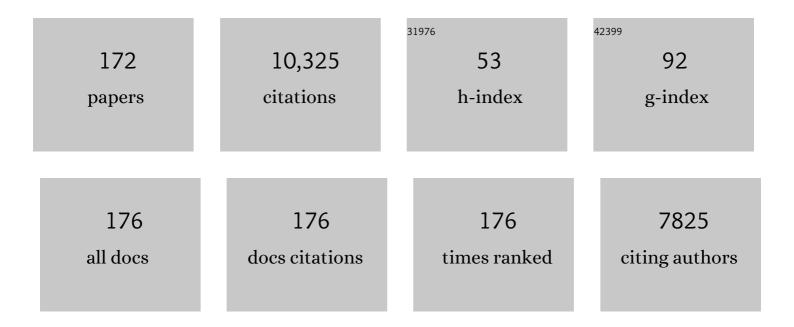
Martin J Warren

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Exploring the onset of <scp>B₁₂</scp> â€based mutualisms using a recently evolved <scp><i>Chlamydomonas</i></scp> auxotroph and <scp>B₁₂</scp> â€producing bacteria. Environmental Microbiology, 2022, 24, 3134-3147.	3.8	14
2	Calculating metalation in cells reveals CobW acquires Coll for vitamin B12 biosynthesis while related proteins prefer ZnII. Nature Communications, 2021, 12, 1195.	12.8	32
3	Red Fluorescence of European Hedgehog (Erinaceus europaeus) Spines Results from Free-Base Porphyrins of Potential Microbial Origin. Journal of Chemical Ecology, 2021, 47, 588-596.	1.8	10
4	Plasmodium falciparum hydroxymethylbilane synthase does not house any cosynthase activity within the haem biosynthetic pathway. Microbiology (United Kingdom), 2021, 167, .	1.8	0
5	Editorial overview: Bacterial microcompartments to the fore as metabolism is put in its place. Current Opinion in Microbiology, 2021, 64, 159-161.	5.1	0
6	Replacement of the Cobalt Center of Vitamin B ₁₂ by Nickel: Nibalamin and Nibyric Acid Prepared from Metalâ€Free B ₁₂ â€Ligands Hydrogenobalamin and Hydrogenobyric Acid. Angewandte Chemie - International Edition, 2020, 59, 20129-20136.	13.8	18
7	Replacement of the Cobalt Center of Vitamin B ₁₂ by Nickel: Nibalamin and Nibyric Acid Prepared from Metalâ€Free B ₁₂ â€Ligands Hydrogenobalamin and Hydrogenobyric Acid. Angewandte Chemie, 2020, 132, 20304-20311.	2.0	2
8	Effect of metabolosome encapsulation peptides on enzyme activity, coaggregation, incorporation, and bacterial microcompartment formation. MicrobiologyOpen, 2020, 9, e1010.	3.0	14
9	Biosynthesis of the modified tetrapyrroles—the pigments of life. Journal of Biological Chemistry, 2020, 295, 6888-6925.	3.4	170
10	Zinc Substitution of Cobalt in Vitaminâ€B12: Zincobyric acid and Zincobalamin as Luminescent Structural B12â€Mimics. Angewandte Chemie - International Edition, 2019, 58, 14568-14572.	13.8	25
11	Zinc Substitution of Cobalt in Vitaminâ€B12: Zincobyric acid and Zincobalamin as Luminescent Structural B12â€Mimics. Angewandte Chemie, 2019, 131, 14710-14714.	2.0	4
12	Bacterial sensors define intracellular free energies for correct enzyme metalation. Nature Chemical Biology, 2019, 15, 241-249.	8.0	112
13	Bacterial Microcompartment-Mediated Ethanolamine Metabolism in Escherichia coli Urinary Tract Infection. Infection and Immunity, 2019, 87, .	2.2	21
14	The Hydrogenobyric Acid Structure Reveals the Corrin Ligand as an Entatic State Module Empowering B ₁₂ Cofactors for Catalysis. Angewandte Chemie - International Edition, 2019, 58, 10756-10760.	13.8	30
15	Biotechnological Advances in Bacterial Microcompartment Technology. Trends in Biotechnology, 2019, 37, 325-336.	9.3	43
16	A Generic Selfâ€Assembly Process in Microcompartments and Synthetic Protein Nanotubes. Small, 2018, 14, e1704020.	10.0	43
17	Engineered synthetic scaffolds for organizing proteins within the bacterial cytoplasm. Nature Chemical Biology, 2018, 14, 142-147.	8.0	128

