

Zhao-Xun Liang

List of Publications by Year in descending order

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78
papers

3,402
citations

117453

34
h-index

155451

55
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84
all docs

84
docs citations

84
times ranked

3274
citing authors

#	ARTICLE	IF	CITATIONS
1	YybT Is a Signaling Protein That Contains a Cyclic Dinucleotide Phosphodiesterase Domain and a GGDEF Domain with ATPase Activity. <i>Journal of Biological Chemistry</i> , 2010, 285, 473-482.	1.6	231
2	Catalytic Mechanism of Cyclic Di-GMP-Specific Phosphodiesterase: a Study of the EAL Domain-Containing RocR from <i>Pseudomonas aeruginosa</i> . <i>Journal of Bacteriology</i> , 2008, 190, 3622-3631.	1.0	224
3	MrkH, a Novel c-di-GMP-Dependent Transcriptional Activator, Controls <i>Klebsiella pneumoniae</i> Biofilm Formation by Regulating Type 3 Fimbriae Expression. <i>PLoS Pathogens</i> , 2011, 7, e1002204.	2.1	195
4	Thermal-activated protein mobility and its correlation with catalysis in thermophilic alcohol dehydrogenase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 9556-9561.	3.3	134
5	Structural bases of hydrogen tunneling in enzymes: progress and puzzles. <i>Current Opinion in Structural Biology</i> , 2004, 14, 648-655.	2.6	104
6	The Functional Role of a Conserved Loop in EAL Domain-Based Cyclic di-GMP-Specific Phosphodiesterase. <i>Journal of Bacteriology</i> , 2009, 191, 4722-4731.	1.0	100
7	Dynamic Docking and Electron Transfer between Zn-myoglobin and Cytochrome b5. <i>Journal of the American Chemical Society</i> , 2002, 124, 6849-6859.	6.6	98
8	Dynamic Docking and Electron-Transfer between Cytochrome b5 and a Suite of Myoglobin Surface-Charge Mutants. Introduction of a Functional-Docking Algorithm for Protein-Protein Complexes. <i>Journal of the American Chemical Society</i> , 2004, 126, 2785-2798.	6.6	88
9	Enzymatic synthesis of c-di-GMP using a thermophilic diguanylate cyclase. <i>Analytical Biochemistry</i> , 2009, 389, 138-142.	1.1	83
10	The expanding roles of c-di-GMP in the biosynthesis of exopolysaccharides and secondary metabolites. <i>Natural Product Reports</i> , 2015, 32, 663-683.	5.2	81
11	A Flavin Cofactor-Binding PAS Domain Regulates c-di-GMP Synthesis in <i>DGC2</i> from <i>Acetobacter xylinum</i> . <i>Biochemistry</i> , 2009, 48, 10275-10285.	1.2	79
12	Binding of Cyclic Diguanylate in the Non-catalytic EAL Domain of FimX Induces a Long-range Conformational Change. <i>Journal of Biological Chemistry</i> , 2011, 286, 2910-2917.	1.6	73
13	Crystal Structure and Amide H/D Exchange of Binary Complexes of Alcohol Dehydrogenase from <i>Bacillus stearothermophilus</i> : An Insight into Thermostability and Cofactor Binding. <i>Biochemistry</i> , 2004, 43, 5266-5277.	1.2	69
14	Multifunctional Antimicrobial Nanofiber Dressings Containing μ -Polylysine for the Eradication of Bacterial Bioburden and Promotion of Wound Healing in Critically Colonized Wounds. <i>ACS Applied Materials & Interfaces</i> , 2020, 12, 15989-16005.	4.0	69
15	Complexity and simplicity in the biosynthesis of enediyne natural products. <i>Natural Product Reports</i> , 2010, 27, 499.	5.2	68
16	Evidence for Increased Local Flexibility in Psychrophilic Alcohol Dehydrogenase Relative to Its Thermophilic Homologue. <i>Biochemistry</i> , 2004, 43, 14676-14683.	1.2	62
17	A cyclic di-GMP binding adaptor protein interacts with a chemotaxis methyltransferase to control flagellar motor switching. <i>Science Signaling</i> , 2016, 9, ra102.	1.6	61
18	Cyclic di-AMP synthesis by the diadenylate cyclase <i>CdaA</i> is modulated by the peptidoglycan biosynthesis enzyme <i>GlmM</i> in <i>actococcus lactis</i> . <i>Molecular Microbiology</i> , 2016, 99, 1015-1027.	1.2	61

