

Robert B Gennis

List of Publications by Year in descending order

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173
papers

7,988
citations

46984

47
h-index

62565

80
g-index

177
all docs

177
docs citations

177
times ranked

4887
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>Escherichia coli</i> amino acid auxotrophic expression host strains for investigating protein structure–function relationships. <i>Journal of Biochemistry</i> , 2021, 169, 387-394.	0.9	6
2	Time-Resolved Electrometric Study of the F ⁺ O Transition in Cytochrome c Oxidase. The Effect of Zn ²⁺ Ions on the Positive Side of the Membrane. <i>Biochemistry (Moscow)</i> , 2021, 86, 105-122.	0.7	7
3	Evolution of the cytochrome <i>bd</i> oxygen reductase superfamily and the function of CydAA TM in Archaea. <i>ISME Journal</i> , 2021, 15, 3534-3548.	4.4	18
4	Identification of a cytochrome bc ₁ -aa ₃ supercomplex in <i>Rhodobacter sphaeroides</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2021, 1862, 148433.	0.5	8
5	Cryo-EM structures of <i>Escherichia coli</i> cytochrome <i>bo</i> ₃ reveal bound phospholipids and ubiquinone-8 in a dynamic substrate binding site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	3.3	25
6	The Monoheme <i>c</i> Subunit of Respiratory Alternative Complex III Is Not Essential for Electron Transfer to Cytochrome <i>aa</i> ₃ in <i>Flavobacterium johnsoniae</i> . <i>Microbiology Spectrum</i> , 2021, 9, e0013521.	1.2	2
7	Specific inhibition of proton pumping by the T315V mutation in the K channel of cytochrome <i>ba</i> from <i>Thermus thermophilus</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2021, 1862, 148450.	0.5	5
8	The three-spin intermediate at the O–O cleavage and proton-pumping junction in heme–Cu oxidases. <i>Science</i> , 2021, 373, 1225-1229.	6.0	13
9	Energy transfer between the nicotinamide nucleotide transhydrogenase and ATP synthase of <i>Escherichia coli</i> . <i>Scientific Reports</i> , 2021, 11, 21234.	1.6	4
10	Structure of the cytochrome <i>aa</i> ₃ -600 heme-copper menaquinol oxidase bound to inhibitor HQNO shows TMO is part of the quinol binding site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 872-876.	3.3	21
11	The oligomeric state of the <i>Caldivirga maquilingsensis</i> type III sulfide:Quinone Oxidoreductase is required for membrane binding. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2020, 1861, 148132.	0.5	2
12	Discovery of Prenyltransferase Inhibitors with <i>In Vitro</i> and <i>In Vivo</i> Antibacterial Activity. <i>ACS Infectious Diseases</i> , 2020, 6, 2979-2993.	1.8	14
13	Role of respiratory <i>NADH</i> oxidation in the regulation of <i>Staphylococcus aureus</i> virulence. <i>EMBO Reports</i> , 2020, 21, e45832.	2.0	16
14	The carboxy-terminal insert in the Q-loop is needed for functionality of <i>Escherichia coli</i> cytochrome <i>bd</i> -I. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2020, 1861, 148175.	0.5	19
15	The Ubiquinol Binding Site of Cytochrome <i>bo</i> ₃ from <i>Escherichia coli</i> Accommodates Menaquinone and Stabilizes a Functional Menasemiquinone. <i>Biochemistry</i> , 2019, 58, 4559-4569.	1.2	6
16	Characterization and X-ray structure of the NADH-dependent coenzyme A disulfide reductase from <i>Thermus thermophilus</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2019, 1860, 148080.	0.5	1
17	Single-particle cryo-EM studies of transmembrane proteins in SMA copolymer nanodiscs. <i>Chemistry and Physics of Lipids</i> , 2019, 221, 114-119.	1.5	34
18	Microcin J25 inhibits ubiquinol oxidase activity of purified cytochrome <i>bd</i> -I from <i>Escherichia coli</i> . <i>Biochimie</i> , 2019, 160, 141-147.	1.3	14

