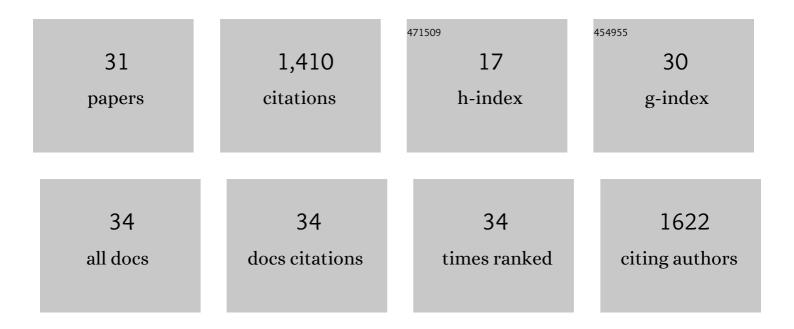
Markos Koutmos

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Antivitamins B12: Synthesis and application as inhibitory ligand of the B12-tailoring enzyme CblC. Methods in Enzymology, 2022, 668, 157-178.	1.0	1
2	Pseudouridine synthase 7 is an opportunistic enzyme that binds and modifies substrates with diverse sequences and structures. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	21
3	Disease-associated mutations in mitochondrial precursor tRNAs affect binding, m1R9 methylation, and tRNA processing by mtRNase P. Rna, 2021, 27, 420-432.	3.5	9
4	Patient mutations in human ATP:cob(I)alamin adenosyltransferase differentially affect its catalytic versus chaperone functions. Journal of Biological Chemistry, 2021, 297, 101373.	3.4	3
5	Mobile loop dynamics in adenosyltransferase control binding and reactivity of coenzyme B ₁₂ . Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 30412-30422.	7.1	18
6	An Interprotein Co–S Coordination Complex in the B ₁₂ -Trafficking Pathway. Journal of the American Chemical Society, 2020, 142, 16334-16345.	13.7	20
7	The Photoactive Excited State of the B ₁₂ -Based Photoreceptor CarH. Journal of Physical Chemistry B, 2020, 124, 10732-10738.	2.6	25
8	Interplay between substrate recognition, 5′ end tRNA processing and methylation activity of human mitochondrial RNase P. Rna, 2019, 25, 1646-1660.	3.5	21
9	Itaconyl-CoA forms a stable biradical in methylmalonyl-CoA mutase and derails its activity and repair. Science, 2019, 366, 589-593.	12.6	71
10	The folate-binding module of <i>Thermus thermophilus</i> cobalamin-dependent methionine synthase displays a distinct variation of the classical TIM barrel: a TIM barrel with a `twist'. Acta Crystallographica Section D: Structural Biology, 2018, 74, 41-51.	2.3	5
11	Coordination chemistry controls the thiol oxidase activity of the B12-trafficking protein CblC. Journal of Biological Chemistry, 2017, 292, 9733-9744.	3.4	19
12	Antivitamin B ₁₂ Inhibition of the Human B ₁₂ â€Processing Enzyme CblC: Crystal Structure of an Inactive Ternary Complex with Glutathione as the Cosubstrate. Angewandte Chemie - International Edition, 2017, 56, 7387-7392.	13.8	30
13	Inhibierung des humanen B ₁₂ â€verarbeitenden Enzyms CblC durch Antivitamineâ€B ₁₂ – Kristallstruktur des inaktiven tern¤en Komplexes mit dem Kosubstrat Glutathion. Angewandte Chemie, 2017, 129, 7493-7498.	2.0	6
14	Molecular recognition of pre-tRNA by <i>Arabidopsis</i> protein-only Ribonuclease P. Rna, 2017, 23, 1860-1873.	3.5	16
15	Loss of the mitochondrial protein-only ribonuclease P complex causes aberrant tRNA processing and lethality in Drosophila. Nucleic Acids Research, 2016, 44, 6409-6422.	14.5	32
16	Nuclear Protein-Only Ribonuclease P2 Structure and Biochemical Characterization Provide Insight into the Conserved Properties of tRNA 5′ End Processing Enzymes. Journal of Molecular Biology, 2016, 428, 26-40.	4.2	31
17	Differential substrate recognition by isozymes of plant protein-only Ribonuclease P. Rna, 2016, 22, 782-792.	3.5	26
18	Evolutionary Analyses and Natural Selection of Betaine-Homocysteine S-Methyltransferase (BHMT) and BHMT2 Genes. PLoS ONE, 2015, 10, e0134084.	2.5	13

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19	Structure of Human B12 Trafficking Protein CblD Reveals Molecular Mimicry and Identifies a New Subfamily of Nitro-FMN Reductases. Journal of Biological Chemistry, 2015, 290, 29155-29166.	3.4	25
20	Specific potassium ion interactions facilitate homocysteine binding to betaine-homocysteine <i>S</i> -methyltransferase. Proteins: Structure, Function and Bioinformatics, 2014, 82, 2552-2564.	2.6	10
21	RNase P enzymes. RNA Biology, 2013, 10, 909-914.	3.1	36
22	Mitochondrial ribonuclease P structure provides insight into the evolution of catalytic strategies for precursor-tRNA 5â€2 processing. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 16149-16154.	7.1	110
23	Structural Basis of Multifunctionality in a Vitamin B12-processing Enzyme. Journal of Biological Chemistry, 2011, 286, 29780-29787.	3.4	78
24	Structural basis for substrate activation and regulation by cystathionine beta-synthase (CBS) domains in cystathionine β-synthase. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 20958-20963.	7.1	341
25	Insights into the reactivation of cobalamin-dependent methionine synthase. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 18527-18532.	7.1	86
26	Structureâ^'Activity Study of New Inhibitors of Human Betaine-Homocysteine S-Methyltransferase. Journal of Medicinal Chemistry, 2009, 52, 3652-3665.	6.4	10
27	Cobalamin-dependent and cobamide-dependent methyltransferases. Current Opinion in Structural Biology, 2008, 18, 658-666.	5.7	149
28	A disulfide-stabilized conformer of methionine synthase reveals an unexpected role for the histidine ligand of the cobalamin cofactor. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 4115-4120.	7.1	47
29	Betaine-Homocysteine S-Methyltransferase-2 Is an S-Methylmethionine-Homocysteine Methyltransferase. Journal of Biological Chemistry, 2008, 283, 8939-8945.	3.4	71
30	Metal active site elasticity linked to activation of homocysteine in methionine synthases. Proceedings of the United States of America, 2008, 105, 3286-3291.	7.1	56
31	Borohydride, Azide, and Chloride Anions As Terminal Ligands on Fe/Mo/S Clusters. Synthesis, Structure and Characterization of [(Cl4-cat)(PPr3) MoFe3S4(X)2]2(Bu4N)4and [(Cl4-cat)(PPr3)MoFe3S4(PPr3)(X)]2(Bu4N)2(X = N3-, BH4-, Cl-) Double-Fused Cubanes. NMR Reactivity Studies of [(Cl4-cat)(PPr3) MoFe3S4(BH4)2]2(Bu4N)4. Inorganic Chemistry. 2006. 45. 3648-3656.	4.0	14