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List of Publications by Year in descending order

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47
papers

7,170
citations

201674

27
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206112

48
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all docs

55
docs citations

55
times ranked

6997
citing authors

#	ARTICLE	IF	CITATIONS
1	Atypical Molecular Basis for Drug Resistance to Mitochondrial Function Inhibitors in Plasmodium falciparum. Antimicrobial Agents and Chemotherapy, 2021, 65, .	3.2	7
2	Divergent Mitochondrial Ribosomes in Unicellular Parasitic Protozoans. Trends in Parasitology, 2020, 36, 318-321.	3.3	6
3	Genetic ablation of the mitoribosome in the malaria parasite Plasmodium falciparum sensitizes it to antimalarials that target mitochondrial functions. Journal of Biological Chemistry, 2020, 295, 7235-7248.	3.4	23
4	para-Aminobenzoate Synthesis versus Salvage in Malaria Parasites. Trends in Parasitology, 2019, 35, 176-178.	3.3	6
5	Mitochondrial type II NADH dehydrogenase of Plasmodium falciparum (PfNDH2) is dispensable in the asexual blood stages. PLoS ONE, 2019, 14, e0214023.	2.5	29
6	The mitochondrial ribosomal protein L13 is critical for the structural and functional integrity of the mitochondrion in Plasmodium falciparum. Journal of Biological Chemistry, 2018, 293, 8128-8137.	3.4	50
7	Novel Defense Peptides from Platelets Kill Malaria Parasites. Trends in Parasitology, 2018, 34, 729-731.	3.3	6
8	Alkoxy carbonate Ester Prodrugs of Preclinical Drug Candidate ELQ-300 for Prophylaxis and Treatment of Malaria. ACS Infectious Diseases, 2017, 3, 728-735.	3.8	38
9	Functional Profiling of a Plasmodium Genome Reveals an Abundance of Essential Genes. Cell, 2017, 170, 260-272.e8.	28.9	471
10	Caged Garcinia Xanthenes, a Novel Chemical Scaffold with Potent Antimalarial Activity. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	15
11	+Targeting Mitochondrial Functions as Antimalarial Regime, What Is Next?. Current Clinical Microbiology Reports, 2017, 4, 175-191.	3.4	12
12	Characterization of a Plasmodium falciparum Orthologue of the Yeast Ubiquinone-Binding Protein, Coq10p. PLoS ONE, 2016, 11, e0152197.	2.5	6
13	ELQ-300 Prodrugs for Enhanced Delivery and Single-Dose Cure of Malaria. Antimicrobial Agents and Chemotherapy, 2015, 59, 5555-5560.	3.2	62
14	Inhibition of Cytochrome bc 1 as a Strategy for Single-Dose, Multi-Stage Antimalarial Therapy. American Journal of Tropical Medicine and Hygiene, 2015, 92, 1195-1201.	1.4	34
15	Genetic Investigation of Tricarboxylic Acid Metabolism during the Plasmodium falciparum Life Cycle. Cell Reports, 2015, 11, 164-174.	6.4	134
16	The Heme Biosynthesis Pathway Is Essential for Plasmodium falciparum Development in Mosquito Stage but Not in Blood Stages. Journal of Biological Chemistry, 2014, 289, 34827-34837.	3.4	133
17	Discovery, Synthesis, and Optimization of Antimalarial 4(1 <i>H</i>)-Quinolone-3-Diarylethers. Journal of Medicinal Chemistry, 2014, 57, 3818-3834.	6.4	100
18	Quinolone-3-Diarylethers: A New Class of Antimalarial Drug. Science Translational Medicine, 2013, 5, 177ra37.	12.4	187