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19	X-ray transparent microfluidic platforms for membrane protein crystallization with microseeds. <i>Lab on A Chip</i> , 2018, 18, 944-954.	3.1	19
20	Structure of the alternative complex III in a supercomplex with cytochrome oxidase. <i>Nature</i> , 2018, 557, 123-126.	13.7	198
21	Unpaired Electron Spin Density Distribution across Reduced [2Fe-2S] Cluster Ligands by ¹³ C- ¹⁵ N-Cysteine Labeling. <i>Inorganic Chemistry</i> , 2018, 57, 741-746.	1.9	7
22	Cytochrome <i>aa</i> ₃ Oxygen Reductase Utilizes the Tunnel Observed in the Crystal Structures To Deliver O ₂ for Catalysis. <i>Biochemistry</i> , 2018, 57, 2150-2161.	1.2	5
23	Role of the tightly bound quinone for the oxygen reaction of cytochrome <i>bo</i> ₃ oxidase from <i>Escherichia coli</i> . <i>FEBS Letters</i> , 2018, 592, 3380-3387.	1.3	8
24	Mechanism of proton transfer through the KC proton pathway in the <i>Vibrio cholerae</i> cbb3 terminal oxidase. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2018, 1859, 1191-1198.	0.5	3
25	The electron distribution in the "activated" state of cytochrome c oxidase. <i>Scientific Reports</i> , 2018, 8, 7502.	1.6	15
26	Functional importance of Glutamate-445 and Glutamate-99 in proton-coupled electron transfer during oxygen reduction by cytochrome <i>bd</i> from <i>Escherichia coli</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2018, 1859, 577-590.	0.5	11
27	Ionophoric effects of the antitubercular drug bedaquiline. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 7326-7331.	3.3	85
28	Type 2 NADH Dehydrogenase Is the Only Point of Entry for Electrons into the <i>Streptococcus agalactiae</i> Respiratory Chain and Is a Potential Drug Target. <i>MBio</i> , 2018, 9, .	1.8	24
29	Bacterial denitrifying nitric oxide reductases and aerobic respiratory terminal oxidases use similar delivery pathways for their molecular substrates. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2018, 1859, 712-724.	0.5	10
30	Dynamics of the K ^B Proton Pathway in Cytochrome <i>ba</i> ₃ from <i>Thermus thermophilus</i> . <i>Israel Journal of Chemistry</i> , 2017, 57, 424-436.	1.0	6
31	Searching for the low affinity ubiquinone binding site in cytochrome <i>bo</i> ₃ from <i>Escherichia coli</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2017, 1858, 366-370.	0.5	8
32	X-ray transparent microfluidic chips for high-throughput screening and optimization of in meso membrane protein crystallization. <i>Biomicrofluidics</i> , 2017, 11, 024118.	1.2	7
33	Location of the Substrate Binding Site of the Cytochrome <i>bo</i> ₃ Ubiquinol Oxidase from <i>Escherichia coli</i> . <i>Journal of the American Chemical Society</i> , 2017, 139, 8346-8354.	6.6	17
34	Critical Role of Water Molecules in Proton Translocation by the Membrane-Bound Transhydrogenase. <i>Structure</i> , 2017, 25, 1111-1119.e3.	1.6	12
35	The ¹³ C Photodissociation and Recombination Dynamics of the W172Y/F282T Ligand Channel Mutant of <i>Rhodobacter sphaeroides</i> <i>aa</i> ₃ Cytochrome <i>c</i> Oxidase. <i>Photochemistry and Photobiology</i> , 2016, 92, 410-419.	1.3	3
36	Proton Dynamics at the Membrane Surface. <i>Biophysical Journal</i> , 2016, 110, 1909-1911.	0.2	17