18 Understanding the Control of a Vitamin B<inf>12</inf> Riboswitch. , 2018, , .

2

#	Article	IF	CITATIONS
19	De novo targeting to the cytoplasmic and luminal side of bacterial microcompartments. Nature Communications, 2018, 9, 3413.	12.8	39

$_{20}$ Construction of Fluorescent Analogs to Follow the Uptake and Distribution of Cobalamin (Vitamin) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 $_{5.2}$

21	<i>Staphylococcus aureus</i> haem biosynthesis and acquisition pathways are linked through haem monooxygenase IsdG. Molecular Microbiology, 2018, 109, 385-400.	2.5	18
22	Vitamin B12. Advances in Food and Nutrition Research, 2018, 83, 215-279.	3.0	105
23	Bacterial microcompartmentâ€directed polyphosphate kinase promotes stable polyphosphate accumulation in <i>E. coli</i> . Biotechnology Journal, 2017, 12, 1600415.	3.5	53
24	Prokaryotic Heme Biosynthesis: Multiple Pathways to a Common Essential Product. Microbiology and Molecular Biology Reviews, 2017, 81, .	6.6	236
25	Elucidation of the biosynthesis of the methane catalyst coenzyme F430. Nature, 2017, 543, 78-82.	27.8	104
26	Construction of Recombinant Pdu Metabolosome Shells for Small Molecule Production in <i>Corynebacterium glutamicum</i> . ACS Synthetic Biology, 2017, 6, 2145-2156.	3.8	41
27	Desulfovibrio vulgaris CbiK P cobaltochelatase: evolution of a haem binding protein orchestrated by the incorporation of two histidine residues. Environmental Microbiology, 2017, 19, 106-118.	3.8	9
28	Classification of polyhedral shapes from individual anisotropically resolved cryo-electron tomography reconstructions. BMC Bioinformatics, 2016, 17, 234.	2.6	4
29	Effect of bio-engineering on size, shape, composition and rigidity of bacterial microcompartments. Scientific Reports, 2016, 6, 36899.	3.3	31
30	Total Synthesis, Structure, and Biological Activity of Adenosylrhodibalamin, the Nonâ€Natural Rhodium Homologue of Coenzyme B ₁₂ . Angewandte Chemie - International Edition, 2016, 55, 11281-11286.	13.8	42
31	Signalling intertextuality in business emails. English for Specific Purposes, 2016, 42, 26-37.	2.8	13
32	Employing bacterial microcompartment technology to engineer a shell-free enzyme-aggregate for enhanced 1,2-propanediol production in Escherichia coli. Metabolic Engineering, 2016, 36, 48-56.	7.0	74
33	Do collocational frameworks have local grammars?. International Journal of Corpus Linguistics, 2016, 21, 1-27.	1.4	7
34	Crystal structure of CobK reveals strand-swapping between Rossmann-fold domains and molecular basis of the reduced precorrin product trap. Scientific Reports, 2015, 5, 16943.	3.3	7
35	<scp><i>S</i></scp> <i>taphylococcus aureus</i> haem biosynthesis: characterisation of the enzymes involved in final steps of the pathway. Molecular Microbiology, 2015, 97, 472-487.	2.5	66
36	FAD binding, cobinamide binding and active site communication in the corrin reductase (CobR). Bioscience Reports, 2014, 34, .	2.4	6

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37	Identification and characterization of the â€~missing' terminal enzyme for siroheme biosynthesis in αâ€proteobacteria. Molecular Microbiology, 2014, 92, 153-163.	2.5	20
38	Structural Insights into Higher Order Assembly and Function of the Bacterial Microcompartment Protein PduA. Journal of Biological Chemistry, 2014, 289, 22377-22384.	3.4	77
