

Jorge C Escalante-Semerena

List of Publications by Year in descending order

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168
papers

6,623
citations

66234

42
h-index

88477

70
g-index

173
all docs

173
docs citations

173
times ranked

4411
citing authors

#	ARTICLE	IF	CITATIONS
1	The biosynthesis of adenosylcobalamin (vitamin B12). <i>Natural Product Reports</i> , 2002, 19, 390-412.	5.2	409
2	Identification of the Protein Acetyltransferase (Pat) Enzyme that Acetylates Acetyl-CoA Synthetase in <i>Salmonella enterica</i> . <i>Journal of Molecular Biology</i> , 2004, 340, 1005-1012.	2.0	243
3	The 1.75 Å... Crystal Structure of Acetyl-CoA Synthetase Bound to Adenosine-5'-propylphosphate and Coenzyme A. <i>Biochemistry</i> , 2003, 42, 2866-2873.	1.2	208
4	N-Lysine Propionylation Controls the Activity of Propionyl-CoA Synthetase. <i>Journal of Biological Chemistry</i> , 2007, 282, 30239-30245.	1.6	176
5	Acylation of Biomolecules in Prokaryotes: a Widespread Strategy for the Control of Biological Function and Metabolic Stress. <i>Microbiology and Molecular Biology Reviews</i> , 2015, 79, 321-346.	2.9	173
6	N ^ε -Lysine Acetylation of a Bacterial Transcription Factor Inhibits Its DNA-Binding Activity. <i>PLoS ONE</i> , 2010, 5, e15123.	1.1	137
7	CobB, a New Member of the SIR2 Family of Eucaryotic Regulatory Proteins, Is Required to Compensate for the Lack of Nicotinate Mononucleotide:5,6-Dimethylbenzimidazole Phosphoribosyltransferase Activity in <i>cobT</i> Mutants during Cobalamin Biosynthesis in <i>Salmonella typhimurium</i> LT2. <i>Journal of Biological Chemistry</i> , 1998, 273, 31788-31794.	1.6	136
8	<i>Salmonella typhimurium</i> LT2 Catabolizes Propionate via the 2-Methylcitric Acid Cycle. <i>Journal of Bacteriology</i> , 1999, 181, 5615-5623.	1.0	134
9	Control of Acetyl-Coenzyme A Synthetase (AcsA) Activity by Acetylation/Deacetylation without NAD + Involvement in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2006, 188, 5460-5468.	1.0	129
10	Construction and use of new cloning vectors for the rapid isolation of recombinant proteins from <i>Escherichia coli</i> . <i>Plasmid</i> , 2008, 59, 231-237.	0.4	122
11	Minimal Functions and Physiological Conditions Required for Growth of <i>Salmonella enterica</i> on Ethanolamine in the Absence of the Metabolosome. <i>Journal of Bacteriology</i> , 2005, 187, 8039-8046.	1.0	115
12	Short-Chain Fatty Acid Activation by Acyl-Coenzyme A Synthetases Requires SIR2 Protein Function in <i>Salmonella enterica</i> and <i>Saccharomyces cerevisiae</i> . <i>Genetics</i> , 2003, 163, 545-555.	1.2	106
13	Characterization of the Propionyl-CoA Synthetase (PrpE) Enzyme of <i>Salmonella enterica</i> : Residue Lys592 Is Required for Propionyl-AMP Synthesis. <i>Biochemistry</i> , 2002, 41, 2379-2387.	1.2	104
14	Control of protein function by reversible N ^ε -lysine acetylation in bacteria. <i>Current Opinion in Microbiology</i> , 2011, 14, 200-204.	2.3	91
15	Spectroscopic and Computational Studies of the ATP:Corrinoid Adenosyltransferase (CobA) from <i>Salmonella enterica</i> : Insights into the Mechanism of Adenosylcobalamin Biosynthesis. <i>Journal of the American Chemical Society</i> , 2005, 127, 8710-8719.	6.6	90
16	Conversion of Cobinamide into Adenosylcobamide in Bacteria and Archaea. <i>Journal of Bacteriology</i> , 2007, 189, 4555-4560.	1.0	89
17	Coenzyme F420 dependence of the methylenetetrahydromethanopterin dehydrogenase of <i>Methanobacterium thermoautotrophicum</i> . <i>Biochemical and Biophysical Research Communications</i> , 1985, 133, 884-890.	1.0	85
18	An In Vitro Reducing System for the Enzymic Conversion of Cobalamin to Adenosylcobalamin. <i>Journal of Biological Chemistry</i> , 2001, 276, 32101-32108.	1.6	83