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19	Enhanced uptake of potassium or glycine betaine or export of cyclic-di-AMP restores osmoresistance in a high cyclic-di-AMP <i>Lactococcus lactis</i> mutant. <i>PLoS Genetics</i> , 2018, 14, e1007574.	1.5	61
20	Unusual Heme-Binding PAS Domain from YybT Family Proteins. <i>Journal of Bacteriology</i> , 2011, 193, 1543-1551.	1.0	60
21	Sequencing and functional annotation of the whole genome of the filamentous fungus <i>Aspergillus westerdijkiae</i> . <i>BMC Genomics</i> , 2016, 17, 633.	1.2	58
22	Structural Insights into the Regulatory Mechanism of the Response Regulator RocR from <i>Pseudomonas aeruginosa</i> in Cyclic Di-GMP Signaling. <i>Journal of Bacteriology</i> , 2012, 194, 4837-4846.	1.0	57
23	Synthesis of (<i>R</i>)-Mellein by a Partially Reducing Iterative Polyketide Synthase. <i>Journal of the American Chemical Society</i> , 2012, 134, 11924-11927.	6.6	49
24	The structure and inhibition of a GGDEF diguanylate cyclase complexed with (c-di-GMP) ₂ at the active site. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2011, 67, 997-1008.	2.5	48
25	Structural Insights into the Distinct Binding Mode of Cyclic Di-AMP with <i>Sa</i> CpaA_RCK. <i>Biochemistry</i> , 2015, 54, 4936-4951.	1.2	48
26	Impact of Protein Flexibility on Hydride-Transfer Parameters in Thermophilic and Psychrophilic Alcohol Dehydrogenases. <i>Journal of the American Chemical Society</i> , 2004, 126, 9500-9501.	6.6	47
27	Identification of a biosynthetic gene cluster for the polyene macrolactam sceliphrolactam in a <i>Streptomyces</i> strain isolated from mangrove sediment. <i>Scientific Reports</i> , 2018, 8, 1594.	1.6	46
28	Structure and Catalytic Mechanism of the Thioesterase CalE7 in Eneidyne Biosynthesis. <i>Journal of Biological Chemistry</i> , 2009, 284, 15739-15749.	1.6	42
29	Progress in Understanding the Molecular Basis Underlying Functional Diversification of Cyclic Dinucleotide Turnover Proteins. <i>Journal of Bacteriology</i> , 2017, 199, .	1.0	41
30	Electrostatic Control of Electron Transfer between Myoglobin and Cytochrome b5: Effect of Methylating the Heme Propionates of Zn-Myoglobin. <i>Journal of the American Chemical Society</i> , 2000, 122, 3552-3553.	6.6	40
31	A Cyclic di-GMP-binding Adaptor Protein Interacts with Histidine Kinase to Regulate Two-component Signaling. <i>Journal of Biological Chemistry</i> , 2016, 291, 16112-16123.	1.6	40
32	Crystal Structure of the Acyltransferase Domain of the Iterative Polyketide Synthase in Eneidyne Biosynthesis. <i>Journal of Biological Chemistry</i> , 2012, 287, 23203-23215.	1.6	38
33	The <i>EAL</i> -like protein <i>STM</i> 1697 regulates virulence phenotypes, motility and biofilm formation in <i>Salmonella typhimurium</i> . <i>Molecular Microbiology</i> , 2013, 90, 1216-1232.	1.2	38
34	Characterization of a Carbonyl-Conjugated Polyene Precursor in 10-Membered Eneidyne Biosynthesis. <i>Journal of the American Chemical Society</i> , 2008, 130, 8142-8143.	6.6	37
35	Visualizing the Perturbation of Cellular Cyclic di-GMP Levels in Bacterial Cells. <i>Journal of the American Chemical Society</i> , 2013, 135, 566-569.	6.6	37
36	Insights into Biofilm Dispersal Regulation from the Crystal Structure of the PAS-GGDEF-EAL Region of RbdA from <i>Pseudomonas aeruginosa</i> . <i>Journal of Bacteriology</i> , 2018, 200, .	1.0	37

