

# Peter J Tonge

## List of Publications by Year in descending order

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220  
papers

9,538  
citations

30070

54  
h-index

51608

86  
g-index

231  
all docs

231  
docs citations

231  
times ranked

8472  
citing authors

#	ARTICLE	IF	CITATIONS
1	Drugâ€‘target residence time: critical information for lead optimization. <i>Current Opinion in Chemical Biology</i> , 2010, 14, 467-474.	6.1	391
2	The isoniazid-NAD adduct is a slow, tight-binding inhibitor of InhA, the <i>Mycobacterium tuberculosis</i> enoyl reductase: Adduct affinity and drug resistance. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 13881-13886.	7.1	298
3	Drug Discovery Using Chemical Systems Biology: Repositioning the Safe Medicine Comtan to Treat Multi-Drug and Extensively Drug Resistant Tuberculosis. <i>PLoS Computational Biology</i> , 2009, 5, e1000423.	3.2	283
4	Inhibitors of FabI, an Enzyme Drug Target in the Bacterial Fatty Acid Biosynthesis Pathway. <i>Accounts of Chemical Research</i> , 2008, 41, 11-20.	15.6	246
5	High Affinity InhA Inhibitors with Activity against Drug-Resistant Strains of <i>Mycobacterium tuberculosis</i> . <i>ACS Chemical Biology</i> , 2006, 1, 43-53.	3.4	234
6	Inhibition of InhA, the Enoyl Reductase from <i>Mycobacterium tuberculosis</i> , by Triclosan and Isoniazidâ€‘. <i>Biochemistry</i> , 2000, 39, 7645-7650.	2.5	226
7	Structural basis and mechanism of enoyl reductase inhibition by triclosan. <i>Journal of Molecular Biology</i> , 1999, 290, 859-865.	4.2	201
8	Observation of Excited-State Proton Transfer in Green Fluorescent Protein using Ultrafast Vibrational Spectroscopy. <i>Journal of the American Chemical Society</i> , 2005, 127, 2864-2865.	13.7	189
9	Drugâ€‘Target Kinetics in Drug Discovery. <i>ACS Chemical Neuroscience</i> , 2018, 9, 29-39.	3.5	189
10	A Machine Learning-Based Method To Improve Docking Scoring Functions and Its Application to Drug Repurposing. <i>Journal of Chemical Information and Modeling</i> , 2011, 51, 408-419.	5.4	175
11	Probing the Ground State Structure of the Green Fluorescent Protein Chromophore Using Raman Spectroscopyâ€‘. <i>Biochemistry</i> , 2000, 39, 4423-4431.	2.5	161
12	A Slow, Tight Binding Inhibitor of InhA, the Enoyl-Acyl Carrier Protein Reductase from <i>Mycobacterium tuberculosis</i> . <i>Journal of Biological Chemistry</i> , 2010, 285, 14330-14337.	3.4	155
13	Novel Trisubstituted Benzimidazoles, Targeting <i>Mtb</i> FtsZ, as a New Class of Antitubercular Agents. <i>Journal of Medicinal Chemistry</i> , 2011, 54, 374-381.	6.4	145
14	Marine natural products from the Turkish sponge <i>Agelas oroides</i> that inhibit the enoyl reductases from <i>Plasmodium falciparum</i> , <i>Mycobacterium tuberculosis</i> and <i>Escherichia coli</i> . <i>Bioorganic and Medicinal Chemistry</i> , 2007, 15, 6834-6845.	3.0	129
15	Translating slow-binding inhibition kinetics into cellular and in vivo effects. <i>Nature Chemical Biology</i> , 2015, 11, 416-423.	8.0	127
16	Roles of Tyrosine 158 and Lysine 165 in the Catalytic Mechanism of InhA, the Enoyl-ACP Reductase from <i>Mycobacterium tuberculosis</i> â€‘. <i>Biochemistry</i> , 1999, 38, 13623-13634.	2.5	117
17	Ultrafast Structural Dynamics in BLUF Domains:â€‘ Transient Infrared Spectroscopy of AppA and Its Mutants. <i>Journal of the American Chemical Society</i> , 2007, 129, 15556-15564.	13.7	113
18	An Alternate Proton Acceptor for Excited-State Proton Transfer in Green Fluorescent Protein:â€‘ Rewiring GFP. <i>Journal of the American Chemical Society</i> , 2008, 130, 1227-1235.	13.7	108

