

Marcus C S Lee

List of Publications by Year in descending order

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Version: 2024-02-01

53
papers

6,783
citations

117625

34
h-index

168389

53
g-index

59
all docs

59
docs citations

59
times ranked

7174
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Spiroindolones, a Potent Compound Class for the Treatment of Malaria. <i>Science</i> , 2010, 329, 1175-1180. | 12.6 | 1,031 |
| 2 | BI-DIRECTIONAL PROTEIN TRANSPORT BETWEEN THE ER AND GOLGI. <i>Annual Review of Cell and Developmental Biology</i> , 2004, 20, 87-123. | 9.4 | 815 |
| 3 | Multiple Cargo Binding Sites on the COPII Subunit Sec24p Ensure Capture of Diverse Membrane Proteins into Transport Vesicles. <i>Cell</i> , 2003, 114, 497-509. | 28.9 | 461 |
| 4 | Sar1p N-Terminal Helix Initiates Membrane Curvature and Completes the Fission of a COPII Vesicle. <i>Cell</i> , 2005, 122, 605-617. | 28.9 | 455 |
| 5 | Targeting Plasmodium PI(4)K to eliminate malaria. <i>Nature</i> , 2013, 504, 248-253. | 27.8 | 377 |
| 6 | A novel multiple-stage antimalarial agent that inhibits protein synthesis. <i>Nature</i> , 2015, 522, 315-320. | 27.8 | 353 |
| 7 | Imaging of <i>Plasmodium</i> Liver Stages to Drive Next-Generation Antimalarial Drug Discovery. <i>Science</i> , 2011, 334, 1372-1377. | 12.6 | 308 |
| 8 | Quantitative assessment of <i>Plasmodium falciparum</i> sexual development reveals potent transmission-blocking activity by methylene blue. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, E1214-23. | 7.1 | 293 |
| 9 | Antimalarial efficacy of MMV390048, an inhibitor of <i>Plasmodium</i> phosphatidylinositol 4-kinase. <i>Science Translational Medicine</i> , 2017, 9, . | 12.4 | 204 |
| 10 | (+)-SJ733, a clinical candidate for malaria that acts through ATP4 to induce rapid host-mediated clearance of <i>Plasmodium</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E5455-62. | 7.1 | 199 |
| 11 | Mapping the malaria parasite druggable genome by using in vitro evolution and chemogenomics. <i>Science</i> , 2018, 359, 191-199. | 12.6 | 194 |
| 12 | Site-specific genome editing in <i>Plasmodium falciparum</i> using engineered zinc-finger nucleases. <i>Nature Methods</i> , 2012, 9, 993-998. | 19.0 | 149 |
| 13 | Profiling the Essential Nature of Lipid Metabolism in Asexual Blood and Gametocyte Stages of <i>Plasmodium falciparum</i> . <i>Cell Host and Microbe</i> , 2015, 18, 371-381. | 11.0 | 144 |
| 14 | Ceramide Biosynthesis Is Required for the Formation of the Oligomeric H ⁺ -ATPase Pma1p in the Yeast Endoplasmic Reticulum. <i>Journal of Biological Chemistry</i> , 2002, 277, 22395-22401. | 3.4 | 124 |
| 15 | KAF156 Is an Antimalarial Clinical Candidate with Potential for Use in Prophylaxis, Treatment, and Prevention of Disease Transmission. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 5060-5067. | 3.2 | 122 |
| 16 | A potent antimalarial benzoxaborole targets a <i>Plasmodium falciparum</i> cleavage and polyadenylation specificity factor homologue. <i>Nature Communications</i> , 2017, 8, 14574. | 12.8 | 110 |
| 17 | UDP-galactose and acetyl-CoA transporters as <i>Plasmodium</i> multidrug resistance genes. <i>Nature Microbiology</i> , 2016, 1, 16166. | 13.3 | 102 |
| 18 | A broad analysis of resistance development in the malaria parasite. <i>Nature Communications</i> , 2016, 7, 11901. | 12.8 | 94 |

