

Ekaterina Y Bezsudnova

List of Publications by Year in descending order

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Version: 2024-02-01

36
papers

513
citations

687363

13
h-index

713466

21
g-index

38
all docs

38
docs citations

38
times ranked

595
citing authors

#	ARTICLE	IF	CITATIONS
1	Denitrification in a binary culture and thiocyanate metabolism in <i>Thiohalophilus thiocyanoxidans</i> gen. nov. sp. nov. – a moderately halophilic chemolithoautotrophic sulfur-oxidizing Gammaproteobacterium from hypersaline lakes. <i>Archives of Microbiology</i> , 2007, 187, 441-450.	2.2	76
2	Thiocyanate hydrolase, the primary enzyme initiating thiocyanate degradation in the novel obligately chemolithoautotrophic halophilic sulfur-oxidizing bacterium <i>Thiohalophilus thiocyanoxidans</i> . <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2007, 1774, 1563-1570.	2.3	42
3	Properties of bacterial and archaeal branched-chain amino acid aminotransferases. <i>Biochemistry (Moscow)</i> , 2017, 82, 1572-1591.	1.5	37
4	Water-soluble cyclopalladated aryl oxime: a potent “green” catalyst. <i>Journal of Organometallic Chemistry</i> , 2001, 622, 38-42.	1.8	33
5	Structural insight into the substrate specificity of PLP fold type IV transaminases. <i>Applied Microbiology and Biotechnology</i> , 2020, 104, 2343-2357.	3.6	32
6	First structure of archaeal branched-chain amino acid aminotransferase from <i>Thermoproteus uzoniensis</i> specific for L-amino acids and R-amines. <i>Extremophiles</i> , 2016, 20, 215-225.	2.3	28
7	Thermostable Branched-Chain Amino Acid Transaminases From the Archaea <i>Geoglobus acetivorans</i> and <i>Archaeoglobus fulgidus</i> : Biochemical and Structural Characterization. <i>Frontiers in Bioengineering and Biotechnology</i> , 2019, 7, 7.	4.1	26
8	Structural insight into the molecular basis of polyextremophilicity of short-chain alcohol dehydrogenase from the hyperthermophilic archaeon <i>Thermococcus sibiricus</i> . <i>Biochimie</i> , 2012, 94, 2628-2638.	2.6	23
9	A Novel highly thermostable branched-chain amino acid aminotransferase from the crenarchaeon <i>Vulcanisaeta moutnovskia</i> . <i>Enzyme and Microbial Technology</i> , 2017, 96, 127-134.	3.2	22
10	Characterization of a Thermostable Short-Chain Alcohol Dehydrogenase from the Hyperthermophilic Archaeon <i>Thermococcus sibiricus</i> . <i>Applied and Environmental Microbiology</i> , 2010, 76, 4096-4098.	3.1	21
11	Experimental and computational studies on the unusual substrate specificity of branched-chain amino acid aminotransferase from <i>Thermoproteus uzoniensis</i> . <i>Archives of Biochemistry and Biophysics</i> , 2016, 607, 27-36.	3.0	20
12	Biochemical and structural insights into PLP fold type IV transaminase from <i>Thermobaculum terrenum</i> . <i>Biochimie</i> , 2019, 158, 130-138.	2.6	19
13	ATP-dependent DNA ligase from <i>Thermococcus</i> sp. 1519 displays a new arrangement of the OB-fold domain. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 1440-1447.	0.7	16
14	The Uncommon Active Site of D-Amino Acid Transaminase from <i>Haliscomenobacter hydrossis</i> : Biochemical and Structural Insights into the New Enzyme. <i>Molecules</i> , 2021, 26, 5053.	3.8	14
15	Identification of branched-chain amino acid aminotransferases active towards (R)-(+)-1-phenylethylamine among PLP fold type IV transaminases. <i>Journal of Biotechnology</i> , 2018, 271, 26-28.	3.8	13
16	Functional characterization of PLP fold type IV transaminase with a mixed type of activity from <i>Haliangium ochraceum</i> . <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2019, 1867, 575-585.	2.3	11
17	Overexpression, purification and crystallization of a thermostable DNA ligase from the archaeon <i>Thermococcus</i> sp. 1519. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2009, 65, 368-371.	0.7	9
18	Expression, purification, crystallization and preliminary crystallographic analysis of a thermostable DNA ligase from the archaeon <i>Thermococcus sibiricus</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2012, 68, 163-165.	0.7	8