39	Recent advances in the biosynthesis of modified tetrapyrroles: the discovery of an alternative pathway for the formation of heme and heme d 1. Cellular and Molecular Life Sciences, 2014, 71, 2837-2863.	5.4	54
40	Towards a cell factory for vitamin B12 production in Bacillus megaterium: bypassing of the cobalamin riboswitch control elements. New Biotechnology, 2014, 31, 553-561.	4.4	38
41	The structure, function and properties of sirohaem decarboxylase – an enzyme with structural homology to a transcription factor family that is part of the alternative haem biosynthesis pathway. Molecular Microbiology, 2014, 93, 247-261.	2.5	14
42	Solution Structure of a Bacterial Microcompartment Targeting Peptide and Its Application in the Construction of an Ethanol Bioreactor. ACS Synthetic Biology, 2014, 3, 454-465.	3.8	175
43	"Preparation is everything― Meetings in professional contexts in Hong Kong. English for Specific Purposes, 2014, 36, 12-26.	2.8	6
44	Characterisation of Desulfovibrio vulgaris haem b synthase, a radical SAM family member. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 1238-1247.	2.3	23
45	Bacterial microcompartments moving into a synthetic biological world. Journal of Biotechnology, 2013, 163, 273-279.	3.8	92
46	"Just spoke to …― The types and directionality of intertextuality in professional discourse. English for Specific Purposes, 2013, 32, 12-24.	2.8	32
47	Characterization of the Enzyme CbiH60 Involved in Anaerobic Ring Contraction of the Cobalamin (Vitamin B12) Biosynthetic Pathway. Journal of Biological Chemistry, 2013, 288, 297-305.	3.4	19
48	Elucidation of the anaerobic pathway for the corrin component of cobalamin (vitamin B) Tj ETQq0 0 0 rgBT /Ove 2013, 110, 14906-14911.	erlock 101 7.1	f 50 307 Td (88
49	Bacillus megaterium Has Both a Functional BluB Protein Required for DMB Synthesis and a Related Flavoprotein That Forms a Stable Radical Species. PLoS ONE, 2013, 8, e55708.	2.5	20
50	Characterization of the evolutionarily conserved iron–sulfur cluster of sirohydrochlorin ferrochelatase from <i>Arabidopsis thaliana</i> . Biochemical Journal, 2012, 444, 227-237.	3.7	19
51	The anaerobic biosynthesis of vitamin B12. Biochemical Society Transactions, 2012, 40, 581-586.	3.4	75
52	Sulfate-Reducing Bacteria Reveal a New Branch of Tetrapyrrole Metabolism. Advances in Microbial Physiology, 2012, 61, 267-295.	2.4	12
53	An enzyme-trap approach allows isolation of intermediates in cobalamin biosynthesis. Nature Chemical Biology, 2012, 8, 933-940.	8.0	62
54	Characterization of <i>Cupriavidus metallidurans</i> CYP116B1 – A thiocarbamate herbicide oxygenating P450–phthalate dioxygenase reductase fusion protein. FEBS Journal, 2012, 279, 1675-1693.	4.7	37

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55	Bacterial ferrochelatase turns human: Tyr13 determines the apparent metal specificity of Bacillus subtilis ferrochelatase. Journal of Biological Inorganic Chemistry, 2011, 16, 235-242.	2.6	14
56	Structure of PduT, a trimeric bacterial microcompartment protein with a 4Fe–4S cluster-binding site. Acta Crystallographica Section D: Biological Crystallography, 2011, 67, 91-96.	2.5	62
57	Evolution in a family of chelatases facilitated by the introduction of active site asymmetry and protein oligomerization. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 97-102.	7.1	43