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37	The unusual redox properties of C-type oxidases. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2016, 1857, 1892-1899.	0.5	27
38	Q-Band Electron-Nuclear Double Resonance Reveals Out-of-Plane Hydrogen Bonds Stabilize an Anionic Ubisemiquinone in Cytochrome bo ₃ from <i>Escherichia coli</i> . <i>Biochemistry</i> , 2016, 55, 5714-5725.	1.2	9
39	CtaM Is Required for Menaquinol Oxidase <i>aa</i> ₃ Function in <i>Staphylococcus aureus</i> . <i>MBio</i> , 2016, 7, .	1.8	34
40	All the O ₂ Consumed by <i>Thermus thermophilus</i> Cytochrome ba ₃ Is Delivered to the Active Site through a Long, Open Hydrophobic Tunnel with Entrances within the Lipid Bilayer. <i>Biochemistry</i> , 2016, 55, 1265-1278.	1.2	17
41	<i>Escherichia coli</i> Auxotroph Host Strains for Amino Acid-Selective Isotope Labeling of Recombinant Proteins. <i>Methods in Enzymology</i> , 2015, 565, 45-66.	0.4	19
42	Antiinfectives targeting enzymes and the proton motive force. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E7073-82.	3.3	138
43	Division of labor in transhydrogenase by alternating proton translocation and hydride transfer. <i>Science</i> , 2015, 347, 178-181.	6.0	36
44	Mutation of a single residue in the <i>ba</i> ₃ oxidase specifically impairs protonation of the pump site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 3397-3402.	3.3	23
45	The two transmembrane helices of CcoP are sufficient for assembly of the cbb ₃ -type heme-copper oxygen reductase from <i>Vibrio cholerae</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2015, 1847, 1231-1239.	0.5	8
46	Review and Hypothesis. New insights into the reaction mechanism of transhydrogenase: Swivelling the dIII component may gate the proton channel. <i>FEBS Letters</i> , 2015, 589, 2027-2033.	1.3	16
47	Plasticity in the High Affinity Menaquinone Binding Site of the Cytochrome <i>aa</i> ₃ -600 Menaquinol Oxidase from <i>Bacillus subtilis</i> . <i>Biochemistry</i> , 2015, 54, 5030-5044.	1.2	9
48	Replacing Arg70 by Histidine in The Cytochrome Aa ₃ 600 Menaquinol Oxidase from <i>Bacillus Subtilis</i> Changes The Nitrogen Interacting with The Semiquinone Formed at The Q _h Site But Does Not Eliminate Catalytic Function. <i>FASEB Journal</i> , 2015, 29, LB110.	0.2	0
49	Conformational coupling between the active site and residues within the KC-channel of the <i>Vibrio cholerae</i> cbb ₃ -type (C-family) oxygen reductase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E4419-E4428.	3.3	8
50	Characterization of the type 2 NADH:menaquinone oxidoreductases from <i>Staphylococcus aureus</i> and the bactericidal action of phenothiazines. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2014, 1837, 954-963.	0.5	41
51	Subunit CydX of <i>Escherichia coli</i> cytochrome <i>bd</i> ubiquinol oxidase is essential for assembly and stability of the di-heme active site. <i>FEBS Letters</i> , 2014, 588, 1537-1541.	1.3	68
52	The K ^C Channel in the <i>cbb</i> ₃ -Type Respiratory Oxygen Reductase from <i>Rhodobacter capsulatus</i> Is Required for Both Chemical and Pumped Protons. <i>Journal of Bacteriology</i> , 2014, 196, 1825-1832.	1.0	2
53	Multitarget Drug Discovery for Tuberculosis and Other Infectious Diseases. <i>Journal of Medicinal Chemistry</i> , 2014, 57, 3126-3139.	2.9	205
54	Kinetics and Intermediates of the Reaction of Fully Reduced <i>Escherichia coli</i> bo ₃ Ubiquinol Oxidase with O ₂ . <i>Biochemistry</i> , 2014, 53, 5393-5404.	1.2	6

