

Matthias Boll

List of Publications by Year in descending order

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93
papers

5,171
citations

94433

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docs citations

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#	ARTICLE	IF	CITATIONS
1	Degradation of dibutyl phthalate by <i>Paenarthrobacter</i> sp. Shss isolated from Saravan landfill, Hyrcanian Forests, Iran. <i>Biodegradation</i> , 2022, 33, 59-70.	3.0	9
2	Enoyl-Coenzyme A Respiration via Formate Cycling in Syntrophic Bacteria. <i>MBio</i> , 2022, 13, e0374021.	4.1	2
3	Genes and enzymes involved in the biodegradation of the quaternary carbon compound pivalate in the denitrifying <i>Thauera humireducens</i> strain <i>PIV</i> . <i>Environmental Microbiology</i> , 2022, , .	3.8	0
4	Structural Evidence for a [4Fe-5S] Intermediate in the Non-Redox Desulfuration of Thiouracil. <i>Angewandte Chemie</i> , 2021, 133, 428-435.	2.0	0
5	Structural Evidence for a [4Fe-5S] Intermediate in the Non-Redox Desulfuration of Thiouracil. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 424-431.	13.8	15
6	Activation of short-chain ketones and isopropanol in sulfate-reducing bacteria. <i>BMC Microbiology</i> , 2021, 21, 50.	3.3	2
7	Functional diversity of prokaryotic HdrA(BC) modules: Role in flavin-based electron bifurcation processes and beyond. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2021, 1862, 148379.	1.0	18
8	Structural Basis of Cyclic 1,3-Diene Forming Acyl-Coenzyme A Dehydrogenases. <i>ChemBioChem</i> , 2021, 22, 3173-3177.	2.6	2
9	The missing enzymatic link in syntrophic methane formation from fatty acids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	7
10	A fully reversible 25-hydroxy steroid kinase involved in oxygen-independent cholesterol side-chain oxidation. <i>Journal of Biological Chemistry</i> , 2021, 297, 101105.	3.4	6
11	Microbial degradation of phthalates: biochemistry and environmental implications. <i>Environmental Microbiology Reports</i> , 2020, 12, 3-15.	2.4	98
12	ATP-dependent hydroxylation of an unactivated primary carbon with water. <i>Nature Communications</i> , 2020, 11, 3906.	12.8	10
13	Tungstoenzymes: Occurrence, Catalytic Diversity and Cofactor Synthesis. <i>Inorganics</i> , 2020, 8, 44.	2.7	29
14	Channeling C1 Metabolism toward S-Adenosylmethionine-Dependent Conversion of Estrogens to Androgens in Estrogen-Degrading Bacteria. <i>MBio</i> , 2020, 11, .	4.1	8
15	Oxygen detoxification by dienoyl-CoA oxidase involving flavin/disulfide cofactors. <i>Molecular Microbiology</i> , 2020, 114, 17-30.	2.5	1
16	An Aerobic Hybrid Phthalate Degradation Pathway via Phthaloyl-Coenzyme A in Denitrifying Bacteria. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	5
17	Catabolic Pathways and Enzymes Involved in the Anaerobic Degradation of Monocyclic Aromatic Compounds. , 2020, , 85-133.		5
18	Catabolic Pathways and Enzymes Involved in the Anaerobic Degradation of Polycyclic Aromatic Hydrocarbons. , 2020, , 135-150.		0