58	Molecular hijacking of siroheme for the synthesis of heme and <i>d</i> ₁ heme. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 18260-18265.	7.1	121
59	NirJ, a radical SAM family member of the <i>d</i> ₁ heme biogenesis cluster. FEBS Letters, 2010, 584, 2461-2466.	2.8	27
60	Metabolic engineering of cobalamin (vitamin B ₁₂) production in <i>Bacillus megaterium</i> . Microbial Biotechnology, 2010, 3, 24-37.	4.2	75
61	Cloning, purification and preliminary crystallographic analysis of cobalamin methyltransferases fromRhodobacter capsulatus. Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 1652-1656.	0.7	2
62	NirF is a periplasmic protein that binds <i>d</i> ₁ heme as part of its essential role in <i>d</i> ₁ heme biogenesis. FEBS Journal, 2010, 277, 4944-4955.	4.7	16
63	A Novel Pathway for the Biosynthesis of Heme in <i>Archaea</i> : Genome-Based Bioinformatic Predictions and Experimental Evidence. Archaea, 2010, 2010, 1-15.	2.3	56
64	A short story about a big magic bug. Bioengineered Bugs, 2010, 1, 85-91.	1.7	53
65	Synthesis of Empty Bacterial Microcompartments, Directed Organelle Protein Incorporation, and Evidence of Filament-Associated Organelle Movement. Molecular Cell, 2010, 38, 305-315.	9.7	200
66	Characterisation of PduS, the pdu Metabolosome Corrin Reductase, and Evidence of Substructural Organisation within the Bacterial Microcompartment. PLoS ONE, 2010, 5, e14009.	2.5	36
67	What can a corpus tell us about multi-word units?. , 2010, , 212-226.		17
68	Demonstration That CobG, the Monooxygenase Associated with the Ring Contraction Process of the Aerobic Cobalamin (Vitamin B12) Biosynthetic Pathway, Contains an Fe-S Center and a Mononuclear Non-heme Iron Center. Journal of Biological Chemistry, 2009, 284, 4796-4805.	3.4	16
69	The <i>Pseudomonasâ€faeruginosa nirE</i> gene encodes the <i> S</i> â€adenosylâ€ <scp>L</scp> â€methionineâ€dependent uroporphyrinogen III methyltransferase required for heme <i>d</i> ₁ biosynthesis. FEBS Journal, 2009, 276, 5973-5982.	4.7	33
70	<i>d</i> ₁ â€fhaem biogenesis – assessing the roles of three <i>nir</i> gene products. FEBS Journal, 2009, 276, 6399-6411.	4.7	40
71	Uncovering the Extent of the Phraseological Tendency: Towards a Systematic Analysis of Concgrams. Applied Linguistics, 2009, 30, 236-252.	2.4	128
72	Iron–sulfur cluster dynamics in biotin synthase: A new [2Fe–2S]1+ cluster. Biochemical and Biophysical Research Communications, 2009, 381, 487-490.	2.1	7

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73	The AAA+ motor complex of subunits CobS and CobT of cobaltochelatase visualized by single particle electron microscopy. Journal of Structural Biology, 2009, 167, 227-234.	2.8	38
74	Structure of a trimeric bacterial microcompartment shell protein, EtuB, associated with ethanol utilization in <i>Clostridium kluyveri</i> . Biochemical Journal, 2009, 423, 199-207.	3.7	80
75	Functional characterization of the early steps of tetrapyrrole biosynthesis and modification in <i>Desulfovibrio vulgaris</i> Hildenborough. Biochemical Journal, 2009, 420, 317-326.	3.7	37
76	Biochemical and Structural Insights into Bacterial Organelle Form and Biogenesis. Journal of Biological Chemistry, 2008, 283, 14366-14375.	3.4	133