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19	Studies of Propionate Toxicity in <i>Salmonella enterica</i> Identify 2-Methylcitrate as a Potent Inhibitor of Cell Growth. <i>Journal of Biological Chemistry</i> , 2001, 276, 19094-19101.	1.6	81
20	The <i>prpE</i> gene of <i>Salmonella typhimurium</i> LT2 encodes propionyl-CoA synthetase. <i>Microbiology (United Kingdom)</i> , 1997, 141, 1007-1010.	0.7	80
21	Reversible N-lysine acetylation regulates the activity of acyl-CoA synthetases involved in anaerobic benzoate catabolism in <i>Rhodospseudomonas palustris</i> . <i>Molecular Microbiology</i> , 2010, 76, 874-888.	1.2	80
22	System-wide Studies of N-Lysine Acetylation in <i>Rhodospseudomonas palustris</i> Reveal Substrate Specificity of Protein Acetyltransferases. <i>Journal of Biological Chemistry</i> , 2012, 287, 15590-15601.	1.6	80
23	In Vitro Conversion of Propionate to Pyruvate by <i>Salmonella enterica</i> Enzymes: 2-Methylcitrate Dehydratase (PrpD) and Aconitase Enzymes Catalyze the Conversion of 2-Methylcitrate to 2-Methylisocitrate. <i>Biochemistry</i> , 2001, 40, 4703-4713.	1.2	79
24	Protein Acetylation in Bacteria. <i>Annual Review of Microbiology</i> , 2019, 73, 111-132.	2.9	78
25	Reduction of Cob(III)alamin to Cob(II)alamin in <i>Salmonella enterica</i> Serovar Typhimurium LT2. <i>Journal of Bacteriology</i> , 2000, 182, 4304-4309.	1.0	75
26	Three-Dimensional Structure of ATP:Corrinoid Adenosyltransferase from <i>Salmonella typhimurium</i> in Its Free State, Complexed with MgATP, or Complexed with Hydroxycobalamin and MgATP. <i>Biochemistry</i> , 2001, 40, 361-374.	1.2	72
27	In <i>Bacillus subtilis</i> , the Sirtuin Protein Deacetylase, Encoded by the <i>sirtN</i> Gene (Formerly <i>prpA</i>) Coenzyme A Synthetase. <i>Journal of Bacteriology</i> , 2009, 191, 1749-1755.	1.0	69
28	Propionyl Coenzyme A Is a Common Intermediate in the 1,2-Propanediol and Propionate Catabolic Pathways Needed for Expression of the <i>prpBCDE</i> Operon during Growth of <i>Salmonella enterica</i> on 1,2-Propanediol. <i>Journal of Bacteriology</i> , 2003, 185, 2802-2810.	1.0	67
29	The <i>eutT</i> Gene of <i>Salmonella enterica</i> Encodes an Oxygen-Labile, Metal-Containing ATP:Corrinoid Adenosyltransferase Enzyme. <i>Journal of Bacteriology</i> , 2004, 186, 5708-5714.	1.0	67
30	Structural Characterization of the Active Site of the PduO-Type ATP:Co(I)rrinoid Adenosyltransferase from <i>Lactobacillus reuteri</i> . <i>Journal of Biological Chemistry</i> , 2007, 282, 2596-2605.	1.6	63
31	CbiZ, an amidohydrolase enzyme required for salvaging the coenzyme B12 precursor cobinamide in archaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 3591-3596.	3.3	62
32	Residue Leu-641 of Acetyl-CoA Synthetase is Critical for the Acetylation of Residue Lys-609 by the Protein Acetyltransferase Enzyme of <i>Salmonella enterica</i> . <i>Journal of Biological Chemistry</i> , 2005, 280, 26200-26205.	1.6	62
33	Methenyl-tetrahydromethanopterin cyclohydrolase in cell extracts of <i>Methanobacterium</i> . <i>Archives of Biochemistry and Biophysics</i> , 1985, 242, 430-439.	1.4	60
34	Purification and Characterization of CobT, the Nicotinate-mononucleotide:5,6-Dimethylbenzimidazole Phosphoribosyltransferase Enzyme from <i>Salmonella typhimurium</i> LT2. <i>Journal of Biological Chemistry</i> , 1997, 272, 17662-17667.	1.6	56
35	Single-enzyme conversion of FMNH ₂ to 5,6-dimethylbenzimidazole, the lower ligand of B12. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 2921-2926.	3.3	56
36	Structural Characterization of a Human-Type Corrinoid Adenosyltransferase Confirms That Coenzyme B ₁₂ Is Synthesized through a Four-Coordinate Intermediate. <i>Biochemistry</i> , 2008, 47, 5755-5766.	1.2	55

