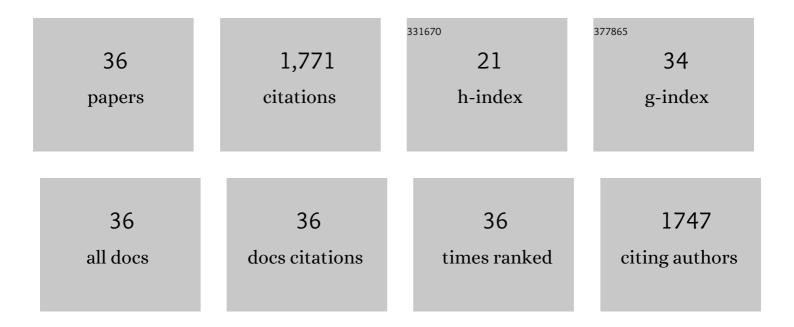
## T Joseph Kappock

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Pterin-Dependent Amino Acid Hydroxylases. Chemical Reviews, 1996, 96, 2659-2756.	47.7	310
2	Diverse carotenoids protect against chemically induced neoplastic transformation. Carcinogenesis, 1991, 12, 671-678.	2.8	252
3	Single-molecule paleoenzymology probes the chemistry of resurrected enzymes. Nature Structural and Molecular Biology, 2011, 18, 592-596.	8.2	182
4	Lipases Provide a New Mechanistic Model for Polyhydroxybutyrate (PHB) Synthases:  Characterization of the Functional Residues in Chromatium vinosum PHB Synthase. Biochemistry, 2000, 39, 3927-3936.	2.5	106
5	A Specialized Citric Acid Cycle Requiring Succinyl-Coenzyme A (CoA):Acetate CoA-Transferase (AarC) Confers Acetic Acid Resistance on the Acidophile <i>Acetobacter aceti</i> . Journal of Bacteriology, 2008, 190, 4933-4940.	2.2	99
6	Modular evolution of the purine biosynthetic pathway. Current Opinion in Chemical Biology, 2000, 4, 567-572.	6.1	93
7	X-ray crystal structure of aminoimidazole ribonucleotide synthetase (PurM), from the Escherichia coli purine biosynthetic pathway at 2.5 Ã resolution. Structure, 1999, 7, 1155-1166.	3.3	68
8	Spectroscopic Characterization of the Catalytically Competent Ferrous Site of the Resting, Activated, and Substrate-Bound Forms of Phenylalanine Hydroxylase. Journal of the American Chemical Society, 1997, 119, 1901-1915.	13.7	65
9	X-ray Crystal Structure of Glycinamide Ribonucleotide Synthetase fromEscherichia coliâ€,‡. Biochemistry, 1998, 37, 15647-15662.	2.5	57
10	Three-Dimensional Structure ofN5-Carboxyaminoimidazole Ribonucleotide Synthetase:Â A Member of the ATP Grasp Protein Superfamilyâ€,‡. Biochemistry, 1999, 38, 15480-15492.	2.5	52
11	Crystal structure of Escherichia coli PurE, an unusual mutase in the purine biosynthetic pathway. Structure, 1999, 7, 1395-1406.	3.3	50
12	[6] Solubilization, cellular uptake, and activity of β-carotene and other carotenoids as inhibitors of neoplastic transformation in cultured cells. Methods in Enzymology, 1993, 214, 55-68.	1.0	46
13	Spectroscopic and Kinetic Properties of Unphosphorylated Rat Hepatic Phenylalanine Hydroxylase Expressed in Escherichia coli. Journal of Biological Chemistry, 1995, 270, 30532-30544.	3.4	43
14	Evidence for the Direct Transfer of the Carboxylate of N5-Carboxyaminoimidazole Ribonucleotide (N5-CAIR) To Generate 4-Carboxy-5-aminoimidazole Ribonucleotide Catalyzed by Escherichia coli PurE, an N5-CAIR Mutase. Biochemistry, 1999, 38, 3012-3018.	2.5	43
15	Structure of a NADH-Insensitive Hexameric Citrate Synthase that Resists Acid Inactivation. Biochemistry, 2006, 45, 13487-13499.	2.5	43
16	Biochemical and Structural Studies of N5-Carboxyaminoimidazole Ribonucleotide Mutase from the Acidophilic Bacterium Acetobacter aceti. Biochemistry, 2006, 45, 8193-8208.	2.5	27
17	Alanine racemase from the acidophile Acetobacter aceti. Protein Expression and Purification, 2007, 51, 39-48.	1.3	25
18	Acidophilic adaptations in the structure ofAcetobacter aceti N5-carboxyaminoimidazole ribonucleotide mutase (PurE). Acta Crystallographica Section D: Biological Crystallography, 2004, 60, 1753-1760.	2.5	24

