.	4.3	54
29	Biocatalytic Enantioselective Oxidative CC Coupling by Aerobic CH Activation. Angewandte Chemie - International Edition, 2011, 50, 1068-1071.	13.8	72
30	Cover Picture: Biocatalytic Enantioselective Oxidative CC Coupling by Aerobic CH Activation (Angew. Chem. Int. Ed. 5/2011). Angewandte Chemie - International Edition, 2011, 50, 967-967.	13.8	0
31	Structural and Mechanistic Studies Reveal the Functional Role of Bicovalent Flavinylation in Berberine Bridge Enzyme. Journal of Biological Chemistry, 2009, 284, 19993-20001.	3.4	43
32	Berberine bridge enzyme catalyzes the six electron oxidation of (S)-reticuline to dehydroscoulerine. Phytochemistry, 2009, 70, 1092-1097.	2.9	28
33	A concerted mechanism for berberine bridge enzyme. Nature Chemical Biology, 2008, 4, 739-741.	8.0	127
34	Functional Roles of the 6-S-Cysteinyl, 8α-N1-Histidyl FAD in Glucooligosaccharide Oxidase from Acremonium strictum. Journal of Biological Chemistry, 2008, 283, 30990-30996.	3.4	32
35	6-S-Cysteinylation of Bi-covalently Attached FAD in Berberine Bridge Enzyme Tunes the Redox Potential for Optimal Activity. Journal of Biological Chemistry, 2007, 282, 24437-24443.	3.4	52
36	Biochemical Evidence That Berberine Bridge Enzyme Belongs to a Novel Family of Flavoproteins Containing a Bi-covalently Attached FAD Cofactor. Journal of Biological Chemistry, 2006, 281, 21276-21285.	3.4	107

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