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19	Nicotinamidase from the thermophilic archaeon <i>Acidilobus saccharovorans</i> : Structural and functional characteristics. <i>Biochemistry (Moscow)</i> , 2014, 79, 54-61.	1.5	8
20	Structure of the dodecamer of the aminopeptidase APDkam598 from the archaeon <i>Desulfurococcus kamchatkensis</i> . <i>Acta Crystallographica Section F, Structural Biology Communications</i> , 2015, 71, 277-285.	0.8	8
21	Structural characterization of geranylgeranyl pyrophosphate synthase GACE1337 from the hyperthermophilic archaeon <i>Geoglobus acetivorans</i> . <i>Extremophiles</i> , 2018, 22, 877-888.	2.3	7
22	Intramolecular hydrogen bonding in the polyextremophilic short-chain dehydrogenase from the archaeon <i>Thermococcus sibiricus</i> and its close structural homologs. <i>Biochimie</i> , 2015, 118, 82-89.	2.6	6
23	Diaminopelargonic acid transaminase from <i>Psychrobacter cryohalolentis</i> is active towards (S)-(-)-1-phenylethylamine, aldehydes and α -diketones. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 9621-9633.	3.6	6
24	Effects of pH and temperature on (S)-amine activity of transaminase from the cold-adapted bacterium <i>Psychrobacter cryohalolentis</i> . <i>Extremophiles</i> , 2020, 24, 537-549.	2.3	6
25	Structures of β -glycosidase from <i>Acidilobus saccharovorans</i> in complexes with tris and glycerol. <i>Doklady Biochemistry and Biophysics</i> , 2013, 449, 99-101.	0.9	5
26	Characterization of a novel M42 aminopeptidase from crenarchaeon <i>Desulfurococcus kamchatkensis</i> . <i>Doklady Biochemistry and Biophysics</i> , 2012, 442, 30-32.	0.9	4
27	NADP-Dependent Aldehyde Dehydrogenase from Archaeon <i>Pyrobaculum</i> sp.1860: Structural and Functional Features. <i>Archaea</i> , 2016, 2016, 1-14.	2.3	3
28	Studies of Peroxidase Refolding in the Presence of Specific Antibodies. <i>Applied Biochemistry and Microbiology</i> , 2003, 39, 446-453.	0.9	2
29	Expression, purification and crystallization of a thermostable short-chain alcohol dehydrogenase from the archaeon <i>Thermococcus sibiricus</i> . <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2010, 66, 655-657.	0.7	2
30	Counterbalance of Stability and Activity Observed for Thermostable Transaminase from <i>Thermobaculum terrenum</i> in the Presence of Organic Solvents. <i>Catalysts</i> , 2020, 10, 1024.	3.5	2
31	Sodium Chloride-Induced Modulation of the Activity and Thermal Stability of Short-Chain Oxidoreductase from the Archaeon <i>Thermococcus sibiricus</i> . <i>Applied Biochemistry and Biotechnology</i> , 2013, 171, 1877-1889.	2.9	1
32	Probing the role of the residues in the active site of the transaminase from <i>Thermobaculum terrenum</i> . <i>PLoS ONE</i> , 2021, 16, e0255098.	2.5	1
33	Structural features of thermostable short-chain alcohol dehydrogenase from hyperthermophilic archaeon <i>Thermococcus sibiricus</i> . <i>Current Opinion in Biotechnology</i> , 2011, 22, S85.	6.6	0
34	Molecular dynamics study of the structural and dynamic characteristics of the polyextremophilic short-chain dehydrogenase from the <i>Thermococcus sibiricus</i> archaeon and its homologues. <i>AIP Conference Proceedings</i> , 2017, . .	0.4	0
35	Effect of Ketosubstrate on the Product Yield in the Transamination Reaction Catalyzed by Transaminase from <i>Thermoproteus uzoniensis</i> . <i>Doklady Biochemistry and Biophysics</i> , 2020, 490, 5-8.	0.9	0
36	A Puzzling Protein from <i>Variovorax paradoxus</i> Has a PLP Fold Type IV Transaminase Structure and Binds PLP without Catalytic Lysine. <i>Crystals</i> , 2022, 12, 619.	2.2	0