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19	The class II benzoyl-Coenzyme A reductase complex from the sulfate-reducing <i>Desulfosarcina cetonica</i> . <i>Environmental Microbiology</i> , 2019, 21, 4241-4252.	3.8	10
20	One-megadalton metalloenzyme complex in <i>Geobacter metallireducens</i> involved in benzene ring reduction beyond the biological redox window. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 2259-2264.	7.1	32
21	Low potential enzymatic hydride transfer via highly cooperative and inversely functionalized flavin cofactors. <i>Nature Communications</i> , 2019, 10, 2074.	12.8	14
22	Enzymes involved in phthalate degradation in sulphate-reducing bacteria. <i>Environmental Microbiology</i> , 2019, 21, 3601-3612.	3.8	22
23	Evolution of a xenobiotic degradation pathway: formation and capture of the labile phthaloyl-CoA intermediate during anaerobic phthalate degradation. <i>Molecular Microbiology</i> , 2018, 108, 614-626.	2.5	15
24	Functional Characterization of Three Specific Acyl-Coenzyme A Synthetases Involved in Anaerobic Cholesterol Degradation in <i>Sterolibacterium denitrificans</i> Chol1S. <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	11
25	A catalytically versatile benzoyl-CoA reductase, key enzyme in the degradation of methyl- and halobenzoates in denitrifying bacteria. <i>Journal of Biological Chemistry</i> , 2018, 293, 10264-10274.	3.4	22
26	Enantioselective Enzymatic Naphthoyl Ring Reduction. <i>Chemistry - A European Journal</i> , 2018, 24, 12505-12508.	3.3	15
27	Four Molybdenum-Dependent Steroid C-25 Hydroxylases: Heterologous Overproduction, Role in Steroid Degradation, and Application for 25-Hydroxyvitamin D ₃ Synthesis. <i>MBio</i> , 2018, 9, .	4.1	16
28	Catabolic Pathways and Enzymes Involved in the Anaerobic Degradation of Polycyclic Aromatic Hydrocarbons. , 2018, , 1-17.		2
29	Catabolic Pathways and Enzymes Involved in the Anaerobic Degradation of Monocyclic Aromatic Compounds. , 2018, , 1-50.		2
30	An unusual strategy for the anoxic biodegradation of phthalate. <i>ISME Journal</i> , 2017, 11, 224-236.	9.8	61
31	A patchwork pathway for oxygenase-independent degradation of side chain containing steroids. <i>Environmental Microbiology</i> , 2017, 19, 4684-4699.	3.8	28
32	Breaking Benzene Aromaticity—Computational Insights into the Mechanism of the Tungsten-Containing Benzoyl-CoA Reductase. <i>Journal of the American Chemical Society</i> , 2017, 139, 14488-14500.	13.7	19
33	Phthaloyl-Coenzyme A decarboxylase from <i>Thauera chlorobenzoica</i> : the prenylated flavin, K ⁺ and Fe ²⁺ -dependent key enzyme of anaerobic phthalate degradation. <i>Environmental Microbiology</i> , 2017, 19, 3734-3744.	3.8	27
34	Promiscuous Defluorinating Enoyl-CoA Hydratases/Hydrolases Allow for Complete Anaerobic Degradation of 2-Fluorobenzoate. <i>Frontiers in Microbiology</i> , 2017, 8, 2579.	3.5	21
35	ATP-Dependent C—F Bond Cleavage Allows the Complete Degradation of 4-Fluoroaromatics without Oxygen. <i>MBio</i> , 2016, 7, .	4.1	35
36	ATP-Dependent Electron Activation Module of Benzoyl-Coenzyme A Reductase from the Hyperthermophilic Archaeon <i>Ferroglobus placidus</i> . <i>Biochemistry</i> , 2016, 55, 5578-5586.	2.5	11

