

Constanze Pinske

List of Publications by Year in descending order

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papers

822
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623188

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27
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37
all docs

37
docs citations

37
times ranked

634
citing authors

#	ARTICLE	IF	CITATIONS
1	Anaerobic Formate and Hydrogen Metabolism. <i>EcoSal Plus</i> , 2016, 7, .	2.1	95
2	Metabolic Deficiencies Revealed in the Biotechnologically Important Model Bacterium <i>Escherichia coli</i> BL21(DE3). <i>PLoS ONE</i> , 2011, 6, e22830.	1.1	61
3	Physiology and Bioenergetics of [NiFe]-Hydrogenase 2-Catalyzed H ₂ -Consuming and H ₂ -Producing Reactions in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2015, 197, 296-306.	1.0	60
4	Exploring the directionality of <i>Escherichia coli</i> formate hydrogenlyase: a membrane-bound enzyme capable of fixing carbon dioxide to organic acid. <i>MicrobiologyOpen</i> , 2016, 5, 721-737.	1.2	60
5	The respiratory molybdo-selenoprotein formate dehydrogenases of <i>Escherichia coli</i> have hydrogen: benzyl viologen oxidoreductase activity. <i>BMC Microbiology</i> , 2011, 11, 173.	1.3	55
6	Efficient electron transfer from hydrogen to benzyl viologen by the [NiFe]-hydrogenases of <i>Escherichia coli</i> is dependent on the coexpression of the iron-sulfur cluster-containing small subunit. <i>Archives of Microbiology</i> , 2011, 193, 893-903.	1.0	51
7	Characterization of <i>Escherichia coli</i> [NiFe]-Hydrogenase Distribution During Fermentative Growth at Different pHs. <i>Cell Biochemistry and Biophysics</i> , 2012, 62, 433-440.	0.9	46
8	Zymographic differentiation of [NiFe]-Hydrogenases 1, 2 and 3 of <i>Escherichia coli</i> K-12. <i>BMC Microbiology</i> , 2012, 12, 134.	1.3	40
9	Dependence on the FOF1-ATP synthase for the activities of the hydrogen-oxidizing hydrogenases 1 and 2 during glucose and glycerol fermentation at high and low pH in <i>Escherichia coli</i> . <i>Journal of Bioenergetics and Biomembranes</i> , 2011, 43, 645-650.	1.0	38
10	A-Type Carrier Protein ErpA Is Essential for Formation of an Active Formate-Nitrate Respiratory Pathway in <i>Escherichia coli</i> K-12. <i>Journal of Bacteriology</i> , 2012, 194, 346-353.	1.0	37
11	Delivery of Iron-Sulfur Clusters to the Hydrogen-Oxidizing [NiFe]-Hydrogenases in <i>Escherichia coli</i> Requires the A-Type Carrier Proteins ErpA and IscA. <i>PLoS ONE</i> , 2012, 7, e31755.	1.1	34
12	The importance of iron in the biosynthesis and assembly of [NiFe]-hydrogenases. <i>Biomolecular Concepts</i> , 2014, 5, 55-70.	1.0	21
13	Development of a cell-free system reveals an oxygen-labile step in the maturation of [NiFe]-hydrogenase 2 of <i>Escherichia coli</i> . <i>FEBS Letters</i> , 2010, 584, 4109-4114.	1.3	19
14	Analysis of hydrogenase 1 levels reveals an intimate link between carbon and hydrogen metabolism in <i>Escherichia coli</i> K-12. <i>Microbiology (United Kingdom)</i> , 2012, 158, 856-868.	0.7	18
15	SlyD-dependent nickel delivery limits maturation of [NiFe]-hydrogenases in late-stationary phase <i>Escherichia coli</i> cells. <i>Metallomics</i> , 2015, 7, 683-690.	1.0	17
16	The role of the ferric-uptake regulator Fur and iron homeostasis in controlling levels of the [NiFe]-hydrogenases in <i>Escherichia coli</i> . <i>International Journal of Hydrogen Energy</i> , 2010, 35, 8938-8944.	3.8	14
17	Levels of control exerted by the Isc iron-sulfur cluster system on biosynthesis of the formate hydrogenlyase complex. <i>Microbiology (United Kingdom)</i> , 2013, 159, 1179-1189.	0.7	14
18	The Ferredoxin-Like Proteins HydN and YsaA Enhance Redox Dye-Linked Activity of the Formate Dehydrogenase H Component of the Formate Hydrogenlyase Complex. <i>Frontiers in Microbiology</i> , 2018, 9, 1238.	1.5	12