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55	Proton pumping by an inactive structural variant of cytochrome c oxidase. <i>Journal of Inorganic Biochemistry</i> , 2014, 140, 6-11.	1.5	11
56	The Semiquinone at the Q _i Site of the bc ₁ Complex Explored Using HYSCORE Spectroscopy and Specific Isotopic Labeling of Ubiquinone in <i>Rhodobacter sphaeroides</i> via ¹³ C Methionine and Construction of a Methionine Auxotroph. <i>Biochemistry</i> , 2014, 53, 6022-6031.	1.2	14
57	Characterization of the Type III sulfide:quinone oxidoreductase from <i>Caldivirga maquilingsis</i> and its membrane binding. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2013, 1827, 266-275.	0.5	34
58	Alternate pathways for NADH oxidation in <i>Thermus thermophilus</i> using type 2 NADH dehydrogenases. <i>Biological Chemistry</i> , 2013, 394, 667-676.	1.2	7
59	Characterization of the nitric oxide reductase from <i>Thermus thermophilus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 12613-12618.	3.3	15
60	Dissection of Hydrogen Bond Interaction Network around an Iron-Sulfur Cluster by Site-Specific Isotope Labeling of Hyperthermophilic Archaeal Rieske-Type Ferredoxin. <i>Journal of the American Chemical Society</i> , 2012, 134, 19731-19738.	6.6	19
61	Interactions of Intermediate Semiquinone with Surrounding Protein Residues at the Q _H Site of Wild-Type and D75H Mutant Cytochrome <i>bc₃</i> from <i>Escherichia coli</i> . <i>Biochemistry</i> , 2012, 51, 3827-3838.	1.2	31
62	Product-controlled steady-state kinetics between cytochrome aa ₃ from <i>Rhodobacter sphaeroides</i> and equine ferrocytochrome c analyzed by a novel spectrophotometric approach. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 1894-1900.	0.5	1
63	Functional Importance of a Pair of Conserved Glutamic Acid Residues and of Ca ²⁺ Binding in the cbb ₃ -Type Oxygen Reductases from <i>Rhodobacter sphaeroides</i> and <i>Vibrio cholerae</i> . <i>Biochemistry</i> , 2012, 51, 7290-7296.	1.2	8
64	Structure Changes upon Deprotonation of the Proton Release Group in the Bacteriorhodopsin Photocycle. <i>Biophysical Journal</i> , 2012, 103, 444-452.	0.2	16
65	Timing of Electron and Proton Transfer in the <i>ba₃</i> Cytochrome <i>c</i> Oxidase from <i>Thermus thermophilus</i> . <i>Biochemistry</i> , 2012, 51, 4507-4517.	1.2	15
66	Cell-free synthesis of cytochrome bo ₃ ubiquinol oxidase in artificial membranes. <i>Analytical Biochemistry</i> , 2012, 423, 39-45.	1.1	20
67	Proton transfer in <i>ba₃</i> cytochrome c oxidase from <i>Thermus thermophilus</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 650-657.	0.5	52
68	Solid-State NMR Study of the Charge-Transfer Complex between Ubiquinone-8 and Disulfide Bond Generating Membrane Protein DsbB. <i>Journal of the American Chemical Society</i> , 2011, 133, 4359-4366.	6.6	20
69	A rapid and robust method for selective isotope labeling of proteins. <i>Methods</i> , 2011, 55, 370-378.	1.9	55
70	211436 Optimization of expression condition and purification of cbb ₃ -type cytochrome c oxidase(Heme) Tj ETQq0,0,0 rgBT /Overlock 1		
71	Differential effects of glutamate-286 mutations in the aa ₃ -type cytochrome c oxidase from <i>Rhodobacter sphaeroides</i> and the cytochrome bo ₃ ubiquinol oxidase from <i>Escherichia coli</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2011, 1807, 1342-1348.	0.5	2
72	The cytochrome bd respiratory oxygen reductases. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2011, 1807, 1398-1413.	0.5	445