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19	Isotopic Labeling and Normal-Mode Analysis of a Model Green Fluorescent Protein Chromophore. <i>Journal of Physical Chemistry B</i> , 2002, 106, 6056-6066.	2.6	107
20	Slow-Onset Inhibition of the FabI Enoyl Reductase from <i>Francisella tularensis</i> : Residence Time and <i>in Vivo</i> Activity. <i>ACS Chemical Biology</i> , 2009, 4, 221-231.	3.4	106
21	Structure-Activity Studies of the Inhibition of FabI, the Enoyl Reductase from <i>Escherichia coli</i> , by Triclosan: Kinetic Analysis of Mutant FabIs. <i>Biochemistry</i> , 2003, 42, 4406-4413.	2.5	105
22	Inhibition of the Bacterial Enoyl Reductase FabI by Triclosan: A Structure-Reactivity Analysis of FabI Inhibition by Triclosan Analogues. <i>Journal of Medicinal Chemistry</i> , 2004, 47, 509-518.	6.4	101
23	Structure of Acyl Carrier Protein Bound to FabI, the FASII Enoyl Reductase from <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2006, 281, 39285-39293.	3.4	101
24	Targeting FtsZ for Antituberculosis Drug Discovery: Noncytotoxic Taxanes as Novel Antituberculosis Agents. <i>Journal of Medicinal Chemistry</i> , 2006, 49, 463-466.	6.4	100
25	Light-Driven Decarboxylation of Wild-Type Green Fluorescent Protein. <i>Journal of the American Chemical Society</i> , 2003, 125, 6919-6926.	13.7	99
26	Direct inhibitors of InhA are active against <i>Mycobacterium tuberculosis</i> . <i>Science Translational Medicine</i> , 2015, 7, 269ra3.	12.4	98
27	Ultrafast Excited and Ground-State Dynamics of the Green Fluorescent Protein Chromophore in Solution. <i>Journal of Physical Chemistry A</i> , 2004, 108, 4587-4598.	2.5	97
28	Synthesis and SAR studies of 1,4-benzoxazine MenB inhibitors: Novel antibacterial agents against <i>Mycobacterium tuberculosis</i> . <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010, 20, 6306-6309.	2.2	89
29	Crystal Structure of <i>Mycobacterium tuberculosis</i> MenB, a Key Enzyme in Vitamin K2 Biosynthesis. <i>Journal of Biological Chemistry</i> , 2003, 278, 42352-42360.	3.4	86
30	Discovery of anti-TB agents that target the cell-division protein FtsZ. <i>Future Medicinal Chemistry</i> , 2010, 2, 1305-1323.	2.3	79
31	Synthesis and Spectroscopic Studies of Model Red Fluorescent Protein Chromophores. <i>Organic Letters</i> , 2002, 4, 1523-1526.	4.6	78
32	<i>Staphylococcus aureus</i> FabI: Inhibition, Substrate Recognition, and Potential Implications for <i>In Vivo</i> Essentiality. <i>Structure</i> , 2012, 20, 802-813.	3.3	78
33	Targeting InhA, the FASII Enoyl-ACP Reductase: SAR Studies on Novel Inhibitor Scaffolds. <i>Current Topics in Medicinal Chemistry</i> , 2012, 12, 672-693.	2.1	76
34	Synthesis and <i>in vitro</i> antimycobacterial activity of B-ring modified diaryl ether InhA inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2008, 18, 3029-3033.	2.2	75
35	Ground state isomerization of a model green fluorescent protein chromophore. <i>FEBS Letters</i> , 2003, 549, 35-38.	2.8	74
36	Proton Relay Reaction in Green Fluorescent Protein (GFP): Polarization-Resolved Ultrafast Vibrational Spectroscopy of Isotopically Edited GFP. <i>Journal of Physical Chemistry B</i> , 2006, 110, 22009-22018.	2.6	73