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37	The cobinamide amidohydrolase (cobyric acid-forming) CbiZ enzyme: a critical activity of the cobamide remodelling system of <i>Rhodobacter sphaeroides</i> . <i>Molecular Microbiology</i> , 2009, 74, 1198-1210.	1.2	52
38	CobD, a Novel Enzyme with l-Threonine-O-3-phosphate Decarboxylase Activity, Is Responsible for the Synthesis of (R)-1-Amino-2-propanol O-2-Phosphate, a Proposed New Intermediate in Cobalamin Biosynthesis in <i>Salmonella typhimurium</i> LT2. <i>Journal of Biological Chemistry</i> , 1998, 273, 2684-2691.	1.6	51
39	Reassessment of the Late Steps of Coenzyme B ₁₂ Synthesis in <i>Salmonella enterica</i> : Evidence that Dephosphorylation of Adenosylcobalamin-5'-Phosphate by the CobC Phosphatase Is the Last Step of the Pathway. <i>Journal of Bacteriology</i> , 2007, 189, 2210-2218.	1.0	50
40	Purification and Characterization of the Bifunctional CobU Enzyme of <i>Salmonella typhimurium</i> LT2. <i>Journal of Biological Chemistry</i> , 1995, 270, 23560-23569.	1.6	47
41	Biochemical and Mutational Analyses of AcuA, the Acetyltransferase Enzyme That Controls the Activity of the Acetyl Coenzyme A Synthetase (AcsA) in <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2008, 190, 5132-5136.	1.0	47
42	ArsAB, a novel enzyme from <i>Sporomusa ovata</i> activates phenolic bases for adenosylcobamide biosynthesis. <i>Molecular Microbiology</i> , 2011, 81, 952-967.	1.2	47
43	Acetoacetyl-CoA synthetase activity is controlled by a protein acetyltransferase with unique domain organization in <i>Streptomyces lividans</i> . <i>Molecular Microbiology</i> , 2013, 87, 152-167.	1.2	47
44	Three-Dimensional Structure of Adenosylcobinamide Kinase/Adenosylcobinamide Phosphate Guanylyltransferase from <i>Salmonella typhimurium</i> Determined to 2.3 Å... Resolution. <i>Biochemistry</i> , 1998, 37, 7686-7695.	1.2	46
45	In <i>Salmonella enterica</i> , the sirtuin-dependent protein acylation/deacylation system (SDPADS) maintains energy homeostasis during growth on low concentrations of acetate. <i>Molecular Microbiology</i> , 2011, 80, 168-183.	1.2	44
