Amanda K Lukens

List of Publications by Year in descending order

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AMANDA K LUKENS

#	Article	IF	CITATIONS
1	The Plasmodium falciparum ABC transporter ABCI3 confers parasite strain-dependent pleiotropic antimalarial drug resistance. Cell Chemical Biology, 2022, 29, 824-839.e6.	2.5	14
2	Chemogenomics identifies acetyl-coenzyme A synthetase as a target for malaria treatment and prevention. Cell Chemical Biology, 2022, 29, 191-201.e8.	2.5	39
3	Adaptive laboratory evolution in S. cerevisiae highlights role of transcription factors in fungal xenobiotic resistance. Communications Biology, 2022, 5, 128.	2.0	8
4	MalDA, Accelerating Malaria Drug Discovery. Trends in Parasitology, 2021, 37, 493-507.	1.5	51
5	Prioritization of Molecular Targets for Antimalarial Drug Discovery. ACS Infectious Diseases, 2021, 7, 2764-2776.	1.8	35
6	The Adaptive Proline Response in <i>P. falciparum</i> Is Independent of <i>Pf</i> elK1 and elF2α Signaling. ACS Infectious Diseases, 2019, 5, 515-520.	1.8	5
7	In vitro selection predicts malaria parasite resistance to dihydroorotate dehydrogenase inhibitors in a mouse infection model. Science Translational Medicine, 2019, 11, .	5.8	30
8	Identification of Collateral Sensitivity to Dihydroorotate Dehydrogenase Inhibitors in <i>Plasmodium falciparum</i> . ACS Infectious Diseases, 2018, 4, 508-515.	1.8	15
9	Mapping the malaria parasite druggable genome by using in vitro evolution and chemogenomics. Science, 2018, 359, 191-199.	6.0	194
10	Quantitative Proteomic Profiling Reveals Novel Plasmodium falciparum Surface Antigens and Possible Vaccine Candidates. Molecular and Cellular Proteomics, 2018, 17, 43-60.	2.5	29
11	Open-source discovery of chemical leads for next-generation chemoprotective antimalarials. Science, 2018, 362, .	6.0	99
12	Intramolecular Diazaâ€Diels–Alder Protocol: A New Diastereoselective and Modular Oneâ€6tep Synthesis of Constrained Polycyclic Frameworks. Chemistry - A European Journal, 2017, 23, 4137-4148.	1.7	15
13	Genome-Wide Association Studies of Drug-Resistance Determinants. Trends in Parasitology, 2017, 33, 214-230.	1.5	16
14	New paradigms for understanding and step changes in treating active and chronic, persistent apicomplexan infections. Scientific Reports, 2016, 6, 29179.	1.6	40
15	<i>Plasmodium falciparum</i> Cyclic Amine Resistance Locus (PfCARL), a Resistance Mechanism for Two Distinct Compound Classes. ACS Infectious Diseases, 2016, 2, 816-826.	1.8	34
16	Diversity-oriented synthesis yields novel multistage antimalarial inhibitors. Nature, 2016, 538, 344-349.	13.7	214
17	Probing the Azaaurone Scaffold against the Hepatic and Erythrocytic Stages of Malaria Parasites. ChemMedChem, 2016, 11, 2194-2204.	1.6	23
18	A broad analysis of resistance development in the malaria parasite. Nature Communications, 2016, 7, 11901.	5.8	94

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19	Diversity-Oriented Synthesis Probe TargetsPlasmodium falciparumCytochrome b Ubiquinone Reduction Site and Synergizes With Oxidation Site Inhibitors. Journal of Infectious Diseases, 2015, 211, 1097-1103.	1.9	29
20	The cytoplasmic prolyl-tRNA synthetase of the malaria parasite is a dual-stage target of febrifugine and its analogs. Science Translational Medicine, 2015, 7, 288ra77.	5.8	82
21	Exploring the 3-piperidin-4-yl-1H-indole scaffold as a novel antimalarial chemotype. European Journal of Medicinal Chemistry, 2015, 102, 320-333.	2.6	31
22	Triaminopyrimidine is a fast-killing and long-acting antimalarial clinical candidate. Nature Communications, 2015, 6, 6715.	5.8	55
23	Clinical Sequencing Uncovers Origins and Evolution of Lassa Virus. Cell, 2015, 162, 738-750.	13.5	230
24	Harnessing evolutionary fitness in <i>Plasmodium falciparum</i> for drug discovery and suppressing resistance. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 799-804.	3.3	54
25	Diversity-Oriented Synthesis-Facilitated Medicinal Chemistry: Toward the Development of Novel Antimalarial Agents. Journal of Medicinal Chemistry, 2014, 57, 8496-8502.	2.9	33
26	Aminoazabenzimidazoles, a Novel Class of Orally Active Antimalarial Agents. Journal of Medicinal Chemistry, 2014, 57, 5702-5713.	2.9	24
27	Polymorphism in dhfr/dhps genes, parasite density and ex vivo response to pyrimethamine in Plasmodium falciparum malaria parasites in Thies, Senegal. International Journal for Parasitology: Drugs and Drug Resistance, 2013, 3, 135-142.	1.4	27
28	Genetic Surveillance Detects Both Clonal and Epidemic Transmission of Malaria following Enhanced Intervention in Senegal. PLoS ONE, 2013, 8, e60780.	1.1	87
29	SNP Genotyping Identifies New Signatures of Selection in a Deep Sample of West African Plasmodium falciparum Malaria Parasites. Molecular Biology and Evolution, 2012, 29, 3249-3253.	3.5	41
30	Sequence-based association and selection scans identify drug resistance loci in the <i>Plasmodium falciparum</i> malaria parasite. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 13052-13057.	3.3	99
31	Diversity-Oriented Synthesis Yields a Novel Lead for the Treatment of Malaria. ACS Medicinal Chemistry Letters, 2012, 3, 112-117.	1.3	52
32	Identification and Functional Validation of the Novel Antimalarial Resistance Locus PF10_0355 in Plasmodium falciparum. PLoS Genetics, 2011, 7, e1001383.	1.5	85
33	Genome-wide SNP genotyping highlights the role of natural selection in Plasmodium falciparumpopulation divergence. Genome Biology, 2008, 9, R171.	3.8	119
34	A genome-wide map of diversity in Plasmodium falciparum. Nature Genetics, 2007, 39, 113-119.	9.4	320
35	In Vivo Transcriptome ofPlasmodium falciparumReveals Overexpression of Transcripts That Encode Surface Proteins. Journal of Infectious Diseases, 2005, 191, 1196-1203.	1.9	92
36	Intrinsic susceptibility of mouse trophoblasts to natural killer cell-mediated attack in vivo. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 16940-16945.	3.3	29