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37	Novel c-di-GMP recognition modes of the mouse innate immune adaptor protein STING. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 352-366.	2.5	36
38	Photoinduced Electron Transfer in a Supramolecular Species Building of Mono-6-p-nitrobenzoyl- β -cyclodextrin with Naphthalene Derivatives. <i>Journal of Organic Chemistry</i> , 2002, 67, 2429-2434.	1.7	31
39	Replenishing the cyclic-di-AMP pool: regulation of diadenylate cyclase activity in bacteria. <i>Current Genetics</i> , 2016, 62, 731-738.	0.8	31
40	Biosynthesis of Tasikamides via Pathway Coupling and Diazonium-Mediated Hydrazone Formation. <i>Journal of the American Chemical Society</i> , 2022, 144, 1622-1633.	6.6	31
41	Dynamic docking and electron transfer between myoglobin and cytochrome b 5. <i>Journal of Biological Inorganic Chemistry</i> , 2002, 7, 580-588.	1.1	30
42	Products of the iterative polyketide synthases in 9- and 10-membered enediyne biosynthesis. <i>Chemical Communications</i> , 2009, , 7399.	2.2	30
43	Insight into Enzymatic Nitrile Reduction: QM/MM Study of the Catalytic Mechanism of QueF Nitrile Reductase. <i>ACS Catalysis</i> , 2015, 5, 3740-3751.	5.5	28
44	Solution Structure of the PAS Domain of a Thermophilic YybT Protein Homolog Reveals a Potential Ligand-binding Site. <i>Journal of Biological Chemistry</i> , 2013, 288, 11949-11959.	1.6	27
45	Structural analyses unravel the molecular mechanism of cyclic di-GMP regulation of bacterial chemotaxis via a PilZ adaptor protein. <i>Journal of Biological Chemistry</i> , 2018, 293, 100-111.	1.6	25
46	Solution Structures of the Acyl Carrier Protein Domain from the Highly Reducing Type I Iterative Polyketide Synthase CalE8. <i>PLoS ONE</i> , 2011, 6, e20549.	1.1	25
47	Evidence for a novel phosphopantetheinyl transferase domain in the polyketide synthase for enediyne biosynthesis. <i>FEBS Letters</i> , 2008, 582, 1097-1103.	1.3	24
48	Sungeidines from a Non-canonical Enediyne Biosynthetic Pathway. <i>Journal of the American Chemical Society</i> , 2020, 142, 1673-1679.	6.6	24
49	Expression, purification and characterization of the acyl carrier protein phosphodiesterase from <i>Pseudomonas Aeruginosa</i> . <i>Protein Expression and Purification</i> , 2010, 71, 132-138.	0.6	23
50	Nitrile reductase as a biocatalyst: opportunities and challenges. <i>Catalysis Science and Technology</i> , 2014, 4, 2871-2876.	2.1	23
51	Functional Divergence of FimX in PilZ Binding and Type IV Pilus Regulation. <i>Journal of Bacteriology</i> , 2012, 194, 5922-5931.	1.0	22
52	Structure of a Diguanylate Cyclase from <i>Thermotoga maritima</i> : Insights into Activation, Feedback Inhibition and Thermostability. <i>PLoS ONE</i> , 2014, 9, e110912.	1.1	22
53	Emerging paradigms for PilZ domain-mediated C-di-GMP signaling. <i>Biochemical Society Transactions</i> , 2019, 47, 381-388.	1.6	22
54	Induced-fit upon Ligand Binding Revealed by Crystal Structures of the Hot-dog Fold Thioesterase in Dynemicin Biosynthesis. <i>Journal of Molecular Biology</i> , 2010, 404, 291-306.	2.0	21

