

John S Blanchard

List of Publications by Year in descending order

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47
papers

2,185
citations

186265

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46
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docs citations

48
times ranked

2752
citing authors

#	ARTICLE	IF	CITATIONS
1	The Phage-Encoded N-Acetyltransferase Rac Mediates Inactivation of <i>Pseudomonas aeruginosa</i> Transcription by Cleavage of the RNA Polymerase Alpha Subunit. <i>Viruses</i> , 2020, 12, 976.	3.3	11
2	The active site of the <i>Mycobacterium tuberculosis</i> branched-chain amino acid biosynthesis enzyme dihydroxyacid dehydratase contains a 2Fe ²⁺ S cluster. <i>Journal of Biological Chemistry</i> , 2019, 294, 13158-13170.	3.4	12
3	Biochemical Characterization of the <i>Mycobacterium smegmatis</i> Threonine Deaminase. <i>Biochemistry</i> , 2018, 57, 6003-6012.	2.5	5
4	Identification of <i>Mycobacterial</i> RplJ/L10 and RpsA/S1 Proteins as Novel Targets for CD4 ⁺ T Cells. <i>Infection and Immunity</i> , 2017, 85, .	2.2	13
5	Mechanism-Based Inhibition of the <i>Mycobacterium tuberculosis</i> Branched-Chain Aminotransferase by <i>scpd</i> - and <i>scpl</i> -Cycloserine. <i>ACS Chemical Biology</i> , 2017, 12, 1235-1244.	3.4	33
6	Bacterial Branched-Chain Amino Acid Biosynthesis: Structures, Mechanisms, and Drugability. <i>Biochemistry</i> , 2017, 56, 5849-5865.	2.5	119
7	Mechanistic Characterization of <i>Escherichia coli</i> <i>scpl</i> -Aspartate Oxidase from Kinetic Isotope Effects. <i>Biochemistry</i> , 2017, 56, 4044-4052.	2.5	5
8	Post-translational Acetylation of MbtA Modulates <i>Mycobacterial</i> Siderophore Biosynthesis. <i>Journal of Biological Chemistry</i> , 2016, 291, 22315-22326.	3.4	34
9	Chemical Mechanism of the Branched-Chain Aminotransferase <i>IlvE</i> from <i>Mycobacterium tuberculosis</i> . <i>Biochemistry</i> , 2016, 55, 6295-6303.	2.5	23
10	Central Role of Pyruvate Kinase in Carbon Co-catabolism of <i>Mycobacterium tuberculosis</i> . <i>Journal of Biological Chemistry</i> , 2016, 291, 7060-7069.	3.4	35
11	Bacterial GCN5-Related <i>N</i> -Acetyltransferases: From Resistance to Regulation. <i>Biochemistry</i> , 2016, 55, 989-1002.	2.5	154
12	Inhibiting the β -Lactamase of <i>Mycobacterium tuberculosis</i> (Mtb) with Novel Boronic Acid Transition-State Inhibitors (BATSIs). <i>ACS Infectious Diseases</i> , 2015, 1, 234-242.	3.8	30
13	Kinetic and Structural Characterization of the Interaction of 6-Methylidene Penem 2 with the β -Lactamase from <i>Mycobacterium tuberculosis</i> . <i>Biochemistry</i> , 2015, 54, 5657-5664.	2.5	20
14	Acetylation of acetyl-CoA synthetase from <i>Mycobacterium tuberculosis</i> leads to specific inactivation of the adenylation reaction. <i>Archives of Biochemistry and Biophysics</i> , 2014, 550-551, 42-49.	3.0	19
15	Mechanism and Regulation of <i>Mycobactin</i> Fatty Acyl-AMP Ligase <i>FadD33</i> . <i>Journal of Biological Chemistry</i> , 2013, 288, 28116-28125.	3.4	35
16	A reversible acetylation system in mycobacteria. <i>FASEB Journal</i> , 2012, 26, 803.1.	0.5	0
17	Reversible Acetylation and Inactivation of <i>Mycobacterium tuberculosis</i> Acetyl-CoA Synthetase Is Dependent on cAMP. <i>Biochemistry</i> , 2011, 50, 5883-5892.	2.5	98
18	Kinetic Evidence for Interdomain Communication in the Allosteric Regulation of β -Isopropylmalate Synthase from <i>Mycobacterium tuberculosis</i> . <i>Biochemistry</i> , 2009, 48, 1996-2004.	2.5	22