46	Studies of Regulation of Expression of the Propionate (<i>prpBCDE</i>) Operon Provide Insights into How <i>Salmonella typhimurium</i> LT2 Integrates Its 1,2-Propanediol and Propionate Catabolic Pathways. <i>Journal of Bacteriology</i> , 1998, 180, 6511-6518.	1.0	44
47	The Tricarballoylate Utilization (<i>tcuRABC</i>) Genes of <i>Salmonella enterica</i> Serovar Typhimurium LT2. <i>Journal of Bacteriology</i> , 2004, 186, 1629-1637.	1.0	43
48	The Three-Dimensional Structures of Nicotinate Mononucleotide:5,6-Dimethylbenzimidazole Phosphoribosyltransferase (CobT) from <i>Salmonella typhimurium</i> Complexed with 5,6-Dimethylbenzimidazole and Its Reaction Products Determined to 1.9 Å... Resolution. <i>Biochemistry</i> , 1999, 38, 16125-16135.	1.2	42
49	<i>prpR</i> , <i>ntrA</i> , and <i>ihf</i> Functions Are Required for Expression of the <i>prpBCDE</i> Operon, Encoding Enzymes That Catabolize Propionate in <i>Salmonella enterica</i> Serovar Typhimurium LT2. <i>Journal of Bacteriology</i> , 2000, 182, 905-910.	1.0	42
50	The FAD-Dependent Tricarballoylate Dehydrogenase (TcuA) Enzyme of <i>Salmonella enterica</i> Converts Tricarballoylate into cis-Aconitate. <i>Journal of Bacteriology</i> , 2006, 188, 5479-5486.	1.0	40
51	A New Pathway for Salvaging the Coenzyme B ₁₂ Precursor Cobinamide in Archaea Requires Cobinamide-Phosphate Synthase (CbiB) Enzyme Activity. <i>Journal of Bacteriology</i> , 2003, 185, 7193-7201.	1.0	39
52	Acetyl Coenzyme A Synthetase Is Acetylated on Multiple Lysine Residues by a Protein Acetyltransferase with a Single Gcn5-Type N-Acetyltransferase (GNAT) Domain in <i>Saccharopolyspora erythraea</i> . <i>Journal of Bacteriology</i> , 2014, 196, 3169-3178.	1.0	39
53	Structural Investigation of the Biosynthesis of Alternative Lower Ligands for Cobamides by Nicotinate Mononucleotide: 5,6-Dimethylbenzimidazole Phosphoribosyltransferase from <i>Salmonella enterica</i> . <i>Journal of Biological Chemistry</i> , 2001, 276, 37612-37620.	1.6	38
54	The last step in coenzyme B ₁₂ synthesis is localized to the cell membrane in bacteria and archaea. <i>Microbiology (United Kingdom)</i> , 2004, 150, 1385-1395.	0.7	38