77	<i>Lactobacillus reuteri</i> DSM 20016 Produces Cobalamin-Dependent Diol Dehydratase in Metabolosomes and Metabolizes 1,2-Propanediol by Disproportionation. Journal of Bacteriology, 2008, 190, 4559-4567.	2.2	131
78	Two Distinct Roles for Two Functional Cobaltochelatases (CbiK) in <i>Desulfovibrio vulgaris</i> Hildenborough. Biochemistry, 2008, 47, 5851-5857.	2.5	23
79	Identification, Characterization, and Structure/Function Analysis of a Corrin Reductase Involved in Adenosylcobalamin Biosynthesis. Journal of Biological Chemistry, 2008, 283, 10813-10821.	3.4	29
80	Structure and function of SirC from Bacillus megaterium: a metal-binding precorrin-2 dehydrogenase. Biochemical Journal, 2008, 415, 257-263.	3.7	19
81	Biosynthesis and Use of Cobalamin (B ₁₂). EcoSal Plus, 2008, 3, .	5.4	18
82	7. // -> ONE country two SYStems //: The discourse intonation patterns of word associations. Studies in Corpus Linguistics, 2008, , 135-153.	0.2	1
83	Enzyme Sequence and Its Relationship to Hyperbaric Stability of Artificial and Natural Fish Lactate Dehydrogenases. PLoS ONE, 2008, 3, e2042.	2.5	34
84	Checking Understandings: Comparing Textbooks and a Corpus of Spoken English in Hong Kong. Language Awareness, 2007, 16, 190-207.	1.3	15
85	Elucidation of Substrate Specificity in the Cobalamin (Vitamin B12) Biosynthetic Methyltransferases. Journal of Biological Chemistry, 2007, 282, 23957-23969.	3.4	26
86	Concgramming: A computer driven approach to learning the phraseology of English. ReCALL, 2007, 19, 287-306.	5.2	18
87	Evolution of enzymes and pathways for the biosynthesis of cofactors. Natural Product Reports, 2007, 24, 972.	10.3	62
88	Iron–sulfur proteins as initiators of radical chemistry. Natural Product Reports, 2007, 24, 1027.	10.3	36
89	Roles of vitamins B5, B8, B9, B12 and molybdenum cofactor at cellular and organismal levels. Natural Product Reports, 2007, 24, 949.	10.3	42
90	Metal and cofactor insertion. Natural Product Reports, 2007, 24, 963.	10.3	38

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91	{ / [Oh] Not a < ^ Lot > }: Discourse Intonation and Vague Language. , 2007, , 182-197.		3
92	Biotin Synthase Mechanism: Mutagenesis of the YNHNLD Conserved Motifâ€,‡. Biochemistry, 2006, 45, 12274-12281.	2.5	15
93	A study of the nuclear trafficking of the splicing factor protein PRPF31 linked to autosomal dominant retinitis pigmentosa (ADRP). Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2006, 1762, 304-311.	3.8	18
94	Interactions of RadB, a DNA Repair Protein in Archaea, with DNA and ATP. Journal of Molecular Biology, 2006, 358, 46-56.	4.2	38
95	Characterization of retinaldehyde dehydrogenase 3. Biochemical Journal, 2006, 394, 67-75.	3.7	39
96	Crystal Structure of the Vitamin B12 Biosynthetic Cobaltochelatase, CbiXS, from Archaeoglobus Fulgidus. Journal of Structural and Functional Genomics, 2006, 7, 37-50.	1.2	30
97	From n-gram to skipgram to concgram. International Journal of Corpus Linguistics, 2006, 11, 411-433.	1.4	244
98	The Substrate Radical of Escherichia coli Oxygen-independent Coproporphyrinogen III Oxidase HemN. Journal of Biological Chemistry, 2006, 281, 15727-15734.	3.4	73
99	Finding the final pieces of the vitamin B12 biosynthetic jigsaw. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 4799-4800.	7.1	18
100	Algae Need Their Vitamins. Eukaryotic Cell, 2006, 5, 1175-1183.	3.4	385