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37	25-Hydroxyvitamin ₃ Synthesis by Enzymatic Steroid Side-Chain Hydroxylation with Water. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 1881-1884.	13.8	20
38	25-Hydroxyvitamin ₃ Synthesis by Enzymatic Steroid Side-Chain Hydroxylation with Water. <i>Angewandte Chemie</i> , 2016, 128, 1913-1916.	2.0	2
39	Fermentative Cyclohexane Carboxylate Formation in <i>Syntrophus aciditrophicus</i> . <i>Journal of Molecular Microbiology and Biotechnology</i> , 2016, 26, 165-179.	1.0	11
40	Anaerobic Microbial Degradation of Hydrocarbons: From Enzymatic Reactions to the Environment. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2016, 26, 5-28.	1.0	615
41	Ethylbenzene Dehydrogenase and Related Molybdenum Enzymes Involved in Oxygen-Independent Alkyl Chain Hydroxylation. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2016, 26, 45-62.	1.0	50
42	Structure and Function of the Unusual Tungsten Enzymes Acetylene Hydratase and Class II Benzoyl-Coenzyme A Reductase. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2016, 26, 119-137.	1.0	29
43	Anaerobic Degradation of Benzene and Polycyclic Aromatic Hydrocarbons. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2016, 26, 92-118.	1.0	218
44	Functional Gene Markers for Fumarate-Adding and Dearomatizing Key Enzymes in Anaerobic Aromatic Hydrocarbon Degradation in Terrestrial Environments. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2016, 26, 180-194.	1.0	52
45	Two distinct old yellow enzymes are involved in naphthyl ring reduction during anaerobic naphthalene degradation. <i>Molecular Microbiology</i> , 2015, 95, 162-172.	2.5	39
46	Enzymes of the benzoyl-coenzyme A degradation pathway in the hyperthermophilic archaeon <i>Ferroplasma acidiphilum</i> . <i>Environmental Microbiology</i> , 2015, 17, 3289-3300.	3.8	18
47	Structural basis of enzymatic benzene ring reduction. <i>Nature Chemical Biology</i> , 2015, 11, 586-591.	8.0	52
48	Unraveling the Specific Regulation of the Central Pathway for Anaerobic Degradation of 3-Methylbenzoate. <i>Journal of Biological Chemistry</i> , 2015, 290, 12165-12183.	3.4	13
49	Glutaryl-coenzyme A dehydrogenase from <i>Geobacter metallireducens</i> interaction with electron transferring flavoprotein and kinetic basis of unidirectional catalysis. <i>FEBS Journal</i> , 2014, 281, 5120-5131.	4.7	6
50	Anaerobic degradation of homocyclic aromatic compounds via arylcarboxyl-coenzyme A esters: organisms, strategies and key enzymes. <i>Environmental Microbiology</i> , 2014, 16, 612-627.	3.8	156
51	A PCR-based assay for the detection of anaerobic naphthalene degradation. <i>FEMS Microbiology Letters</i> , 2014, 354, 55-59.	1.8	18
52	Enzymes Involved in a Novel Anaerobic Cyclohexane Carboxylic Acid Degradation Pathway. <i>Journal of Bacteriology</i> , 2014, 196, 3667-3674.	2.2	26
53	The Benzoyl-Coenzyme A Reductase and 2-Hydroxyacyl-Coenzyme A Dehydratase Radical Enzyme Family. <i>ChemBioChem</i> , 2014, 15, 2188-2194.	2.6	40
54	Characterization of the <i>mbd</i> cluster encoding the anaerobic 3-methylbenzoyl-CoA central pathway. <i>Environmental Microbiology</i> , 2013, 15, 148-166.	3.8	37