Т ЈОЅЕРН КАРРОСК

#	Article	IF	CITATIONS
19	You are lost without a map: Navigating the sea of protein structures. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2015, 1854, 258-268.	2.3	24
20	N5-CAIR Mutase: Role of a CO2Binding Site and Substrate Movement in Catalysisâ€,‡. Biochemistry, 2007, 46, 2842-2855.	2.5	23
21	Crystal Structures of <i>Acetobacter aceti</i> Succinyl-Coenzyme A (CoA):Acetate CoA-Transferase Reveal Specificity Determinants and Illustrate the Mechanism Used by Class I CoA-Transferases. Biochemistry, 2012, 51, 8422-8434.	2.5	22
22	<i>Treponema denticola</i> PurE Is a Bacterial AIR Carboxylase. Biochemistry, 2011, 50, 4623-4637.	2.5	19
23	Formyl 0enzyme A (CoA):0xalate CoAâ€ŧransferase from the acidophile <i>Acetobacter aceti</i> has a distinctive electrostatic surface and inherent acid stability. Protein Science, 2012, 21, 686-696.	7.6	18
24	Function and X-Ray crystal structure of Escherichia coli YfdE. PLoS ONE, 2013, 8, e67901.	2.5	13
25	Cloning and transcriptional analysis of Crepis alpina fatty acid desaturases affecting the biosynthesis of crepenynic acid. Journal of Experimental Botany, 2007, 58, 1421-1432.	4.8	12
26	Atomic-resolution crystal structure of thioredoxin from the acidophilic bacterium Acetobacter aceti. Protein Science, 2006, 16, 92-98.	7.6	11
27	Functional analysis of the acetic acid resistance (aar) gene cluster in Acetobacter aceti strain 1023. Acetic Acid Bacteria, 2013, 2, 3.	1.0	11
28	The Partial Substrate Dethiaacetyl-Coenzyme A Mimics All Critical Carbon Acid Reactions in the Condensation Half-Reaction Catalyzed by <i>Thermoplasma acidophilum</i> Citrate Synthase. Biochemistry, 2009, 48, 7878-7891.	2.5	9
29	An active site–tail interaction in the structure of hexahistidine-tagged <i>Thermoplasma acidophilum</i> citrate synthase. Acta Crystallographica Section F, Structural Biology Communications, 2015, 71, 1292-1299.	0.8	7
30	Multiple Active Site Histidine Protonation States in Acetobacter aceti N5-Carboxyaminoimidazole Ribonucleotide Mutase Detected by REDOR NMR. Biochemistry, 2007, 46, 9507-9512.	2.5	4
31	Draft Genome Sequence of Acetobacter aceti Strain 1023, a Vinegar Factory Isolate. Genome Announcements, 2014, 2, .	0.8	4
32	Metal stopping reagents facilitate discontinuous activity assays of the de novo purine biosynthesis enzyme PurE. Analytical Biochemistry, 2014, 452, 43-45.	2.4	3
33	Functional Dissection of the Bipartite Active Site of the Class I Coenzyme A (CoA)-Transferase Succinyl-CoA:Acetate CoA-Transferase. Frontiers in Chemistry, 2016, 4, 23.	3.6	3
34	The Purine Machine Scores a Base Hit. ACS Chemical Biology, 2008, 3, 460-462.	3.4	2
35	Altered Pathway Routing in a Class of Salmonella enterica Serovar Typhimurium Mutants Defective in Aminoimidazole Ribonucleotide Synthetase. Journal of Bacteriology, 2001, 183, 2234-2240.	2.2	1
36	A Biosynthetic Enzyme Worms Its Way out of a Conserved Mechanism. Structure, 2013, 21, 1719-1720.	3.3	0