Steven E Ealick

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

35	1,291	17	35
papers	citations	h-index	g-index
35	1,475 ext. citations	9.4	4.05
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
35	Thiamin biosynthesis in prokaryotes. <i>Archives of Microbiology</i> , 1999 , 171, 293-300	3	237
34	Structural analyses reveal two distinct families of nucleoside phosphorylases. <i>Biochemical Journal</i> , 2002 , 361, 1-25	3.8	172
33	Diphthamide biosynthesis requires an organic radical generated by an iron-sulphur enzyme. <i>Nature</i> , 2010 , 465, 891-6	50.4	153
32	Structural insights into the mechanism of the PLP synthase holoenzyme from Thermotoga maritima. <i>Biochemistry</i> , 2006 , 45, 14609-20	3.2	71
31	Crystal structure of 4-methyl-5-beta-hydroxyethylthiazole kinase from Bacillus subtilis at 1.5 A resolution. <i>Biochemistry</i> , 2000 , 39, 7868-77	3.2	62
30	Radical S-adenosylmethionine (SAM) enzymes in cofactor biosynthesis: a treasure trove of complex organic radical rearrangement reactions. <i>Journal of Biological Chemistry</i> , 2015 , 290, 3980-6	5.4	53
29	Structure of a human S-adenosylmethionine decarboxylase self-processing ester intermediate and mechanism of putrescine stimulation of processing as revealed by the H243A mutant. <i>Biochemistry</i> , 2001 , 40, 9495-504	3.2	52
28	Co-opting sulphur-carrier proteins from primary metabolic pathways for 2-thiosugar biosynthesis. <i>Nature</i> , 2014 , 510, 427-31	50.4	51
27	X-ray crystal structure of glycinamide ribonucleotide synthetase from Escherichia coli. <i>Biochemistry</i> , 1998 , 37, 15647-62	3.2	49
26	Structural studies of viperin, an antiviral radical SAM enzyme. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, 6806-6811	11.5	46
25	Structural characterization of the enzyme-substrate, enzyme-intermediate, and enzyme-product complexes of thiamin phosphate synthase. <i>Biochemistry</i> , 2001 , 40, 10103-14	3.2	42
24	Crystal structure of thiamin phosphate synthase from Bacillus subtilis at 1.25 A resolution. <i>Biochemistry</i> , 1999 , 38, 6460-70	3.2	39
23	Organometallic and radical intermediates reveal mechanism of diphthamide biosynthesis. <i>Science</i> , 2018 , 359, 1247-1250	33.3	32
22	Crystal structure of thiaminase-I from Bacillus thiaminolyticus at 2.0 A resolution. <i>Biochemistry</i> , 1998 , 37, 15981-9	3.2	32
21	Anaerobic 5-Hydroxybenzimidazole Formation from Aminoimidazole Ribotide: An Unanticipated Intersection of Thiamin and Vitamin BBiosynthesis. <i>Journal of the American Chemical Society</i> , 2015 , 137, 10444-7	16.4	21
20	Non-canonical active site architecture of the radical SAM thiamin pyrimidine synthase. <i>Nature Communications</i> , 2015 , 6, 6480	17.4	21
19	From Suicide Enzyme to Catalyst: The Iron-Dependent Sulfide Transfer in Methanococcus jannaschii Thiamin Thiazole Biosynthesis. <i>Journal of the American Chemical Society</i> , 2016 , 138, 3639-42	16.4	21

18	Substrate-Dependent Cleavage Site Selection by Unconventional Radical S-Adenosylmethionine Enzymes in Diphthamide Biosynthesis. <i>Journal of the American Chemical Society</i> , 2017 , 139, 5680-5683	16.4	17
17	Lyme disease spirochaete Borrelia burgdorferi does not require thiamin. <i>Nature Microbiology</i> , 2016 , 2, 16213	26.6	16
16	Structural Basis for Iron-Mediated Sulfur Transfer in Archael and Yeast Thiazole Synthases. <i>Biochemistry</i> , 2016 , 55, 1826-38	3.2	14
15	Lysine relay mechanism coordinates intermediate transfer in vitamin B6 biosynthesis. <i>Nature Chemical Biology</i> , 2017 , 13, 290-294	11.7	13
14	Structural Basis of the Substrate Selectivity of Viperin. <i>Biochemistry</i> , 2020 , 59, 652-662	3.2	12
13	Different polyamine pathways from bacteria have replaced eukaryotic spermidine biosynthesis in ciliates Tetrahymena thermophila and Paramecium tetaurelia. <i>Molecular Microbiology</i> , 2015 , 97, 791-807	7 ^{4.1}	11
12	Crystal Structures of the Iron-Sulfur Cluster-Dependent Quinolinate Synthase in Complex with Dihydroxyacetone Phosphate, Iminoaspartate Analogues, and Quinolinate. <i>Biochemistry</i> , 2016 , 55, 4135	. 3 ,2	10
11	Ethenoguanines undergo glycosylation by nucleoside 2Vdeoxyribosyltransferases at non-natural sites. <i>PLoS ONE</i> , 2014 , 9, e115082	3.7	9
10	Burkholderia glumae ToxA Is a Dual-Specificity Methyltransferase That Catalyzes the Last Two Steps of Toxoflavin Biosynthesis. <i>Biochemistry</i> , 2016 , 55, 2748-59	3.2	8
9	Polyketide Ring Expansion Mediated by a Thioesterase, Chain Elongation and Cyclization Domain, in Azinomycin Biosynthesis: Characterization of AziB and AziG. <i>Biochemistry</i> , 2016 , 55, 704-14	3.2	6
8	Towards the structural characterization of the human methyltransferome. <i>Current Opinion in Structural Biology</i> , 2018 , 53, 12-21	8.1	4
7	Menaquinone Biosynthesis: Biochemical and Structural Studies of Chorismate Dehydratase. <i>Biochemistry</i> , 2019 , 58, 1837-1840	3.2	3
6	The Crystal Structure of Dph2 in Complex with Elongation Factor 2 Reveals the Structural Basis for the First Step of Diphthamide Biosynthesis. <i>Biochemistry</i> , 2019 , 58, 4343-4351	3.2	3
5	Biochemical Characterization and Structural Basis of Reactivity and Regioselectivity Differences between Burkholderia thailandensis and Burkholderia glumae 1,6-Didesmethyltoxoflavin N-Methyltransferase. <i>Biochemistry</i> , 2017 , 56, 3934-3944	3.2	3
4	Biochemical and structural characterization of Klebsiella pneumoniae oxamate amidohydrolase in the uric acid degradation pathway. <i>Acta Crystallographica Section D: Structural Biology</i> , 2016 , 72, 808-16	5.5	3
3	Structural basis of elongation factor 2 switching. Current Research in Structural Biology, 2020 , 2, 25-34	2.8	3
2	The use of Trichomonas vaginalis purine nucleoside phosphorylase to activate fludarabine in the treatment of solid tumors. <i>Cancer Chemotherapy and Pharmacology</i> , 2020 , 85, 573-583	3.5	2
1	Trapping and structural characterisation of a covalent intermediate in vitamin B biosynthesis catalysed by the Pdx1 PLP synthase RSC Chemical Biology, 2022, 3, 227-230	3	