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55	Identification and characterization of 2-naphthoyl-coenzyme A reductase, the prototype of a novel class of dearomatizing reductases. <i>Molecular Microbiology</i> , 2013, 88, 1032-1039.	2.5	52
56	Cyclohexanecarboxyl-Coenzyme A (CoA) and Cyclohex-1-ene-1-Carboxyl-CoA Dehydrogenases, Two Enzymes Involved in the Fermentation of Benzoate and Crotonate in <i>Syntrophus aciditrophicus</i> . <i>Journal of Bacteriology</i> , 2013, 195, 3193-3200.	2.2	27
57	ATP-dependent/independent enzymatic ring reductions involved in the anaerobic catabolism of naphthalene. <i>Environmental Microbiology</i> , 2013, 15, 1832-1841.	3.8	35
58	Identification and Characterization of a Succinyl-Coenzyme A (CoA):Benzoate CoA Transferase in <i>Geobacter metallireducens</i> . <i>Journal of Bacteriology</i> , 2012, 194, 2501-2508.	2.2	25
59	Anaerobic degradation of 4-methylbenzoate via a specific 4-methylbenzoyl-CoA pathway. <i>Environmental Microbiology</i> , 2012, 14, 1118-1132.	3.8	27
60	Microbial degradation of aromatic compounds – from one strategy to four. <i>Nature Reviews Microbiology</i> , 2011, 9, 803-816.	28.6	952
61	Occurrence, genes and expression of the W/Se-containing class II benzoyl-coenzyme A reductases in anaerobic bacteria. <i>Environmental Microbiology</i> , 2011, 13, 696-709.	3.8	65
62	Enzymes involved in the anaerobic degradation of <i>meta</i> -substituted halobenzoates. <i>Molecular Microbiology</i> , 2011, 82, 758-769.	2.5	39
63	Conversion of a decarboxylating to a non-decarboxylating glutaryl-coenzyme A dehydrogenase by site-directed mutagenesis. <i>FEBS Letters</i> , 2011, 585, 1317-1321.	2.8	11
64	Combined Application of PCR-Based Functional Assays for the Detection of Aromatic-Compound-Degrading Anaerobes. <i>Applied and Environmental Microbiology</i> , 2011, 77, 5056-5061.	3.1	55
65	Reversible Biological Birch Reduction at an Extremely Low Redox Potential. <i>Journal of the American Chemical Society</i> , 2010, 132, 9850-9856.	13.7	85
66	Structural Basis for Promoting and Preventing Decarboxylation in Glutaryl-Coenzyme A Dehydrogenases. <i>Biochemistry</i> , 2010, 49, 5350-5357.	2.5	14
67	Decarboxylating and Nondecarboxylating Glutaryl-Coenzyme A Dehydrogenases in the Aromatic Metabolism of Obligately Anaerobic Bacteria. <i>Journal of Bacteriology</i> , 2009, 191, 4401-4409.	2.2	40
68	Differential Membrane Proteome Analysis Reveals Novel Proteins Involved in the Degradation of Aromatic Compounds in <i>Geobacter metallireducens</i> . <i>Molecular and Cellular Proteomics</i> , 2009, 8, 2159-2169.	3.8	25
69	Identification and characterization of the tungsten-containing class of benzoyl-coenzyme A reductases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 17687-17692.	7.1	112
70	Purification and Characterization of Active-Site Components of the Putative <i>p</i> -Cresol Methylhydroxylase Membrane Complex from <i>Geobacter metallireducens</i> . <i>Journal of Bacteriology</i> , 2008, 190, 6493-6500.	2.2	25
71	6-Oxocyclohex-1-ene-1-carboxyl-coenzyme A hydrolases from obligately anaerobic bacteria: characterization and identification of its gene as a functional marker for aromatic compounds degrading anaerobes. <i>Environmental Microbiology</i> , 2008, 10, 1547-1556.	3.8	99
72	Mechanism of Enzymatic Birch Reduction: Stereochemical Course and Exchange Reactions of Benzoyl-CoA Reductase. <i>Journal of the American Chemical Society</i> , 2008, 130, 14050-14051.	13.7	46