101	Algae acquire vitamin B12 through a symbiotic relationship with bacteria. Nature, 2005, 438, 90-93.	27.8	1,258
102	Crystallization and preliminary structure analysis of CobE, an essential protein of cobalamin (vitamin) Tj ETQq0 0 442-444.	0 rgBT /0 0.7	verlock 10 Tf 2
103	//CAN i help you //. International Journal of Corpus Linguistics, 2005, 10, 85-107.	1.4	9
104	Identification and Characterization of the Terminal Enzyme of Siroheme Biosynthesis from Arabidopsis thaliana. Journal of Biological Chemistry, 2005, 280, 4713-4721.	3.4	42
105	Identification and Characterization of a Novel Vitamin B12 (Cobalamin) Biosynthetic Enzyme (CobZ) from Rhodobacter capsulatus, Containing Flavin, Heme, and Fe-S Cofactors. Journal of Biological Chemistry, 2005, 280, 1086-1094.	3.4	52
106	Peer assessment of language proficiency. Language Testing, 2005, 22, 93-121.	3.2	126
107	King George III and porphyria: an elemental hypothesis and investigation. Lancet, The, 2005, 366, 332-335.	13.7	38
108	The dimeric form of flavocytochrome P450 BM3 is catalytically functional as a fatty acid hydroxylase. FEBS Letters, 2005, 579, 5582-5588.	2.8	107

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109	Purification, characterisation and intracellular localisation of aryl hydrocarbon interacting protein-like 1 (AIPL1) and effects of mutations associated with inherited retinal dystrophies. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2004, 1690, 141-149.	3.8	8
110	Structure/Function Studies on a S-Adenosyl-l-methionine-dependent Uroporphyrinogen III C Methyltransferase (SUMT), a Key Regulatory Enzyme of Tetrapyrrole Biosynthesis. Journal of Molecular Biology, 2004, 344, 419-433.	4.2	56
111	CysG structure reveals tetrapyrrole-binding features and novel regulation of siroheme biosynthesis. Nature Structural and Molecular Biology, 2003, 10, 1064-1073.	8.2	78
112	The language learner as language researcher: putting corpus linguistics on the timetable. System, 2003, 31, 173-186.	3.4	61
113	Characterization of the Cobaltochelatase CbiXL. Journal of Biological Chemistry, 2003, 278, 41900-41907.	3.4	49
114	A Story of Chelatase Evolution. Journal of Biological Chemistry, 2003, 278, 22388-22395.	3.4	93
115	X-ray structure of a putative reaction intermediate of 5-aminolaevulinic acid dehydratase. Biochemical Journal, 2003, 373, 733-738.	3.7	29
116	Identification and functional analysis of enzymes required for precorrin-2 dehydrogenation and metal ion insertion in the biosynthesis of sirohaem and cobalamin in Bacillus megaterium. Biochemical Journal, 2003, 370, 505-516.	3.7	93
117	Indirectness, inexplicitness and vagueness made clearer. Pragmatics, 2003, 13, 381-400.	1.0	35
118	The biosynthesis of adenosylcobalamin (vitamin B12). Natural Product Reports, 2002, 19, 390-412.	10.3	409
119	Characterisation of two genes for guanylate cyclase activator protein (GCAP1 and GCAP2) in the Japanese pufferfish, Fugu rubripes. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2002, 1577, 73-80.	2.4	1
120	The structure of Saccharomyces cerevisiae Met8p, a bifunctional dehydrogenase and ferrochelatase. EMBO Journal, 2002, 21, 2068-2075.	7.8	63
121	Isolation and characterization of 14 additional genes specifying the anaerobic biosynthesis of cobalamin (vitamin B12) in Propionibacterium freudenreichii (P. shermanii) The GenBank accession numbers for the sequences reported in this paper are AY033235, AY033236, U13043 and U51164 Microbiology (United Kingdom), 2002, 148, 1845-1853.	1.8	46