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19	Bioenergetic aspects of archaeal and bacterial hydrogen metabolism. <i>Advances in Microbial Physiology</i> , 2019, 74, 487-514.	1.0	12
20	Expanding the substrates for a bacterial hydrogenlyase reaction. <i>Microbiology (United Kingdom)</i> , 2017, 163, 649-653.	0.7	12
21	A single amino acid exchange converts FocA into a unidirectional efflux channel for formate. <i>Microbiology (United Kingdom)</i> , 2022, 168, .	0.7	12
22	Iron restriction induces preferential down-regulation of H ₂ -consuming over H ₂ -evolving reactions during fermentative growth of <i>Escherichia coli</i> . <i>BMC Microbiology</i> , 2011, 11, 196.	1.3	10
23	Differential effects of <i>isc</i> operon mutations on the biosynthesis and activity of key anaerobic metalloenzymes in <i>Escherichia coli</i> . <i>Microbiology (United Kingdom)</i> , 2017, 163, 878-890.	0.7	9
24	Integration of an [FeFe]-hydrogenase into the anaerobic metabolism of <i>Escherichia coli</i> . <i>Biotechnology Reports (Amsterdam, Netherlands)</i> , 2015, 8, 94-104.	2.1	8
25	The dual-function chaperone HycH improves assembly of the formate hydrogenlyase complex. <i>Biochemical Journal</i> , 2017, 474, 2937-2950.	1.7	8
26	Delimiting the Function of the C-Terminal Extension of the <i>Escherichia coli</i> [NiFe]-Hydrogenase 2 Large Subunit Precursor. <i>Frontiers in Microbiology</i> , 2019, 10, 2223.	1.5	8
27	Dissection of the Hydrogen Metabolism of the Enterobacterium <i>Trabulsirella guamensis</i> : Identification of a Formate-Dependent and Essential Formate Hydrogenlyase Complex Exhibiting Phylogenetic Similarity to Complex I. <i>Journal of Bacteriology</i> , 2019, 201, .	1.0	8
28	Chromogenic assessment of the three molybdo-selenoprotein formate dehydrogenases in <i>Escherichia coli</i> . <i>Biochemistry and Biophysics Reports</i> , 2015, 1, 62-67.	0.7	7
29	pH and a mixed carbon-substrate spectrum influence FocA- and FocB-dependent, formate-driven H ₂ production in <i>Escherichia coli</i> . <i>FEMS Microbiology Letters</i> , 2018, 365, .	0.7	7
30	Amino acid variants of the HybB membrane subunit of <i>Escherichia coli</i> [NiFe]-hydrogenase support a role in proton transfer. <i>FEBS Letters</i> , 2019, 593, 2194-2203.	1.3	7
31	Influence of <i>C₄Dcu</i> transporters on hydrogenase and formate dehydrogenase activities in stationary phase-grown fermenting <i>Escherichia coli</i> . <i>IUBMB Life</i> , 2020, 72, 1680-1685.	1.5	7
32	Insights Into the Redox Sensitivity of Chloroflexi Hup-Hydrogenase Derived From Studies in <i>Escherichia coli</i> : Merits and Pitfalls of Heterologous [NiFe]-Hydrogenase Synthesis. <i>Frontiers in Microbiology</i> , 2018, 9, 2837.	1.5	4
33	The Extended C-Terminal α -Helix of the HypC Chaperone Restricts Recognition of Large Subunit Precursors by the Hyp-Scaffold Machinery during [NiFe]-Hydrogenase Maturation in <i>Escherichia coli</i> . <i>Journal of Molecular Microbiology and Biotechnology</i> , 2018, 28, 87-97.	1.0	3
34	Susceptibility of the Formate Hydrogenlyase Reaction to the Protonophore CCCP Depends on the Total Hydrogenase Composition. <i>Inorganics</i> , 2020, 8, 38.	1.2	3
35	The N-terminal domains of the paralogous HycE and NuoCD govern assembly of the respective formate hydrogenlyase and NADH dehydrogenase complexes. <i>FEBS Open Bio</i> , 2020, 10, 371-385.	1.0	2