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37	Mechanism-based inhibitors of MenE, an acyl-CoA synthetase involved in bacterial menaquinone biosynthesis. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2008, 18, 5963-5966.	2.2	73
38	Forces, bond lengths, and reactivity: fundamental insight into the mechanism of enzyme catalysis. <i>Biochemistry</i> , 1992, 31, 9122-9125.	2.5	71
39	Crystal Structures of Mycobacterium tuberculosis KasA Show Mode of Action within Cell Wall Biosynthesis and its Inhibition by Thiolactomycin. <i>Structure</i> , 2009, 17, 1004-1013.	3.3	66
40	Radiosynthesis and Bioimaging of the Tuberculosis Chemotherapeutics Isoniazid, Rifampicin and Pyrazinamide in Baboons. <i>Journal of Medicinal Chemistry</i> , 2010, 53, 2882-2891.	6.4	66
41	FtsZ: A Novel Target for Tuberculosis Drug Discovery. <i>Current Topics in Medicinal Chemistry</i> , 2007, 7, 527-543.	2.1	65
42	Proteins in Action: Femtosecond to Millisecond Structural Dynamics of a Photoactive Flavoprotein. <i>Journal of the American Chemical Society</i> , 2013, 135, 16168-16174.	13.7	65
43	A Structural and Energetic Model for the Slow-Onset Inhibition of the <i>Mycobacterium tuberculosis</i> Enoyl-ACP Reductase InhA. <i>ACS Chemical Biology</i> , 2014, 9, 986-993.	3.4	63
44	Rational Design of Broad Spectrum Antibacterial Activity Based on a Clinically Relevant Enoyl-Acyl Carrier Protein (ACP) Reductase Inhibitor. <i>Journal of Biological Chemistry</i> , 2014, 289, 15987-16005.	3.4	63
45	Positron Emission Tomography Imaging with 2- <sup>18</sup> F-Aminobenzoic Acid Detects <i>Staphylococcus aureus</i> Infections and Monitors Drug Response. <i>ACS Infectious Diseases</i> , 2018, 4, 1635-1644.	3.8	63
46	Structure and Mechanism of MbtI, the Salicylate Synthase from <i>Mycobacterium tuberculosis</i> . <i>Biochemistry</i> , 2007, 46, 954-964.	2.5	62
47	Mechanism of the Intramolecular Claisen Condensation Reaction Catalyzed by MenB, a Crotonase Superfamily Member. <i>Biochemistry</i> , 2011, 50, 9532-9544.	2.5	62
48	Role of Glutamate 144 and Glutamate 164 in the Catalytic Mechanism of Enoyl-CoA Hydratase. <i>Biochemistry</i> , 1999, 38, 9508-9516.	2.5	61
49	Ultrafast Vibrational Spectroscopy of the Flavin Chromophore. <i>Journal of Physical Chemistry B</i> , 2006, 110, 20107-20110.	2.6	61
50	Mechanism and Inhibition of saFabI, the Enoyl Reductase from <i>Staphylococcus aureus</i> . <i>Biochemistry</i> , 2008, 47, 4228-4236.	2.5	61
51	Noninvasive Determination of 2- <sup>18</sup> F-Fluoroisonicotinic Acid Hydrazide Pharmacokinetics by Positron Emission Tomography in <i>Mycobacterium tuberculosis</i> -Infected Mice. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 6284-6290.	3.2	60
52	Enoyl-Coenzyme A Hydratase-Catalyzed Exchange of the .alpha.-Protons of Coenzyme A Thiol Esters: A Model for an Enolized Intermediate in the Enzyme-Catalyzed Elimination?. <i>Biochemistry</i> , 1994, 33, 14733-14742.	2.5	59
53	Excited state dynamics in the green fluorescent protein. <i>Journal of Photochemistry and Photobiology A: Chemistry</i> , 2009, 205, 1-11.	3.9	59
54	Rational Optimization of Drug-Target Residence Time: Insights from Inhibitor Binding to the <i>Staphylococcus aureus</i> FabI Enzyme-Product Complex. <i>Biochemistry</i> , 2013, 52, 4217-4228.	2.5	58

