Andrey Galkin

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Cold-Active Serine Alkaline Protease from the Psychrotrophic Bacterium <i>Shewanella</i> Strain Ac10: Gene Cloning and Enzyme Purification and Characterization. Applied and Environmental Microbiology, 1999, 65, 611-617.	3.1	101
2	Cold-active esterase from Psychrobacter sp. Ant300: gene cloning, characterization, and the effects of Glyâ†'Pro substitution near the active site on its catalytic activity and stability. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2004, 1696, 59-65.	2.3	92
3	Structural Insight into Arginine Degradation by Arginine Deiminase, an Antibacterial and Parasite Drug Target. Journal of Biological Chemistry, 2004, 279, 14001-14008.	3.4	68
4	Rational design of a trispecific antibody targeting the HIV-1 Env with elevated anti-viral activity. Nature Communications, 2018, 9, 877.	12.8	65
5	Crystal Structures Representing the Michaelis Complex and the Thiouronium Reaction Intermediate of Pseudomonas aeruginosa Arginine Deiminase. Journal of Biological Chemistry, 2005, 280, 34080-34087.	3.4	58
6	Kinetic Analysis of Pseudomonas aeruginosa Arginine Deiminase Mutants and Alternate Substrates Provides Insight into Structural Determinants of Function. Biochemistry, 2006, 45, 1162-1172.	2.5	58
7	Arginine Deiminase Uses an Active-Site Cysteine in Nucleophilic Catalysis ofl-Arginine Hydrolysis. Journal of the American Chemical Society, 2004, 126, 5374-5375.	13.7	56
8	Structural Basis for Inactivation of Giardia lamblia Carbamate Kinase by Disulfiram. Journal of Biological Chemistry, 2014, 289, 10502-10509.	3.4	51
9	Conversion of \hat{l}_{\pm} -keto acids to d-amino acids by coupling of four enzyme reactions. Journal of Bioscience and Bioengineering, 1997, 83, 299-300.	0.9	47
10	Characterization, Kinetics, and Crystal Structures of Fructose-1,6-bisphosphate Aldolase from the Human Parasite, Giardia lamblia. Journal of Biological Chemistry, 2007, 282, 4859-4867.	3.4	45
11	High-Throughput <i>Giardia lamblia</i> Viability Assay Using Bioluminescent ATP Content Measurements. Antimicrobial Agents and Chemotherapy, 2011, 55, 667-675.	3.2	43
12	Cold-Adapted Alanine Dehydrogenases from Two Antarctic Bacterial Strains: Gene Cloning, Protein Characterization, and Comparison with Mesophilic and Thermophilic Counterparts. Applied and Environmental Microbiology, 1999, 65, 4014-4020.	3.1	38
13	Inactivation of Microbial Arginine Deiminases by <scp>l</scp> -Canavanine. Journal of the American Chemical Society, 2008, 130, 1918-1931.	13.7	37
14	Discovery of Novel Antigiardiasis Drug Candidates. Antimicrobial Agents and Chemotherapy, 2014, 58, 7303-7311.	3.2	33
15	Cold-active DnaK of an Antarctic psychrotroph Shewanella sp. Ac10 supporting the growth of dnaK-null mutant of Escherichia coli at cold temperatures. Extremophiles, 2005, 9, 145-150.	2.3	30
16	Mechanisms of catalysis and inhibition operative in the arginine deiminase from the human pathogen Giardia lamblia. Bioorganic Chemistry, 2009, 37, 149-161.	4.1	30
17	Structural Insights into the Substrate Binding and Stereoselectivity of <i>Giardia</i> Fructose-1,6-bisphosphate Aldolase [,] . Biochemistry, 2009, 48, 3186-3196.	2.5	30
18	Rational design, synthesis and evaluation of first generation inhibitors of the Giardia lamblia fructose-1,6-biphosphate aldolase. Journal of Inorganic Biochemistry, 2011, 105, 509-517.	3.5	23

#	Article	IF	CITATIONS
19	The HI0073/HI0074 protein pair from Haemophilus influenzae is a member of a new nucleotidyltransferase family: Structure, sequence analyses, and solution studies. Proteins: Structure, Function and Bioinformatics, 2002, 50, 249-260.	2.6	22
20	A Homogenous Luminescence Assay Reveals Novel Inhibitors for Giardia LambliaCarbamate Kinase. Current Chemical Genomics, 2012, 6, 93-102.	2.0	16
21	Purification and characterization of alanine dehydrogenase from a marine bacterium, Vibrio proteolyticus. Journal of Molecular Catalysis B: Enzymatic, 2003, 23, 373-378.	1.8	13
22	Xâ€ray structure and kinetic properties of ornithine transcarbamoylase from the human parasite ⟨i⟩Giardia lamblia⟨ i⟩. Proteins: Structure, Function and Bioinformatics, 2009, 76, 1049-1053.	2.6	13
23	X-ray structure and characterization of carbamate kinase from the human parasite <i>Giardia lamblia</i> . Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 386-390.	0.7	11
24	Conversion of cofactor specificities of alanine dehydrogenases by site-directed mutagenesis. Journal of Molecular Catalysis B: Enzymatic, 2004, 30, 173-176.	1.8	10
25	Structure of HI0073 from Haemophilus influenzae, the nucleotide-binding domain of a two-protein nucleotidyl transferase. Proteins: Structure, Function and Bioinformatics, 2005, 60, 807-811.	2.6	10
26	Improvement of thermostability of cold-active serine alkaline protease from the psychrotrophic bacterium Shewanella sp. strain Ac10 by rational mutagenesis. Journal of Molecular Catalysis B: Enzymatic, 2003, 22, 113-117.	1.8	8
27	Discovery and Preclinical Development of Antigiardiasis Fumagillol Derivatives. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	6
28	X-ray structure of HI0817 fromHaemophilus influenzae: Protein of unknown function with a novel fold. Proteins: Structure, Function and Bioinformatics, 2004, 57, 874-877.	2.6	4
29	Crystal Structures of Carbamate Kinase from Giardia lamblia Bound with Citric Acid and AMP-PNP. PLoS ONE, 2013, 8, e64004.	2.5	4
30	HIV-1 gp120–CD4-Induced Antibody Complex Elicits CD4 Binding Site–Specific Antibody Response in Mice. Journal of Immunology, 2020, 204, 1543-1561.	0.8	4