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73	Oriented immobilization and electron transfer to the cytochrome c oxidase. <i>Journal of Solid State Electrochemistry</i> , 2011, 15, 105-114.	1.2	31
74	Aerobic respiratory chain of <i>Escherichia coli</i> is not allowed to work in fully uncoupled mode. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 17320-17324.	3.3	121
75	Kinetic design of the respiratory oxidases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 11057-11062.	3.3	36
76	Entrance of the proton pathway in <i>cbb₃</i> -type heme-copper oxidases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 17661-17666.	3.3	35
77	Exploring by Pulsed EPR the Electronic Structure of Ubisemiquinone Bound at the QH Site of Cytochrome bo ₃ from <i>Escherichia coli</i> with in Vivo ¹³ C-Labeled Methyl and Methoxy Substituents. <i>Journal of Biological Chemistry</i> , 2011, 286, 10105-10114.	1.6	20
78	Adaptation of aerobic respiration to low O ₂ environments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 14109-14114.	3.3	119
79	Partial Steps of Charge Translocation in the Nonpumping N139L Mutant of <i>Rhodobacter sphaeroides</i> Cytochrome c Oxidase with a Blocked D-Channel. <i>Biochemistry</i> , 2010, 49, 3060-3073.	1.2	30
80	Nitroxide spin labels as EPR reporters of the relaxation and magnetic properties of the heme-copper site in cytochrome bo ₃ , <i>E. coli</i> . <i>Journal of Biological Inorganic Chemistry</i> , 2010, 15, 1255-1264.	1.1	6
81	Blocking the K-pathway still allows rapid one-electron reduction of the binuclear center during the anaerobic reduction of the aa ₃ -type cytochrome c oxidase from <i>Rhodobacter sphaeroides</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2010, 1797, 619-624.	0.5	15
82	The quinone-binding sites of the cytochrome bo ₃ ubiquinol oxidase from <i>Escherichia coli</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2010, 1797, 1924-1932.	0.5	41
83	Heme-heme and heme-ligand interactions in the di-heme oxygen-reducing site of cytochrome bd from <i>Escherichia coli</i> revealed by nanosecond absorption spectroscopy. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2010, 1797, 1657-1664.	0.5	36
84	Functional interactions between membrane-bound transporters and membranes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 15763-15767.	3.3	27
85	Characterization of the Semiquinone Radical Stabilized by the Cytochrome aa ₃ -600 Menaquinol Oxidase of <i>Bacillus subtilis</i> . <i>Journal of Biological Chemistry</i> , 2010, 285, 18241-18251.	1.6	24
86	Decoupling Mutations in the D-Channel of the aa ₃ -Type Cytochrome c Oxidase from <i>Rhodobacter sphaeroides</i> Suggest That a Continuous Hydrogen-Bonded Chain of Waters Is Essential for Proton Pumping. <i>Biochemistry</i> , 2010, 49, 4476-4482.	1.2	28
87	The Diheme Cytochrome c ₄ from <i>Vibrio cholerae</i> Is a Natural Electron Donor to the Respiratory cbb ₃ Oxygen Reductase. <i>Biochemistry</i> , 2010, 49, 7494-7503.	1.2	34
88	Conformational transitions and molecular hysteresis of cytochrome c oxidase: Varying the redox state by electronic wiring. <i>Soft Matter</i> , 2010, 6, 5523.	1.2	21
89	The cytochrome ba ₃ oxygen reductase from <i>Thermus thermophilus</i> uses a single input channel for proton delivery to the active site and for proton pumping. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 16169-16173.	3.3	102
90	Critical structural role of R481 in cytochrome c oxidase from <i>Rhodobacter sphaeroides</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2009, 1787, 1272-1275.	0.5	14