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55	Electronic Rearrangement Induced by Substrate Analog Binding to the Enoyl-CoA Hydratase Active Site: Evidence for Substrate Activation. <i>Biochemistry</i> , 1994, 33, 12635-12643.	2.5	56
56	H-Bonding in Alcohols Is Reflected in the C-H Bond Strength: Variation of C-D Vibrational Frequency and Fractionation Factor. <i>Journal of the American Chemical Society</i> , 2000, 122, 11660-11669.	13.7	56
57	Targeting Fatty Acid Biosynthesis for the Development of Novel Chemotherapeutics against <i>Mycobacterium tuberculosis</i> : Evaluation of A-Ring-Modified Diphenyl Ethers as High-Affinity InhA Inhibitors. <i>Antimicrobial Agents and Chemotherapy</i> , 2007, 51, 3562-3567.	3.2	54
58	Fourier-transform infra-red studies of the alkaline isomerization of mitochondrial cytochrome c and the ionization of carboxylic acids. <i>Biochemical Journal</i> , 1989, 258, 599-605.	3.7	53
59	Lysine 190 Is the Catalytic Base in MenF, the Menaquinone-Specific Isochorismate Synthase from <i>Escherichia coli</i> : Implications for an Enzyme Family. <i>Biochemistry</i> , 2007, 46, 946-953.	2.5	53
60	Length of the acyl carbonyl bond in acyl-serine proteases correlates with reactivity. <i>Biochemistry</i> , 1990, 29, 10723-10727.	2.5	51
61	Unlocking the Secrets of Enzyme Power Using Raman Spectroscopy. <i>Accounts of Chemical Research</i> , 1995, 28, 8-13.	15.6	51
62	Photoexcitation of the Blue Light Using FAD Photoreceptor AppA Results in Ultrafast Changes to the Protein Matrix. <i>Journal of the American Chemical Society</i> , 2011, 133, 16893-16900.	13.7	51
63	Stable Analogues of OSB-AMP: Potent Inhibitors of MenE, the Succinylbenzoate-CoA Synthetase from Bacterial Menaquinone Biosynthesis. <i>ChemBioChem</i> , 2012, 13, 129-136.	2.6	51
64	Structural Basis for the Recognition of Mycolic Acid Precursors by KasA, a Condensing Enzyme and Drug Target from <i>Mycobacterium Tuberculosis</i> . <i>Journal of Biological Chemistry</i> , 2013, 288, 34190-34204.	3.4	48
65	Time-Dependent Diaryl Ether Inhibitors of InhA: Structure-Activity Relationship Studies of Enzyme Inhibition, Antibacterial Activity, and in vivo Efficacy. <i>ChemMedChem</i> , 2014, 9, 776-791.	3.2	48
66	Infrared spectroscopy reveals multi-step multi-timescale photoactivation in the photoconvertible protein archetype dronpa. <i>Nature Chemistry</i> , 2018, 10, 845-852.	13.6	48
67	Determination of [ <sup>11</sup> C]Rifampin Pharmacokinetics within <i>Mycobacterium tuberculosis</i> -Infected Mice by Using Dynamic Positron Emission Tomography Bioimaging. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 5768-5774.	3.2	47
68	Evaluating the Contribution of Transition-State Destabilization to Changes in the Residence Time of Triazole-Based InhA Inhibitors. <i>Journal of the American Chemical Society</i> , 2017, 139, 3417-3429.	13.7	46
69	Evidence for electrophilic catalysis in the 4-chlorobenzoyl-CoA dehalogenase reaction: UV, Raman, and <sup>13</sup> C-NMR spectral studies of dehalogenase complexes of benzoyl-CoA adducts. <i>Biochemistry</i> , 1995, 34, 13881-13888.	2.5	45
70	Raman Study of the Polarizing Forces Promoting Catalysis in 4-Chlorobenzoate-CoA Dehalogenase. <i>Biochemistry</i> , 1997, 36, 10192-10199.	2.5	45
71	Development of Modern InhA Inhibitors to Combat Drug Resistant Strains of <i>Mycobacterium tuberculosis</i> . <i>Current Topics in Medicinal Chemistry</i> , 2007, 7, 489-498.	2.1	42
72	Slow Onset Inhibition of Bacterial <sup>12</sup> -Ketoacyl-acyl Carrier Protein Synthases by Thiolactomycin. <i>Journal of Biological Chemistry</i> , 2010, 285, 6161-6169.	3.4	42