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|----|---|-----|-----------|
| 19 | Molecular mechanisms of COPII vesicle formation. <i>Seminars in Cell and Developmental Biology</i> , 2007, 18, 424-434. | 5.0 | 79 |
| 20 | Genomewide Analysis Reveals Novel Pathways Affecting Endoplasmic Reticulum Homeostasis, Protein Modification and Quality Control. <i>Genetics</i> , 2009, 182, 757-769. | 2.9 | 62 |
| 21 | Insights into the intracellular localization, protein associations and artemisinin resistance properties of <i>Plasmodium falciparum</i> K13. <i>PLoS Pathogens</i> , 2020, 16, e1008482. | 4.7 | 60 |
| 22 | Wherever I may roam: Protein and membrane trafficking in <i>P. falciparum</i> -infected red blood cells. <i>Molecular and Biochemical Parasitology</i> , 2012, 186, 95-116. | 1.1 | 56 |
| 23 | CRISPR-Cas9 modified <i>pfmdr1</i> protects <i>Plasmodium falciparum</i> asexual blood stages and gametocytes against a class of piperazine-containing compounds but potentiates artemisinin-based combination therapy partner drugs. <i>Molecular Microbiology</i> , 2016, 101, 381-393. | 2.5 | 56 |
| 24 | Inhibition of Resistance-Refractory <i>P. falciparum</i> Kinase PKG Delivers Prophylactic, Blood Stage, and Transmission-Blocking Antiplasmodial Activity. <i>Cell Chemical Biology</i> , 2020, 27, 806-816.e8. | 5.2 | 56 |
| 25 | Identification and Characterization of a Prevacuolar Compartment in Stigmas of <i>Nicotiana glauca</i> . <i>Plant Cell</i> , 1999, 11, 1499-1508. | 6.6 | 54 |
| 26 | Combining Stage Specificity and Metabolomic Profiling to Advance Antimalarial Drug Discovery. <i>Cell Chemical Biology</i> , 2020, 27, 158-171.e3. | 5.2 | 54 |
| 27 | A novel two-chain proteinase inhibitor generated by circularization of a multidomain precursor protein. <i>Nature Structural Biology</i> , 1999, 6, 526-530. | 9.7 | 51 |
| 28 | Inhibiting Endoplasmic Reticulum (ER)-associated Degradation of Misfolded Yor1p Does Not Permit ER Export Despite the Presence of a Diacidic Sorting Signal. <i>Molecular Biology of the Cell</i> , 2007, 18, 3398-3413. | 2.1 | 51 |
| 29 | MalDA, Accelerating Malaria Drug Discovery. <i>Trends in Parasitology</i> , 2021, 37, 493-507. | 3.3 | 51 |
| 30 | A Method for Rapid Genetic Integration into <i>Plasmodium falciparum</i> Utilizing Mycobacteriophage Bxb1 Integrase. <i>Methods in Molecular Biology</i> , 2010, 634, 87-100. | 0.9 | 50 |
| 31 | <i>Plasmodium falciparum</i> Sec24 marks transitional ER that exports a model cargo via a diacidic motif. <i>Molecular Microbiology</i> , 2008, 68, 1535-1546. | 2.5 | 49 |
| 32 | The Antimalarial Natural Product Salinipostin A Identifies Essential $\hat{1}\pm/\hat{1}^2$ Serine Hydrolases Involved in Lipid Metabolism in <i>P. falciparum</i> Parasites. <i>Cell Chemical Biology</i> , 2020, 27, 143-157.e5. | 5.2 | 48 |
| 33 | Identification and Mechanistic Understanding of Dihydroorotate Dehydrogenase Point Mutations in <i>Plasmodium falciparum</i> that Confer <i>In Vitro</i> Resistance to the Clinical Candidate DSM265. <i>ACS Infectious Diseases</i> , 2019, 5, 90-101. | 3.8 | 43 |
| 34 | Chemogenomics identifies acetyl-coenzyme A synthetase as a target for malaria treatment and prevention. <i>Cell Chemical Biology</i> , 2022, 29, 191-201.e8. | 5.2 | 39 |
| 35 | Cutting back malaria: CRISPR/Cas9 genome editing of <i>Plasmodium</i> . <i>Briefings in Functional Genomics</i> , 2019, 18, 281-289. | 2.7 | 38 |
| 36 | <i>N</i> -Aryl-2-aminobenzimidazoles: Novel, Efficacious, Antimalarial Lead Compounds. <i>Journal of Medicinal Chemistry</i> , 2014, 57, 6642-6652. | 6.4 | 37 |