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19	Steady-State and Pre-steady-State Kinetic Analysis of <i>Mycobacterium smegmatis</i> Cysteine Ligase (MshC). <i>Biochemistry</i> , 2007, 46, 11421-11429.	2.5	28
20	Kinetic analysis of the effects of monovalent cations and divalent metals on the activity of <i>Mycobacterium tuberculosis</i> Î±-isopropylmalate synthase. <i>Archives of Biochemistry and Biophysics</i> , 2006, 451, 141-148.	3.0	23
21	Kinetic and Chemical Mechanism of Î±-Isopropylmalate Synthase from <i>Mycobacterium tuberculosis</i> . <i>Biochemistry</i> , 2006, 45, 8988-8999.	2.5	50
22	Enzymology of Bacterial Lysine Biosynthesis. <i>Advances in Enzymology and Related Areas of Molecular Biology</i> , 2006, 72, 279-324.	1.3	62
23	Slow-onset Feedback Inhibition: Inhibition of <i>Mycobacterium tuberculosis</i> Î±-Isopropylmalate Synthase by Leucine. <i>Journal of the American Chemical Society</i> , 2005, 127, 10004-10005.	13.7	39
24	Kinetic and Mechanistic Characterization of Recombinant <i>Lactobacillus viridescens</i> FemX (UDP-N-acetylmuramoyl Pentapeptide-lysine N6-Alanyltransferase). <i>Journal of Biological Chemistry</i> , 2003, 278, 22861-22867.	3.4	36
25	Steady-State and Pre-Steady-State Kinetic Analysis of <i>Mycobacterium tuberculosis</i> Pantothenate Synthetase. <i>Biochemistry</i> , 2001, 40, 12904-12912.	2.5	93
26	<i>Mycobacterium tuberculosis</i> Mycothione Reductase: pH Dependence of the Kinetic Parameters and Kinetic Isotope Effects. <i>Biochemistry</i> , 2001, 40, 5119-5126.	2.5	61
27	<i>Mycobacterium tuberculosis</i> Lipoamide Dehydrogenase Is Encoded by Rv0462 and Not by the <i>hlpD</i> or <i>pdB</i> Genes. <i>Biochemistry</i> , 2001, 40, 11353-11363.	2.5	44
28	For the record: The three-dimensional structure of the ternary complex of <i>Corynebacterium glutamicum</i> diaminopimelate dehydrogenase-NADPH-L-lysine-2-aminomethylene-pimelate. <i>Protein Science</i> , 2000, 9, 2034-2037.	2.5	29
29	Enzyme-Catalyzed Acylation of Homoserine: Mechanistic Characterization of the <i>Haemophilus influenzae</i> met2-Encoded Homoserine Transacetylase. <i>Biochemistry</i> , 2000, 39, 8556-8564.	2.5	38
30	<i>Rhodococcus</i> l-Phenylalanine Dehydrogenase: Kinetics, Mechanism, and Structural Basis for Catalytic Specificity. <i>Biochemistry</i> , 2000, 39, 9174-9187.	2.5	108
31	Mechanistic Analysis of the <i>argE</i> -Encoded N-Acetylornithine Deacetylase. <i>Biochemistry</i> , 2000, 39, 1285-1293.	2.5	55
32	Vinylogous Amide Analogues of Diaminopimelic Acid (DAP) as Inhibitors of Enzymes Involved in Bacterial Lysine Biosynthesis. <i>Organic Letters</i> , 2000, 2, 3857-3860.	4.6	19
33	Chemical Mechanism of <i>Haemophilus influenzae</i> Diaminopimelate Epimerase. <i>Biochemistry</i> , 1999, 38, 4416-4422.	2.5	64
34	The Dual Biosynthetic Capability of N-Acetylornithine Aminotransferase in Arginine and Lysine Biosynthesis. <i>Biochemistry</i> , 1999, 38, 3019-3024.	2.5	73
35	Enzyme-Catalyzed Acylation of Homoserine: Mechanistic Characterization of the <i>Escherichia coli</i> metA-Encoded Homoserine Transsuccinylase. <i>Biochemistry</i> , 1999, 38, 14416-14423.	2.5	54
36	Expression, Purification, and Characterization of <i>Mycobacterium tuberculosis</i> Mycothione Reductase. <i>Biochemistry</i> , 1999, 38, 11827-11833.	2.5	94

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37	The ying and yang of rifampicin. <i>Nature Medicine</i> , 1998, 4, 14-15.	30.7	24
38	Substrate binding and conformational changes of <i>Clostridium glutamicum</i> diaminopimelate dehydrogenase revealed by hydrogen/deuterium exchange and electrospray mass spectrometry. <i>Protein Science</i> , 1998, 7, 293-299.	7.6	24
39	Structural Symmetry: The Three-Dimensional Structure of Haemophilus Influenzae Diaminopimelate Epimerase. <i>Biochemistry</i> , 1998, 37, 16452-16458.	2.5	75
40	Hydrolysis of N-Succinyl-L,diaminopimelic Acid by the Haemophilus influenzae dapE-Encoded Desuccinylase: Metal Activation, Solvent Isotope Effects, and Kinetic Mechanism. <i>Biochemistry</i> , 1998, 37, 10478-10487.	2.5	56
41	The Requirement for Manganese and Oxygen in the Isoniazid-Dependent Inactivation of Mycobacterium tuberculosis Enoyl Reductase. <i>Journal of the American Chemical Society</i> , 1997, 119, 2331-2332.	13.7	60
42	Three-Dimensional Structure of Tetrahydrodipicolinate N-Succinyltransferase. <i>Biochemistry</i> , 1997, 36, 489-494.	2.5	71
43	Interaction of Pyridine Nucleotide Substrates with Escherichia coli Dihydrodipicolinate Reductase: Thermodynamic and Structural Analysis of Binary Complexes. <i>Biochemistry</i> , 1996, 35, 13294-13302.	2.5	42
44	Expression, purification, and crystallization of meso-diaminopimelate dehydrogenase from Corynebacterium glutamicum. <i>Proteins: Structure, Function and Bioinformatics</i> , 1996, 25, 514-516.	2.6	2
45	The Biochemistry and Enzymology of Amino Acid Dehydrogenases. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 1994, 29, 415-467.	5.2	92
46	Production of trimethylamine from structurally related trimethylammonium compounds by resting cell suspensions of γ -butyrobetaine- and D,L-carnitine-grown <i>Acinetobacter calcoaceticus</i> and <i>Pseudomonas putida</i> . <i>Archives of Microbiology</i> , 1983, 135, 305-310.	2.2	12
47	Use of isotope effects to deduce the chemical mechanism of fumarase. <i>Biochemistry</i> , 1980, 19, 4506-4513.	2.5	89