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73	Ultrafast Dynamics of Protein Proton Transfer on Short Hydrogen Bond Potential Energy Surfaces: S65T/H148D GFP.. Journal of the American Chemical Society, 2010, 132, 1452-1453.	13.7	42
74	FTIR studies of hydrogen bonding between $\hat{1}\pm, \hat{1}^2$ -unsaturated esters and alcohols. Journal of Molecular Structure, 1996, 379, 135-142.	3.6	41
75	BLUF Domain Function Does Not Require a Metastable Radical Intermediate State. Journal of the American Chemical Society, 2014, 136, 4605-4615.	13.7	41
76	CoA Adducts of 4-Oxo-4-phenylbut-2-enoates: Inhibitors of MenB from the <i>M. tuberculosis</i> Menaquinone Biosynthesis Pathway. ACS Medicinal Chemistry Letters, 2011, 2, 818-823.	2.8	40
77	Structure of Hexadienoyl-CoA Bound to Enoyl-CoA Hydratase Determined by Transferred Nuclear Overhauser Effect Measurements: A Mechanistic Predictions Based on the X-ray Structure of 4-(Chlorobenzoyl)-CoA Dehalogenase. Biochemistry, 1997, 36, 2211-2220.	2.5	39
78	Asparagine deprivation mediated by <i>Salmonella</i> asparaginase causes suppression of activation-induced T cell metabolic reprogramming. Journal of Leukocyte Biology, 2016, 99, 387-398.	3.3	39
79	Resonance Raman and Fourier transform infrared spectroscopic studies of the acyl carbonyl group in [3-(5-methyl-2-thienyl)acryloyl]chymotrypsin: evidence for artifacts in the spectra obtained by both techniques. Biochemistry, 1991, 30, 4790-4795.	2.5	38
80	Photoactivation of the BLUF Protein PixD Probed by the Site-Specific Incorporation of Fluorotyrosine Residues. Journal of the American Chemical Society, 2017, 139, 14638-14648.	13.7	38
81	Ultrafast Infrared Spectroscopy of an Isotope-Labeled Photoactivatable Flavoprotein. Biochemistry, 2011, 50, 1321-1328.	2.5	36
82	Femtosecond to Millisecond Dynamics of Light Induced Allostery in the <i>Avena sativa</i> LOV Domain. Journal of Physical Chemistry B, 2017, 121, 1010-1019.	2.6	36
83	A Virtual Screen Discovers Novel, Fragment-Sized Inhibitors of <i>Mycobacterium tuberculosis</i> InhA. Journal of Chemical Information and Modeling, 2015, 55, 645-659.	5.4	35
84	Substrate Recognition by $\hat{1}^2$ -Ketoacyl-ACP Synthases. Biochemistry, 2011, 50, 10678-10686.	2.5	34
85	Critical role of reverse transcriptase in the inhibitory mechanism of CNI-H0294 on HIV-1 nuclear translocation.. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 11859-11864.	7.1	33
86	Excited State Structure and Dynamics of the Neutral and Anionic Flavin Radical Revealed by Ultrafast Transient Mid-IR to Visible Spectroscopy. Journal of Physical Chemistry B, 2012, 116, 5810-5818.	2.6	33
87	Vibrationally Resolved Photoabsorption Spectroscopy of Red Fluorescent Protein Chromophore Anions. Physical Review Letters, 2003, 90, 118103.	7.8	32
88	Insight through Molecular Mechanics Poisson-Boltzmann Surface Area Calculations into the Binding Affinity of Triclosan and Three Analogues for FabI, the E. coli Enoyl Reductase. Journal of Medicinal Chemistry, 2006, 49, 4574-4580.	6.4	32
89	Thiolactomycin-based $\hat{1}^2$ -Ketoacyl-AcpM Synthase A (KasA) Inhibitors. Journal of Biological Chemistry, 2013, 288, 6045-6052.	3.4	32
90	Pharmacokinetic-pharmacodynamic models that incorporate drug-target binding kinetics. Current Opinion in Chemical Biology, 2019, 50, 120-127.	6.1	31