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55	Biochemical and Thermodynamic Analyses of <i>Salmonella enterica</i> Pat, a Multidomain, Multimeric μ -Lysine Acetyltransferase Involved in Carbon and Energy Metabolism. <i>MBio</i> , 2011, 2, .	1.8	38
56	Syntheses and characterization of vitamin B12-Pt(II) conjugates and their adenosylation in an enzymatic assay. <i>Journal of Biological Inorganic Chemistry</i> , 2008, 13, 335-347.	1.1	37
57	Biologically Active Isoforms of CobB Sirtuin Deacetylase in <i>Salmonella enterica</i> and <i>Erwinia amylovora</i> . <i>Journal of Bacteriology</i> , 2010, 192, 6200-6208.	1.0	37
58	New high-cloning-efficiency vectors for complementation studies and recombinant protein overproduction in <i>Escherichia coli</i> and <i>Salmonella enterica</i> . <i>Plasmid</i> , 2016, 86, 1-6.	0.4	37
59	Identification of an Alternative Nucleoside Triphosphate: 5-Deoxyadenosylcobinamide Phosphate Nucleotidyltransferase in <i>Methanobacterium thermoautotrophicum</i> . <i>Journal of Bacteriology</i> , 2000, 182, 4227-4233.	1.0	36
60	Kinetic and Spectroscopic Studies of the ATP:Corrinoid Adenosyltransferase PduO from <i>Lactobacillus reuteri</i> : Substrate Specificity and Insights into the Mechanism of Co(II)corrinoid Reduction. <i>Biochemistry</i> , 2008, 47, 9007-9015.	1.2	36
61	The <i>eutD</i> Gene of <i>Salmonella enterica</i> Encodes a Protein with Phosphotransacetylase Enzyme Activity. <i>Journal of Bacteriology</i> , 2004, 186, 1890-1892.	1.0	35
62	Purification and Initial Biochemical Characterization of ATP:Cob(I)alamin Adenosyltransferase (EutT) Enzyme of <i>Salmonella enterica</i> . <i>Journal of Biological Chemistry</i> , 2006, 281, 16971-16977.	1.6	34
63	Residue Phe112 of the Human-Type Corrinoid Adenosyltransferase (PduO) Enzyme of <i>Lactobacillus reuteri</i> Is Critical to the Formation of the Four-Coordinate Co(II) Corrinoid Substrate and to the Activity of the Enzyme. <i>Biochemistry</i> , 2009, 48, 3138-3145.	1.2	34
64	Multiple roles of ATP:cob(I)alamin adenosyltransferases in the conversion of B12 to coenzyme B12. <i>Applied Microbiology and Biotechnology</i> , 2010, 88, 41-48.	1.7	34
65	A new pathway for the synthesis of ribazole-phosphate in <i>Listeria innocua</i> . <i>Molecular Microbiology</i> , 2010, 77, 1429-1438.	1.2	34
66	In <i>Salmonella enterica</i> , 2-Methylcitrate Blocks Gluconeogenesis. <i>Journal of Bacteriology</i> , 2010, 192, 771-778.	1.0	34
67	Three-Dimensional Structure of Adenosylcobinamide Kinase/Adenosylcobinamide Phosphate Guanylyltransferase (CobU) Complexed with GMP: Evidence for a Substrate-Induced Transferase Active Site. <i>Biochemistry</i> , 1999, 38, 12995-13005.	1.2	33
68	Structural and Functional Analyses of the Human-Type Corrinoid Adenosyltransferase (PduO) from <i>Lactobacillus reuteri</i> . <i>Biochemistry</i> , 2007, 46, 13829-13836.	1.2	33
69	Structure-Guided Expansion of the Substrate Range of Methylmalonyl Coenzyme A Synthetase (MatB) of <i>Rhodospseudomonas palustris</i> . <i>Applied and Environmental Microbiology</i> , 2012, 78, 6619-6629.	1.4	33
70	The <i>EutQ</i> and <i>EutP</i> proteins are novel acetate kinases involved in ethanolamine catabolism: physiological implications for the function of the ethanolamine metabolosome in <i>Salmonella enterica</i> . <i>Molecular Microbiology</i> , 2016, 99, 497-511.	1.2	33
71	Dihydroflavin-driven Adenosylation of 4-Coordinate Co(II) Corrinoids. <i>Journal of Biological Chemistry</i> , 2010, 285, 2911-2917.	1.6	32
72	The ATP:Co(II)corrinoid Adenosyltransferase (CobA) Enzyme of <i>Salmonella enterica</i> Requires the 2'-OH Group of ATP for Function and Yields Inorganic Triphosphate as Its Reaction Byproduct. <i>Journal of Biological Chemistry</i> , 2002, 277, 33127-33131.	1.6	31