122	Identification and Functional Consequences of a New Mutation (E155G) in the Gene for GCAP1 That Causes Autosomal Dominant Cone Dystrophy. American Journal of Human Genetics, 2001, 69, 471-480.	6.2	115
123	â€~She kows more about Hong Kong than you do isn't it': tags in Hong Kong conversational English. Journal of Pragmatics, 2001, 33, 1419-1439.	1.5	22
124	The use of vague language in intercultural conversations in Hong Kong. English World-wide, 2001, 22, 81-104.	0.5	32
125	Optimization of Met8p crystals through protein-storage buffer manipulation. Acta Crystallographica Section D: Biological Crystallography, 2001, 57, 867-869.	2.5	2
126	The Functions of Actually in a Corpus of Intercultural Conversations. International Journal of Corpus Linguistics, 2001, 6, 257-280.	1.4	12

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127	The Enigma of Cobalamin (Vitamin B12) Biosynthesis inPorphyromonas gingivalis. Journal of Biological Chemistry, 2000, 275, 40316-40323.	3.4	95
128	Characterization of the Rhodobacter sphaeroides 5-aminolaevulinic acid synthase isoenzymes, HemA and HemT, isolated from recombinant Escherichia coli. FEBS Journal, 1999, 265, 290-299.	0.2	33
129	The schiff base complex of yeast 5â€aminolaevulinic acid dehydratase with laevulinic acid. Protein Science, 1999, 8, 1250-1256.	7.6	51
130	Vitamin B12: Insights into Biosynthesis's Mount Improbable. Bioorganic Chemistry, 1999, 27, 100-118.	4.1	42
131	Common Chelatase Design in the Branched Tetrapyrrole Pathways of Heme and Anaerobic Cobalamin Synthesisâ€. Biochemistry, 1999, 38, 10660-10669.	2.5	105
132	Peer and Teacher Assessment of the Oral and Written Tasks of a Group Project. Assessment and Evaluation in Higher Education, 1999, 24, 301-314.	5.6	73
133	X-ray Structure of 5-Aminolevulinic Acid Dehydratase fromEscherichia coliComplexed with the Inhibitor Levulinic Acid at 2.0 à Resolutionâ€. Biochemistry, 1999, 38, 4266-4276.	2.5	96
134	The role of Saccharomyces cerevisiae Met1p and Met8p in sirohaem and cobalamin biosynthesis. Biochemical Journal, 1999, 338, 701.	3.7	21
135	The X-ray structure of a cobalamin biosynthetic enzyme, cobalt-precorrin-4 methyltransferase. Nature Structural Biology, 1998, 5, 585-592.	9.7	60
136	Cobalamin (vitamin B12) biosynthesis . Cloning, expression and crystallisation of the Bacillus megaterium S-adenosyl- L-methionine-dependent cobalt-precorrin-4 transmethylase CbiF. FEBS Journal, 1998, 254, 341-346.	0.2	8
137	Lead poisoning, haem synthesis and 5-aminolaevulinic acid dehydratase. Trends in Biochemical Sciences, 1998, 23, 217-221.	7.5	141
138	GCAP1(Y99C) Mutant Is Constitutively Active in Autosomal Dominant Cone Dystrophy. Molecular Cell, 1998, 2, 129-133.	9.7	150
139	Recombinant Expression, Purification, and Characterization of Three Isoenzymes of Aspartate Aminotransferase fromArabidopsis thaliana. Protein Expression and Purification, 1998, 12, 381-389.	1.3	39
140	Cobalamin (vitamin B12) biosynthesis: identification and characterization of a Bacillus megaterium cobl operon. Biochemical Journal, 1998, 335, 159-166.	3.7	88