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91	Discovery of a cofactor-independent inhibitor of <i>Mycobacterium tuberculosis</i> InhA. Life Science Alliance, 2018, 1, e201800025.	2.8	31
92	Inhibiting enoyl-ACP reductase (FabI) across pathogenic microorganisms by linear sesquiterpene lactones from <i>Anthemis auriculata</i> . Phytomedicine, 2008, 15, 1125-1129.	5.3	30
93	Potential of Lichen Secondary Metabolites against <i>Plasmodium</i> Liver Stage Parasites with FAS-II as the Potential Target. Journal of Natural Products, 2013, 76, 1064-1070.	3.0	30
94	Rational Modulation of the Induced-Fit Conformational Change for Slow-Onset Inhibition in <i>Mycobacterium tuberculosis</i> InhA. Biochemistry, 2015, 54, 4683-4691.	2.5	30
95	Synthesis of Crotonyl-OxyCoA: A Mechanistic Probe of the Reaction Catalyzed by Enoyl-CoA Hydratase. Journal of the American Chemical Society, 2001, 123, 506-507.	13.7	29
96	Involvement of Glycine 141 in Substrate Activation by Enoyl-CoA Hydratase. Biochemistry, 2001, 40, 1725-1733.	2.5	29
97	Substituted diphenyl ethers as a broad-spectrum platform for the development of chemotherapeutics for the treatment of tularaemia. Journal of Antimicrobial Chemotherapy, 2009, 64, 1052-1061.	3.0	29
98	Mechanism and Inhibition of the FabV Enoyl-ACP Reductase from <i>Burkholderia mallei</i> . Biochemistry, 2010, 49, 1281-1289.	2.5	29
99	Structural and Functional Studies of Fatty Acyl Adenylate Ligases from <i>E. coli</i> and <i>L. pneumophila</i> . Journal of Molecular Biology, 2011, 406, 313-324.	4.2	29
100	Mechanism and inhibition of the FabI enoyl-ACP reductase from <i>Burkholderia pseudomallei</i> . Journal of Antimicrobial Chemotherapy, 2011, 66, 564-573.	3.0	29
101	Unraveling the Mechanism of a LOV Domain Optogenetic Sensor: A Glutamine Lever Induces Unfolding of the $\beta$ ± Helix. ACS Chemical Biology, 2020, 15, 2752-2765.	3.4	29
102	Stereoselectivity of Enoyl-CoA Hydratase Results from Preferential Activation of One of Two Bound Substrate Conformers. Chemistry and Biology, 2002, 9, 1247-1255.	6.0	28
103	Gas-phase absorption properties of DsRed model chromophores. Physical Chemistry Chemical Physics, 2003, 5, 3021-3026.	2.8	28
104	Characterizing septum inhibition in <i>Mycobacterium tuberculosis</i> for novel drug discovery. Tuberculosis, 2008, 88, 420-429.	1.9	28
105	Thiolactomycin-Based Inhibitors of Bacterial $\beta$ -Ketoacyl-ACP Synthases with in Vivo Activity. Journal of Medicinal Chemistry, 2016, 59, 5377-5390.	6.4	28
106	Direct observation of the titration of substrate carbonyl groups in the active site of $\alpha$ -chymotrypsin by resonance Raman spectroscopy. Biochemistry, 1989, 28, 6701-6709.	2.5	27
107	Active Site Heterogeneity in Dimethyl Sulfoxide Reductase from <i>Rhodobacter capsulatus</i> Revealed by Raman Spectroscopy. Biochemistry, 2001, 40, 440-448.	2.5	27
108	An Ordered Water Channel in <i>Staphylococcus aureus</i> FabI: Unraveling the Mechanism of Substrate Recognition and Reduction. Biochemistry, 2015, 54, 1943-1955.	2.5	27