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73	Aromatizing Cyclohexa-1,5-diene-1-carbonyl-Coenzyme A Oxidase. <i>Journal of Biological Chemistry</i> , 2008, 283, 20713-20721.	3.4	30
74	Cyclohexa-1,5-Diene-1-Carbonyl-Coenzyme A (CoA) Hydratases of <i>Geobacter metallireducens</i> and <i>Syntrophus aciditrophicus</i> : Evidence for a Common Benzoyl-CoA Degradation Pathway in Facultative and Strict Anaerobes. <i>Journal of Bacteriology</i> , 2007, 189, 1055-1060.	2.2	64
75	Gene clusters involved in anaerobic benzoate degradation of <i>Geobacter metallireducens</i> . <i>Molecular Microbiology</i> , 2005, 58, 1238-1252.	2.5	147
76	Unusual reactions involved in anaerobic metabolism of phenolic compounds. <i>Biological Chemistry</i> , 2005, 386, 989-997.	2.5	45
77	Key enzymes in the anaerobic aromatic metabolism catalysing Birch-like reductions. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2005, 1707, 34-50.	1.0	76
78	Dearomatizing Benzene Ring Reductases. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2005, 10, 132-142.	1.0	92
79	The <i>bzd</i> Gene Cluster, Coding for Anaerobic Benzoate Catabolism, in <i>Azoarcus</i> sp. Strain CIB. <i>Journal of Bacteriology</i> , 2004, 186, 5762-5774.	2.2	111
80	Selenocysteine-Containing Proteins in Anaerobic Benzoate Metabolism of <i>Desulfococcus multivorans</i> . <i>Journal of Bacteriology</i> , 2004, 186, 2156-2163.	2.2	69
81	Structure of a Xanthine Oxidase-Related 4-Hydroxybenzoyl-CoA Reductase with an Additional [4Fe-4S] Cluster and an Inverted Electron Flow. <i>Structure</i> , 2004, 12, 2249-2256.	3.3	62
82	Substrate Binding and Reduction of Benzoyl-CoA Reductase: Evidence for Nucleotide-Dependent Conformational Changes. <i>Biochemistry</i> , 2004, 43, 1376-1385.	2.5	20
83	Properties of 2-Oxoglutarate:Ferredoxin Oxidoreductase from <i>Thauera aromatica</i> and Its Role in Enzymatic Reduction of the Aromatic Ring. <i>Journal of Bacteriology</i> , 2002, 184, 3975-3983.	2.2	51
84	A Birch-like Mechanism in Enzymatic Benzoyl-CoA Reduction: A Kinetic Study of Substrate Analogues Combined with an ab Initio Model. <i>Biochemistry</i> , 2002, 41, 1752-1758.	2.5	57
85	Single Turnover EPR Studies of Benzoyl-CoA Reductase. <i>Biochemistry</i> , 2001, 40, 7612-7620.	2.5	24
86	Redox Centers of 4-Hydroxybenzoyl-CoA Reductase, a Member of the Xanthine Oxidase Family of Molybdenum-containing Enzymes. <i>Journal of Biological Chemistry</i> , 2001, 276, 47853-47862.	3.4	37
87	Mechanism of ATP-driven electron transfer catalyzed by the benzene ring-reducing enzyme benzoyl-CoA reductase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 13619-13624.	7.1	44
88	EPR and Mössbauer Studies of Benzoyl-CoA Reductase. <i>Journal of Biological Chemistry</i> , 2000, 275, 31857-31868.	3.4	38
89	Nonaromatic Products from Anoxic Conversion of Benzoyl-CoA with Benzoyl-CoA Reductase and Cyclohexa-1,5-diene-1-carbonyl-CoA Hydratase. <i>Journal of Biological Chemistry</i> , 2000, 275, 21889-21895.	3.4	56
90	Identification and characterization of the natural electron donor ferredoxin and of FAD as a possible prosthetic group of benzoyl-CoA reductase (dearomatizing), a key enzyme of anaerobic aromatic metabolism. <i>FEBS Journal</i> , 1998, 251, 946-954.	0.2	58

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91	Genes coding for the benzoyl-CoA pathway of anaerobic aromatic metabolism in the bacterium <i>Thauera aromatica</i> . FEBS Journal, 1998, 256, 148-154.	0.2	81
92	Benzoyl-CoA Reductase (Dearomatizing), A Key Enzyme of Anaerobic Aromatic Metabolism. FEBS Journal, 1997, 244, 840-851.	0.2	88
93	Benzoyl-Coenzyme A Reductase (Dearomatizing), a Key Enzyme of Anaerobic Aromatic Metabolism. FEBS Journal, 1995, 234, 921-933.	0.2	205