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19	Mitochondrial RNA polymerase is an essential enzyme in erythrocytic stages of <i>Plasmodium falciparum</i> . <i>Molecular and Biochemical Parasitology</i> , 2012, 185, 48-51.	1.1	10
20	Variation among <i>Plasmodium falciparum</i> Strains in Their Reliance on Mitochondrial Electron Transport Chain Function. <i>Eukaryotic Cell</i> , 2011, 10, 1053-1061.	3.4	59
21	Comparative genomics of the pathogenic ciliate <i>Ichthyophthirius multifiliis</i> , its free-living relatives and a host species provide insights into adoption of a parasitic lifestyle and prospects for disease control. <i>Genome Biology</i> , 2011, 12, R100.	9.6	102
22	Yeast dihydroorotate dehydrogenase as a new selectable marker for <i>Plasmodium falciparum</i> transfection. <i>Molecular and Biochemical Parasitology</i> , 2011, 177, 29-34.	1.1	94
23	ATP Synthase Complex of <i>Plasmodium falciparum</i> . <i>Journal of Biological Chemistry</i> , 2011, 286, 41312-41322.	3.4	69
24	Hemozoin-free <i>Plasmodium falciparum</i> mitochondria for physiological and drug susceptibility studies. <i>Molecular and Biochemical Parasitology</i> , 2010, 174, 150-153.	1.1	27
25	Branched tricarboxylic acid metabolism in <i>Plasmodium falciparum</i> . <i>Nature</i> , 2010, 466, 774-778.	27.8	111
26	Highly Divergent Mitochondrial ATP Synthase Complexes in <i>Tetrahymena thermophila</i> . <i>PLoS Biology</i> , 2010, 8, e1000418.	5.6	72
27	Mitochondrial Evolution and Functions in Malaria Parasites. <i>Annual Review of Microbiology</i> , 2009, 63, 249-267.	7.3	207
28	Mitochondria in malaria and related parasites: ancient, diverse and streamlined. <i>Journal of Bioenergetics and Biomembranes</i> , 2008, 40, 425-33.	2.3	47
29	The validity of mitochondrial dehydrogenases as antimalarial drug targets. <i>Trends in Parasitology</i> , 2008, 24, 8-9.	3.3	25
30	Mitochondrial Drug Targets in Apicomplexan Parasites. <i>Current Drug Targets</i> , 2007, 8, 49-60.	2.1	100
31	Specific role of mitochondrial electron transport in blood-stage <i>Plasmodium falciparum</i> . <i>Nature</i> , 2007, 446, 88-91.	27.8	441
32	Uncovering the Molecular Mode of Action of the Antimalarial Drug Atovaquone Using a Bacterial System. <i>Journal of Biological Chemistry</i> , 2005, 280, 27458-27465.	3.4	83
33	The inhibition of calcium signaling in T lymphocytes from old mice results from enhanced activation of the mitochondrial permeability transition pore. <i>Mechanisms of Ageing and Development</i> , 2002, 123, 707-724.	4.6	34
34	Genome sequence of the human malaria parasite <i>Plasmodium falciparum</i> . <i>Nature</i> , 2002, 419, 498-511.	27.8	3,881
35	Atovaquone resistance in malaria parasites. <i>Drug Resistance Updates</i> , 2000, 3, 283-287.	14.4	69
36	Intrinsic uncoupling of cytochromecoxidase may cause the maternally inherited mitochondrial diseases MELAS and LHON. <i>FEBS Letters</i> , 1998, 433, 93-97.	2.8	20

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37	Flexibility of the Neck Region of the Rieske Iron-Sulfur Protein Is Functionally Important in the Cytochrome bc ₁ Complex. <i>Journal of Biological Chemistry</i> , 1998, 273, 27953-27959.	3.4	94
38	The Involvement of Serine 175 and Alanine 185 of Cytochrome b of <i>Rhodobacter sphaeroides</i> Cytochrome bc ₁ Complex in Interaction with Iron-Sulfur Protein. <i>Journal of Biological Chemistry</i> , 1997, 272, 23722-23728.	3.4	19
39	The Involvement of Threonine 160 of Cytochrome b of <i>Rhodobacter sphaeroides</i> Cytochrome bc ₁ Complex in Quinone Binding and Interaction with Subunit IV. <i>Journal of Biological Chemistry</i> , 1995, 270, 28668-28675.	3.4	28
40	Molecular Genetic and Protein Chemical Characterization of the Cytochrome ba ₃ from <i>Thermus thermophilus</i> HB8. <i>Journal of Biological Chemistry</i> , 1995, 270, 20345-20358.	3.4	74
41	An enhanced broad-host-range vector for Gram-negative bacteria: Avoiding tetracycline phototoxicity during the growth of photosynthetic bacteria. <i>Gene</i> , 1995, 156, 85-88.	2.2	25
42	Molecular modeling studies on the proposed NaCl-induced dimerization of <i>Chromatium vinosum</i> high-potential iron protein. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 1993, 1142, 93-98.	1.0	6
43	Cytochrome caa ₃ from the thermophilic bacterium <i>Thermus thermophilus</i> : A member of the heme-copper oxidase superfamily. <i>Journal of Bioenergetics and Biomembranes</i> , 1993, 25, 103-114.	2.3	26
44	[50] Recovery and cloning of genomic DNA fragments from dried agarose gels. <i>Methods in Enzymology</i> , 1993, 218, 695-704.	1.0	7
45	Plasmid-associated aggregation in <i>Thermus thermophilus</i> HB8. <i>Plasmid</i> , 1990, 24, 45-56.	1.4	16
46	Respiratory proteins from extremely thermophilic, aerobic bacteria. <i>Biochimica Et Biophysica Acta - Reviews on Bioenergetics</i> , 1986, 853, 153-185.	0.2	85
47	Activation of Pyruvate Oxidase and Interaction with Membrane Components. , 1982, , 83-92.		2