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109	Mechanism of MenE Inhibition by Acyl-Adenylate Analogues and Discovery of Novel Antibacterial Agents. <i>Biochemistry</i> , 2015, 54, 6514-6524.	2.5	27
110	A quantitative mechanistic PK/PD model directly connects Btk target engagement and in vivo efficacy. <i>Chemical Science</i> , 2017, 8, 3434-3443.	7.4	27
111	Mechanism of the AppA <sub>BLUF</sub> Photocycle Probed by Site-Specific Incorporation of Fluorotyrosine Residues: Effect of the Y21 pK <sub>a</sub> on the Forward and Reverse Ground-State Reactions. <i>Journal of the American Chemical Society</i> , 2016, 138, 926-935.	13.7	26
112	Antibacterial Activity and Mode of Action of a Sulfonamide-Based Class of Oxaborole Leucyl-tRNA-Synthetase Inhibitors. <i>ACS Infectious Diseases</i> , 2019, 5, 1231-1238.	3.8	26
113	Quantifying the Interactions between Biomolecules: Guidelines for Assay Design and Data Analysis. <i>ACS Infectious Diseases</i> , 2019, 5, 796-808.	3.8	26
114	Medium-Chain Acyl-Coenzyme A Dehydrogenase Bound to a Product Analogue, Hexadienoyl-Coenzyme A: Effects on Reduction Potential, pK <sub>a</sub> , and Polarization. <i>Biochemistry</i> , 2000, 39, 13982-13992.	2.5	25
115	Ultrafast Structural Dynamics of BlsA, a Photoreceptor from the Pathogenic Bacterium <i>Acinetobacter baumannii</i> . <i>Journal of Physical Chemistry Letters</i> , 2014, 5, 220-224.	4.6	25
116	Stereospecificity of the Reaction Catalyzed by Enoyl-CoA Hydratase. <i>Journal of the American Chemical Society</i> , 2000, 122, 3987-3994.	13.7	24
117	Correlating drug target kinetics and in vivo pharmacodynamics: long residence time inhibitors of the FabI enoyl-ACP reductase. <i>Chemical Science</i> , 2016, 7, 5945-5954.	7.4	24
118	Antitubercular activity of 1,2,3-triazolyl fatty acid derivatives. <i>European Journal of Medicinal Chemistry</i> , 2017, 125, 842-852.	5.5	24
119	Protein Photochromism Observed by Ultrafast Vibrational Spectroscopy. <i>Journal of Physical Chemistry B</i> , 2013, 117, 11954-11959.	2.6	23
120	Complete Proton Transfer Cycle in GFP and Its T203V and S205V Mutants. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 9303-9307.	13.8	23
121	A Methyl 4-Oxo-4-phenylbut-2-enoate with in Vivo Activity against MRSA That Inhibits MenB in the Bacterial Menaquinone Biosynthesis Pathway. <i>ACS Infectious Diseases</i> , 2016, 2, 329-340.	3.8	22
122	Functional dynamics of a single tryptophan residue in a BLUF protein revealed by fluorescence spectroscopy. <i>Scientific Reports</i> , 2020, 10, 2061.	3.3	22
123	Molecular structures of cis- and trans-S-Ethyl thiocrotonate. A combined vibrational spectroscopic and ab initio SCF-MO study. <i>Journal of the Chemical Society, Faraday Transactions</i> , 1994, 90, 3491.	1.7	21
124	Localized electron polarization in a substrate analog binding to the active site of enoyl-CoA hydratase: Raman spectroscopic and conformational analyses of rotamers of hexadienoyl thioesters. <i>Biospectroscopy</i> , 1995, 1, 387-394.	0.6	21
125	Ultrafast transient mid IR to visible spectroscopy of fully reduced flavins. <i>Physical Chemistry Chemical Physics</i> , 2011, 13, 17642.	2.8	21
126	Vibrational Assignment of the Ultrafast Infrared Spectrum of the Photoactivatable Flavoprotein AppA. <i>Journal of Physical Chemistry B</i> , 2012, 116, 10722-10729.	2.6	21

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127	The Francisella tularensis FabI Enoyl-Acyl Carrier Protein Reductase Gene Is Essential to Bacterial Viability and Is Expressed during Infection. <i>Journal of Bacteriology</i> , 2013, 195, 351-358.	2.2	21
128	Markedly different acyl papain structures deacylate at similar rates: resonance Raman spectroscopic and kinetic evidence. <i>Journal of the American Chemical Society</i> , 1991, 113, 4297-4303.	13.7	20
129	4-Hydroxycinnamoyl-CoA: An Ionizable Probe of the Active Site of the Medium Chain Acyl-CoA Dehydrogenase. <i>Biochemistry</i> , 2000, 39, 92-101.	2.5	20
130	Variation in LOV Photoreceptor Activation Dynamics Probed by Time-Resolved Infrared Spectroscopy. <i>Biochemistry</i> , 2018, 57, 620-630.	2.5	20
131	Structure-kinetic relationships that control the residence time of drug-target complexes: insights from molecular structure and dynamics. <i>Current Opinion in Chemical Biology</i> , 2018, 44, 101-109.	6.1	20
132	Photophysics of the Blue Light Using Flavin Domain. <i>Accounts of Chemical Research</i> , 2022, 55, 402-414.	15.6	19
133	Chemistry of enzyme-substrate complexes revealed by resonance Raman spectroscopy. <i>Chemical Society Reviews</i> , 1990, 19, 293-316.	38.1	18
134	Time-Resolved Emission Spectra of Green Fluorescent Protein. <i>Photochemistry and Photobiology</i> , 2006, 82, 373.	2.5	18
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136	Structure of the <i>Yersinia pestis</i> FabV Enoyl-ACP Reductase and Its Interaction with Two 2-Pyridone Inhibitors. <i>Structure</i> , 2012, 20, 89-100.	3.3	18
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