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55	Regulation of flagellar motor switching by c-di-GMP phosphodiesterases in <i>Pseudomonas aeruginosa</i> . <i>Journal of Biological Chemistry</i> , 2019, 294, 13789-13799.	1.6	20
56	Sol-gel immobilization of a thermophilic diguanylate cyclase for enzymatic production of cyclic-di-GMP. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2010, 67, 98-103.	1.8	19
57	Integrated Genomic and Metabolomic Approach to the Discovery of Potential Anti-Quorum Sensing Natural Products from Microbes Associated with Marine Samples from Singapore. <i>Marine Drugs</i> , 2019, 17, 72.	2.2	16
58	Discovery, biosynthesis and antifungal mechanism of the polyene-polyol meijiemycin. <i>Chemical Communications</i> , 2020, 56, 822-825.	2.2	16
59	A study on the laser flash photolysis of phenothiazine and its N-alkyl derivatives. <i>Journal of Photochemistry and Photobiology A: Chemistry</i> , 1996, 93, 27-31.	2.0	15
60	MS/MS-Based Molecular Networking Approach for the Detection of Aplysiatoxin-Related Compounds in Environmental Marine Cyanobacteria. <i>Marine Drugs</i> , 2018, 16, 505.	2.2	14
61	Pathway Retrofitting Yields Insights into the Biosynthesis of Anthraquinone-Fused Enediynes. <i>Journal of the American Chemical Society</i> , 2021, 143, 11500-11509.	6.6	13
62	Dynamic swimming pattern of <i>Pseudomonas aeruginosa</i> near a vertical wall during initial attachment stages of biofilm formation. <i>Scientific Reports</i> , 2021, 11, 1952.	1.6	12
63	Rigidifying Acyl Carrier Protein Domain in Iterative Type I PKS CalE8 Does Not Affect Its Function. <i>Biophysical Journal</i> , 2012, 103, 1037-1044.	0.2	10
64	Complete Genome Sequence of the Filamentous Fungus <i>Aspergillus westerdijkiae</i> Reveals the Putative Biosynthetic Gene Cluster of Ochratoxin A. <i>Genome Announcements</i> , 2016, 4, .	0.8	10
65	Cyclic di-AMP Oversight of Counter-Ion Osmolyte Pools Impacts Intrinsic Cefuroxime Resistance in <i>Lactococcus lactis</i> . <i>MBio</i> , 2021, 12, .	1.8	10
66	Crystallization studies of the murine c-di-GMP sensor protein STING. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 906-910.	0.7	9
67	Insights into the programmed ketoreduction of partially reducing polyketide synthases: stereo- and substrate-specificity of the ketoreductase domain. <i>Organic and Biomolecular Chemistry</i> , 2014, 12, 8542-8549.	1.5	9
68	The MapZ-Mediated Methylation of Chemoreceptors Contributes to Pathogenicity of <i>Pseudomonas aeruginosa</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 67.	1.5	8
69	Taming the flagellar motor of pseudomonads with a nucleotide messenger. <i>Environmental Microbiology</i> , 2020, 22, 2496-2513.	1.8	8
70	Finally! The structural secrets of a c-di-GMP phosphodiesterase revealed. <i>Molecular Microbiology</i> , 2014, 91, 1-5.	1.2	6
71	Rational Design of Fluorescent Biosensor for Cyclic di-GMP. <i>ChemBioChem</i> , 2011, 12, 2753-2758.	1.3	5
72	Crystallization and preliminary X-ray diffraction studies of Xanthomonas campestris PNPase in the presence of c-di-GMP. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 1247-1250.	0.7	5

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73	Crystallization and preliminary X-ray diffraction characterization of the XccFimXEAAL ^c -di-GMP and XccFimXEAAL ^c -di-GMP ^a -XccPilZ complexes from <i>Xanthomonas campestris</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 301-305.	0.7	5
74	Expression, purification and preliminary crystallographic analysis of <i>Pseudomonas aeruginosa</i> RocR protein. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009, 65, 1035-1038.	0.7	4
75	Enzymatic Production of c-di-GMP Using a Thermophilic Diguanylate Cyclase. <i>Methods in Molecular Biology</i> , 2017, 1657, 11-22.	0.4	4
76	Draft Genome Sequence of <i>Nocardia jinanensis</i> , an Opportunistic Bacterial Pathogen That Causes Cellulitis. <i>Genome Announcements</i> , 2016, 4, .	0.8	2
77	Osmoregulation via Cyclic di-AMP Signaling. , 2020, , 177-189.		1
78	YybT is a signaling protein that contains a cyclic dinucleotide phosphodiesterase domain and a GGDEF domain with ATPase activity.. <i>Journal of Biological Chemistry</i> , 2011, 286, 29441.	1.6	0