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|----|---|------|-----------|
| 37 | Prioritization of Molecular Targets for Antimalarial Drug Discovery. <i>ACS Infectious Diseases</i> , 2021, 7, 2764-2776. | 3.8 | 35 |
| 38 | Identification of a novel four-domain member of the proteinase inhibitor II family from the stigmas of <i>Nicotiana glauca</i> . <i>Plant Molecular Biology</i> , 2000, 42, 329-333. | 3.9 | 33 |
| 39 | CELL BIOLOGY: BAR Domains Go on a Bender. <i>Science</i> , 2004, 303, 479-480. | 12.6 | 32 |
| 40 | Overexpression of plasmepsin II and plasmepsin III does not directly cause reduction in <i>Plasmodium falciparum</i> sensitivity to artesunate, chloroquine and piperaquine. <i>International Journal for Parasitology: Drugs and Drug Resistance</i> , 2019, 9, 16-22. | 3.4 | 32 |
| 41 | Pan-active imidazolopiperazine antimalarials target the <i>Plasmodium falciparum</i> intracellular secretory pathway. <i>Nature Communications</i> , 2020, 11, 1780. | 12.8 | 27 |
| 42 | Reaction hijacking of tyrosine tRNA synthetase as a new whole-of-life-cycle antimalarial strategy. <i>Science</i> , 2022, 376, 1074-1079. | 12.6 | 25 |
| 43 | Structure of a putative ancestral protein encoded by a single sequence repeat from a multidomain proteinase inhibitor gene from <i>Nicotiana glauca</i> . <i>Structure</i> , 1999, 7, 793-802. | 3.3 | 21 |
| 44 | The solution structure of C1-T1, a two-domain proteinase inhibitor derived from a circular precursor protein from <i>Nicotiana glauca</i> Edited by P. E. Wright. <i>Journal of Molecular Biology</i> , 2001, 306, 69-79. | 4.2 | 20 |
| 45 | Nedd8 hydrolysis by UCH proteases in <i>Plasmodium</i> parasites. <i>PLoS Pathogens</i> , 2019, 15, e1008086. | 4.7 | 19 |
| 46 | An integrated strategy for efficient vector construction and multi-gene expression in <i>Plasmodium falciparum</i> . <i>Malaria Journal</i> , 2013, 12, 373. | 2.3 | 18 |
| 47 | PfMFR3: A Multidrug-Resistant Modulator in <i>Plasmodium falciparum</i> . <i>ACS Infectious Diseases</i> , 2021, 7, 811-825. | 3.8 | 16 |
| 48 | Arresting malaria parasite egress from infected red blood cells. <i>Nature Chemical Biology</i> , 2008, 4, 161-162. | 8.0 | 9 |
| 49 | Defining multiplicity of vector uptake in transfected <i>Plasmodium</i> parasites. <i>Scientific Reports</i> , 2020, 10, 10894. | 3.3 | 9 |
| 50 | The Key Glycolytic Enzyme Phosphofructokinase Is Involved in Resistance to Antiplasmodial Glycosides. <i>MBio</i> , 2020, 11, . | 4.1 | 5 |
| 51 | Lumefantrine attenuates <i>Plasmodium falciparum</i> artemisinin resistance during the early ring stage. <i>International Journal for Parasitology: Drugs and Drug Resistance</i> , 2021, 17, 186-190. | 3.4 | 3 |
| 52 | Body weight satisfaction and disordered eating among youth who are active in sport in Singapore. <i>Pedagogs, Psychology, Medical-Biological Problems of Physical Training and Sports</i> , 2015, 19, 51-58. | 0.4 | 1 |
| 53 | Scientists on a RAMPAGE to find apicomplexan transcription start sites. <i>Nature Reviews Microbiology</i> , 2021, 19, 483-483. | 28.6 | 0 |