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73	2-Methylcitrate-dependent activation of the propionate catabolic operon (prpBCDE) of <i>Salmonella enterica</i> by the PrpR protein. <i>Microbiology (United Kingdom)</i> , 2004, 150, 3877-3887.	0.7	31
74	Acetate excretion during growth of <i>Salmonella enterica</i> on ethanolamine requires phosphotransacetylase (EutD) activity, and acetate recapture requires acetyl-CoA synthetase (Acs) and phosphotransacetylase (Pta) activities. <i>Microbiology (United Kingdom)</i> , 2005, 151, 3793-3801.	0.7	30
75	A Toxin Involved in <i>Salmonella</i> Persistence Regulates Its Activity by Acetylating Its Cognate Antitoxin, a Modification Reversed by CobB Sirtuin Deacetylase. <i>MBio</i> , 2017, 8, .	1.8	30
76	Biochemistry and Molecular Genetics of Cobalamin Biosynthesis1. <i>Progress in Molecular Biology and Translational Science</i> , 1997, 56, 347-384.	1.9	29
77	The acnD Genes of <i>Shewanella oneidensis</i> and <i>Vibrio cholerae</i> Encode a New Fe/S-Dependent 2-Methylcitrate Dehydratase Enzyme That Requires prpF Function In Vivo. <i>Journal of Bacteriology</i> , 2004, 186, 454-462.	1.0	29
78	The CbiB Protein of <i>Salmonella enterica</i> Is an Integral Membrane Protein Involved in the Last Step of the De Novo Corrin Ring Biosynthetic Pathway. <i>Journal of Bacteriology</i> , 2007, 189, 7697-7708.	1.0	29
79	<i>Salmonella enterica</i> Requires ApbC Function for Growth on Tricarballoylate: Evidence of Functional Redundancy between ApbC and IscU. <i>Journal of Bacteriology</i> , 2008, 190, 4596-4602.	1.0	29
80	Structural Insights into the Mechanism of Four-Coordinate Cob(II)alamin Formation in the Active Site of the <i>Salmonella enterica</i> ATP:Co(I)rrinoid Adenosyltransferase Enzyme: Critical Role of Residues Phe91 and Trp93. <i>Biochemistry</i> , 2012, 51, 9647-9657.	1.2	29
81	Three-Dimensional Structure of the L-Threonine-O-3-phosphate Decarboxylase (CobD) Enzyme from <i>Salmonella enterica</i> . <i>Biochemistry</i> , 2002, 41, 4798-4808.	1.2	28
82	ABC Transporter for Corrinoids in <i>Halobacterium</i> sp. Strain NRC-1. <i>Journal of Bacteriology</i> , 2005, 187, 5901-5909.	1.0	28
83	Formation of the Dimethylbenzimidazole Ligand of Coenzyme B12 under Physiological Conditions by a Facile Oxidative Cascade. <i>Organic Letters</i> , 2003, 5, 2211-2213.	2.4	27
84	Computer-assisted Docking of Flavodoxin with the ATP:Co(I)rrinoid Adenosyltransferase (CobA) Enzyme Reveals Residues Critical for Protein-Protein Interactions but Not for Catalysis. <i>Journal of Biological Chemistry</i> , 2005, 280, 40948-40956.	1.6	27
85	The genome of <i>Rhodobacter sphaeroides</i> strain 2.4.1 encodes functional cobinamide salvaging systems of archaeal and bacterial origins. <i>Molecular Microbiology</i> , 2008, 70, 824-836.	1.2	27
86	Small-Molecule Acetylation by GCN5-Related N ⁶ -Acetyltransferases in Bacteria. <i>Microbiology and Molecular Biology Reviews</i> , 2020, 84, .	2.9	27
87	Analysis of the Adenosylcobinamide Kinase/Adenosylcobinamide-phosphate Guanylyltransferase (CobU) Enzyme of <i>Salmonella typhimurium</i> LT2. <i>Journal of Biological Chemistry</i> , 2000, 275, 27576-27586.	1.6	26
88	In Vivo Analysis of Cobinamide Salvaging in <i>Rhodobacter sphaeroides</i> Strain 2.4.1. <i>Journal of Bacteriology</i> , 2009, 191, 3842-3851.	1.0	26
89	Structural Studies of the L-Threonine-O-3-phosphate Decarboxylase (CobD) Enzyme from <i>Salmonella enterica</i> : The Apo, Substrate, and Product Aldimine Complexes. <i>Biochemistry</i> , 2002, 41, 9079-9089.	1.2	25
90	Pentaerythritol propoxylate: a new crystallization agent and cryoprotectant induces crystal growth of 2-methylcitrate dehydratase. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2002, 58, 306-309.	2.5	25