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91	Properties of Arg481 Mutants of the <i>aa₃</i> -Type Cytochrome <i>c</i> Oxidase from <i>Rhodobacter sphaeroides</i> Suggest That neither R481 nor the Nearby D-Propionate of Heme <i>a₃</i> Is Likely To Be the Proton Loading Site of the Proton Pump. <i>Biochemistry</i> , 2009, 48, 7123-7131.	1.2	33
92	Cytochrome <i>c</i> oxidase: exciting progress and remaining mysteries. <i>Journal of Bioenergetics and Biomembranes</i> , 2008, 40, 521-531.	1.0	252
93	Diversity of the Heme-Copper Superfamily in Archaea: Insights from Genomics and Structural Modeling. , 2008, 45, 1-31.		124
94	The fully oxidized form of the cytochrome <i>bd</i> quinol oxidase from <i>E. coli</i> does not participate in the catalytic cycle: Direct evidence from rapid kinetics studies. <i>FEBS Letters</i> , 2008, 582, 3705-3709.	1.3	33
95	Impaired proton pumping in cytochrome <i>c</i> oxidase upon structural alteration of the D pathway. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2008, 1777, 897-903.	0.5	43
96	Strong Excitonic Interactions in the Oxygen-Reducing Site of <i>bd</i> -Type Oxidase: The Fe-to-Fe Distance between Hemes <i>d</i> and <i>b₅₉₅</i> is 10 Å.... <i>Biochemistry</i> , 2008, 47, 1752-1759.	1.2	41
97	Identification of the Nitrogen Donor Hydrogen Bonded with the Semiquinone at the Q _H Site of the Cytochrome <i>bo₃</i> from <i>Escherichia coli</i> . <i>Journal of the American Chemical Society</i> , 2008, 130, 15768-15769.	6.6	28
98	Characterization of Mutants That Change the Hydrogen Bonding of the Semiquinone Radical at the QH Site of the Cytochrome <i>bo₃</i> from <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2007, 282, 8777-8785.	1.6	29
99	Comparative Genomics and Site-Directed Mutagenesis Support the Existence of Only One Input Channel for Protons in the C-Family (<i>cbb₃</i> Oxidase) of Heme-Copper Oxygen Reductases. <i>Biochemistry</i> , 2007, 46, 9963-9972.	1.2	70
100	Flash-Photolysis of Fully Reduced and Mixed-Valence CO-Bound <i>Rhodobacter sphaeroides</i> Cytochrome <i>c</i> Oxidase: Heme Spectral Shifts. <i>Biochemistry</i> , 2007, 46, 12568-12578.	1.2	11
101	Glutamate 107 in Subunit I of the Cytochrome <i>bd</i> Quinol Oxidase from <i>Escherichia coli</i> Protonated and near the Heme <i>d</i> /Heme <i>b₅₉₅</i> Binuclear Center. <i>Biochemistry</i> , 2007, 46, 3270-3278.	1.2	31
102	A New Ruthenium Complex To Study Single-Electron Reduction of the Pulsed OH State of Detergent-Solubilized Cytochrome Oxidase. <i>Biochemistry</i> , 2007, 46, 14610-14618.	1.2	26
103	Controlled uncoupling and recoupling of proton pumping in cytochrome <i>c</i> oxidase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 317-322.	3.3	89
104	Single-electron photoreduction of the PM intermediate of cytochrome <i>c</i> oxidase. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2006, 1757, 1122-1132.	0.5	38
105	Replacing Asn207 by Aspartate at the Neck of the D Channel in the <i>aa₃</i> -Type Cytochrome <i>c</i> Oxidase from <i>Rhodobacter sphaeroides</i> Results in Decoupling the Proton Pump. <i>Biochemistry</i> , 2006, 45, 14064-14074.	1.2	44
106	Water as a Cofactor in the Unidirectional Light-Driven Proton Transfer Steps in Bacteriorhodopsin. <i>Photochemistry and Photobiology</i> , 2006, 82, 1398-1405.	1.3	9
107	Magic-angle spinning solid-state NMR of a 144 kDa membrane protein complex: <i>E. coli</i> cytochrome <i>bo₃</i> oxidase. <i>Journal of Biomolecular NMR</i> , 2006, 36, 55-71.	1.6	75
108	Characterization of the Exchangeable Protons in the Immediate Vicinity of the Semiquinone Radical at the QH Site of the Cytochrome <i>bo₃</i> from <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2006, 281, 16879-16887.	1.6	39

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109	Spectral and Kinetic Equivalence of Oxidized Cytochrome c Oxidase as Isolated and "Activated" by Reoxidation. <i>Journal of Biological Chemistry</i> , 2006, 281, 30319-30325.	1.6	45
110	Respiration in Archaea and Bacteria: Diversity of Prokaryotic Electron Transport Carriers. Davide Zannoni, <i>Advances in Photosynthesis and Respiration (Series Editor, Govindjee)</i> , Kluwer Academic Publishers, Dordrecht, The Netherlands, Volume 15, 2004, 350 pp, ISBN 1-4020-2001-5, Price EUR 175.00, USD 193.00, GBP 121.00.. <i>Photosynthesis Research</i> , 2005, 83, 363-364.	1.6	0
111	Time-resolved electrometric and optical studies on cytochrome bd suggest a mechanism of electron-proton coupling in the di-heme active site. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 3657-3662.	3.3	76
112	Helix Switching of a Key Active-Site Residue in the Cytochromecbb3Oxidases. <i>Biochemistry</i> , 2005, 44, 10766-10775.	1.2	56
113	Transmembrane Charge Separation during the Ferryl-oxo "Oxidized Transition in a Nonpumping Mutant of Cytochrome c Oxidase. <i>Journal of Biological Chemistry</i> , 2004, 279, 52558-52565.	1.6	75
114	Some recent contributions of FTIR difference spectroscopy to the study of cytochrome oxidase1. <i>FEBS Letters</i> , 2003, 555, 2-7.	1.3	26
115	The Entry Point of the K-Proton-Transfer Pathway in CytochromecOxidase. <i>Biochemistry</i> , 2002, 41, 10794-10798.	1.2	68
116	A Mutation in Subunit I of Cytochrome Oxidase from <i>Rhodobacter sphaeroides</i> Results in an Increase in Steady-State Activity but Completely Eliminates Proton Pumping. <i>Biochemistry</i> , 2002, 41, 13417-13423.	1.2	122
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