141	Cobalamin (vitamin B12) biosynthesis: functional characterization of the Bacillus megaterium cbi genes required to convert uroporphyrinogen III into cobyrinic acid a,c-diamide. Biochemical Journal, 1998, 335, 167-173.	3.7	37
142	Siroheme Biosynthesis in Higher Plants. Journal of Biological Chemistry, 1997, 272, 2744-2752.	3.4	52
143	Having second thoughts: Student perceptions before and after a peer assessment exercise. Studies in Higher Education, 1997, 22, 233-239.	4.5	149
144	Dipyrromethane cofactor assembly of porphobilinogen deaminase: Formation of apoenzyme and preparation of holoenzyme. Methods in Enzymology, 1997, 281, 317-327.	1.0	7

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145	Reconstitution of the Holoenzyme Form of Escherichia coli Porphobilinogen Deaminase from Apoenzyme with Porphobilinogen and Preuroporphyrinogen:  A Study Using Circular Dichroism Spectroscopy. Biochemistry, 1997, 36, 9273-9282.	2.5	20
146	X-ray structure of 5-aminolaevulinate dehydratase, a hybrid aldolase. Nature Structural Biology, 1997, 4, 1025-1031.	9.7	135
147	Chloroplastic aspartate aminotransferase from Arabidopsis thaliana: an examination of the relationship between the structure of the gene and the spatial structure of the protein. Biochemical Journal, 1996, 319, 969-976.	3.7	28
148	Discovery that the assembly of the dipyrromethane cofactor of porphobilinogen deaminase holoenzyme proceeds initially by the reaction of preuroporphyrinogen with the apoenzyme. Biochemical Journal, 1996, 316, 373-376.	3.7	30
149	Evidence for a covalent intermediate in the <i>S</i> -adenosyl- <scp>l</scp> -methionine-dependent transmethylation reaction catalysed by sirohaem synthase. Biochemical Journal, 1996, 313, 415-421.	3.7	12
150	The maddening business of King George III and porphyria. Trends in Biochemical Sciences, 1996, 21, 229-234.	7.5	16
151	The three-dimensional structure ofEscherichia coliporphobilinogen deaminase at 1.76-Ã resolution. , 1996, 25, 48-78.		40
152	Isolation, characterisation and expression of a cDNA clone encoding plastid aspartate aminotransferase from Arabidopsis thaliana. Plant Molecular Biology, 1995, 27, 1227-1233.	3.9	34
153	Evidence for conformational changes in Escherichia coli porphobilinogen deaminase during stepwise pyrrole chain elongation monitored by increased reactivity of cysteine-134 to alkylation by N-ethylmaleimide. Biochemistry, 1995, 34, 11288-11295.	2.5	24
154	The threeâ€dimensional structures of mutants of porphobilinogen deaminase: Toward an understanding of the structural basis of acute intermittent porphyria. Protein Science, 1994, 3, 1644-1650.	7.6	77
155	5-Amino-6-hydroxy 3,4,5,6-tetrahydropyran-2-one (HAT): A stable, cyclic form of glutamate 1-semialdehyde, the natural precursor for tetrapyrroles. Tetrahedron Letters, 1993, 34, 1177-1180.	1.4	17
156	Enzymic synthesis and structure of precorrin-3, a trimethyldipyrrocorphin intermediate in vitamin B12 biosynthesis. Biochemistry, 1992, 31, 603-609.	2.5	55
157	Expression of 9Salmonella typhimuriumenzymes for cobinamide synthesis Identification of the 11-methyl and 20-methyl transferases of corrin biosynthesis. FEBS Letters, 1992, 301, 73-78.	2.8	52
158	Crystallization and preliminary X-ray investigation of Escherichia coli porphobilinogen deaminase. Journal of Molecular Biology, 1992, 224, 269-271.	4.2	21
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