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91	Î±-5,6-Dimethylbenzimidazole adenine dinucleotide (Î±-DAD), a putative new intermediate of coenzyme B12 biosynthesis in <i>Salmonella typhimurium</i> . <i>Microbiology (United Kingdom)</i> , 2003, 149, 983-990.	0.7	25
92	In <i>Salmonella enterica</i> , the Gcn5-Related Acetyltransferase MddA (Formerly YncA) Acetylates Methionine Sulfoximine and Methionine Sulfone, Blocking Their Toxic Effects. <i>Journal of Bacteriology</i> , 2015, 197, 314-325.	1.0	23
93	The <i>cbiS</i> Gene of the Archaeon <i>Methanopyrus kandleri</i> AV19 Encodes a Bifunctional Enzyme with Adenosylcobinamide Amidohydrolase and Î±-Ribazole-Phosphate Phosphatase Activities. <i>Journal of Bacteriology</i> , 2006, 188, 4227-4235.	1.0	22
94	The Coenzyme B12 Analog 5-Deoxyadenosylcobinamide-GDP Supports Catalysis by Methylmalonyl-CoA Mutase in the Absence of Trans-ligand Coordination. <i>Journal of Biological Chemistry</i> , 2001, 276, 1015-1019.	1.6	21
95	Capture of a Labile Substrate by Expulsion of Water Molecules from the Active Site of Nicotinate Mononucleotide:5,6-Dimethylbenzimidazole Phosphoribosyltransferase (CobT) from <i>Salmonella enterica</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 41120-41127.	1.6	21
96	The three-dimensional crystal structure of the PrpF protein of <i>Shewanella oneidensis</i> complexed with trans-aconitate: Insights into its biological function. <i>Protein Science</i> , 2007, 16, 1274-1284.	3.1	21
97	Structure of Sir2Tm bound to a propionylated peptide. <i>Protein Science</i> , 2011, 20, 131-139.	3.1	21
98	The missing link in coenzyme A biosynthesis: PanM (formerly YhhK), a yeast GCN5 acetyltransferase homologue triggers aspartate decarboxylase (PanD) maturation in <i>Salmonella enterica</i> . <i>Molecular Microbiology</i> , 2012, 84, 608-619.	1.2	21
99	The Acetylation Motif in AMP-Forming Acyl Coenzyme A Synthetases Contains Residues Critical for Acetylation and Recognition by the Protein Acetyltransferase Pat of <i>Rhodospseudomonas palustris</i> . <i>Journal of Bacteriology</i> , 2014, 196, 1496-1504.	1.0	21
100	The <i>SMUL_1544</i> Gene Product Governs Norcobamide Biosynthesis in the Tetrachloroethene-Respiring Bacterium <i>Sulfurospirillum multivorans</i> . <i>Journal of Bacteriology</i> , 2016, 198, 2236-2243.	1.0	20
101	<i>Bacillus megaterium</i> Has Both a Functional BluB Protein Required for DMB Synthesis and a Related Flavoprotein That Forms a Stable Radical Species. <i>PLoS ONE</i> , 2013, 8, e55708.	1.1	20
102	Tricarballoylate Catabolism in <i>Salmonella enterica</i> . The TcuB Protein Uses 4Fe-4S Clusters and Heme to Transfer Electrons from FADH ₂ in the Tricarballoylate Dehydrogenase (TcuA) Enzyme to Electron Acceptors in the Cell Membrane. <i>Biochemistry</i> , 2007, 46, 9107-9115.	1.2	19
103	Deciphering the Regulatory Circuitry That Controls Reversible Lysine Acetylation in <i>Salmonella enterica</i> . <i>MBio</i> , 2015, 6, e00891.	1.8	19
104	Mutation of Phosphotransacetylase but Not Isocitrate Lyase Reduces the Virulence of <i>Salmonella enterica</i> Serovar Typhimurium in Mice. <i>Infection and Immunity</i> , 2006, 74, 2498-2502.	1.0	18
105	Biosynthesis and Use of Cobalamin (B ₁₂). <i>EcoSal Plus</i> , 2008, 3, .	2.1	18
106	The EutT Enzyme of <i>Salmonella enterica</i> Is a Unique ATP:Cob(I)alamin Adenosyltransferase Metalloprotein That Requires Ferrous Ions for Maximal Activity. <i>Journal of Bacteriology</i> , 2014, 196, 903-910.	1.0	18
107	Functional Analysis of the Nicotinate Mononucleotide:5,6-Dimethylbenzimidazole Phosphoribosyltransferase (CobT) Enzyme, Involved in the Late Steps of Coenzyme B ₁₂ Biosynthesis in <i>Salmonella enterica</i> . <i>Journal of Bacteriology</i> , 2010, 192, 145-154.	1.0	17
108	N-Î±-Lysine Acetylation Control Conserved in All Three Life Domains. <i>Microbe Magazine</i> , 2010, 5, 340-344.	0.4	17

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109	Dissecting cobamide diversity through structural and functional analyses of the base-activating CobT enzyme of <i>Salmonella enterica</i> . <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2014, 1840, 464-475.	1.1	16
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