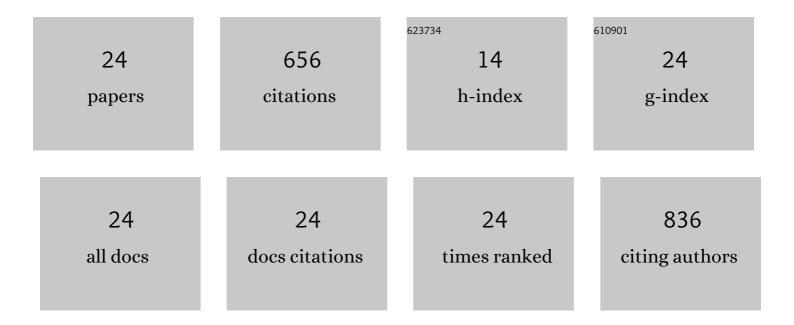
Lars-Göran MÃ¥rtensson

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6053035/publications.pdf

Version: 2024-02-01



#	Article	IF	CITATIONS
1	Comparison of Electron Paramagnetic Resonance Methods to Determine Distances between Spin Labels on Human Carbonic Anhydrase II. Biophysical Journal, 2001, 80, 2886-2897.	0.5	74
2	Adsorption of Human Carbonic Anhydrase II Variants to Silica Nanoparticles Occur Stepwise:Â Binding Is Followed by Successive Conformational Changes to a Molten-Globule-like State. Langmuir, 2000, 16, 8470-8479.	3.5	61
3	Cis-transisomerization is rate-determining in the reactivation of denatured human carbonic anhydrase II as evidenced by proline isomerase. FEBS Letters, 1992, 296, 90-94.	2.8	57
4	Structural Mapping of an Aggregation Nucleation Site in a Molten Globule Intermediate. Journal of Biological Chemistry, 1999, 274, 32897-32903.	3.4	52
5	Robust and convenient analysis of protein thermal and chemical stability. Protein Science, 2015, 24, 2055-2062.	7.6	51
6	Dramatic Stabilization of the Native State of Human Carbonic Anhydrase II by an Engineered Disulfide Bondâ€. Biochemistry, 2002, 41, 15867-15875.	2.5	41
7	Folding around the C-terminus of human carbonic anhydrase II Kinetic characterization by use of a chemically reactive SH-group introduced by protein engineering. FEBS Letters, 1991, 289, 117-122.	2.8	36
8	High-Resolution Probing of Local Conformational Changes in Proteins by the Use of Multiple Labeling: Unfolding and Self-Assembly of Human Carbonic Anhydrase II Monitored by Spin, Fluorescent, and Chemical Reactivity Probes. Biophysical Journal, 2001, 80, 2867-2885.	0.5	35
9	Critical biophysical properties in the <i>Pseudomonas aeruginosa</i> efflux gene regulator MexR are targeted by mutations conferring multidrug resistance. Protein Science, 2010, 19, 680-692.	7.6	32
10	Role of an evolutionarily invariant serine for the stability of human carbonic anhydrase II. BBA - Proteins and Proteomics, 1992, 1118, 179-186.	2.1	29
11	Redesign of human carbonic anhydrase II for increased esterase activity and specificity towards esters with long acyl chains. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2006, 1764, 1601-1606.	2.3	25
12	Characterization of a novel sequence variant, TPMT*28, in the human thiopurine methyltransferase gene. Pharmacogenetics and Genomics, 2010, 20, 700-707.	1.5	25
13	Methotrexate binds to recombinant thiopurine S-methyltransferase and inhibits enzyme activity after high-dose infusions in childhood leukaemia. European Journal of Clinical Pharmacology, 2013, 69, 1641-1649.	1.9	21
14	One amino acid makes a difference–Characterization of a new TPMT allele and the influence of SAM on TPMT stability. Scientific Reports, 2017, 7, 46428.	3.3	16
15	Subtle Differences in Dissociation Rates of Interactions between Destabilized Human Carbonic Anhydrase II Mutants and Immobilized Benzenesulfonamide Inhibitors Probed by a Surface Plasmon Resonance Biosensor. Analytical Biochemistry, 2001, 296, 188-196.	2.4	15
16	A test of proposed rules for helix capping: Implications for protein design. Protein Science, 2009, 11, 516-521.	7.6	13
17	Methylation of selenocysteine catalysed by thiopurine S-methyltransferase. Biochimica Et Biophysica Acta - General Subjects, 2019, 1863, 182-190.	2.4	13
18	Denaturant-Assisted Formation of a Stabilizing Disulfide Bridge from Engineered Cysteines in Nonideal Conformationsâ€. Biochemistry, 2005, 44, 3487-3493.	2.5	12

#	Article	IF	CITATIONS
19	Comprehensive study of thiopurine methyltransferase genotype, phenotype, and genotype-phenotype discrepancies in Sweden. Biochemical Pharmacology, 2019, 164, 263-272.	4.4	11
20	Circumnavigating Misfolding Traps in the Energy Landscape through Protein Engineering:Â Suppression of Molten Globule and Aggregation in Carbonic Anhydraseâ€. Biochemistry, 2004, 43, 6803-6807.	2.5	10
21	Pharmacogenetic studies of thiopurine methyltransferase genotypeâ€phenotype concordance and effect of methotrexate on thiopurine metabolism. Basic and Clinical Pharmacology and Toxicology, 2021, 128, 52-65.	2.5	8
22	Thermodynamic Interrogation of a Folding Disease. Mutant Mapping of Position 107 in Human Carbonic Anhydrase II Linked to Marble Brain Disease. Biochemistry, 2008, 47, 1288-1298.	2.5	7
23	Structural Characteristics Determine the Cause of the Low Enzyme Activity of Two Thiopurine <i>S</i> -Methyltransferase Allelic Variants: A Biophysical Characterization of TPMT*2 and TPMT*5. Biochemistry, 2012, 51, 5912-5920.	2.5	7
24	In Vitro Protein Stability of Two Naturally Occurring Thiopurine <i>S</i> -Methyltransferase Variants: Biophysical Characterization of TPMT*6 and TPMT*8. ACS Omega, 2017, 2, 4991-4999.	3.5	5