Maja Malmberg

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Polymorphisms in Plasmodium falciparum Chloroquine Resistance Transporter and Multidrug Resistance 1 Genes: Parasite Risk Factors That Affect Treatment Outcomes for P. falciparum Malaria After Artemether-Lumefantrine and Artesunate-Amodiaquine. American Journal of Tropical Medicine and Hygiene, 2014, 91, 833-843. | 1.4 | 204 |
| 2 | Plasmodium falciparum Drug Resistance Phenotype as Assessed by Patient Antimalarial Drug Levels and Its Association With pfmdr1 Polymorphisms. Journal of Infectious Diseases, 2013, 207, 842-847. | 4.0 | 99 |
| 3 | Novel Polymorphisms in Plasmodium falciparum ABC Transporter Genes Are Associated with Major ACT Antimalarial Drug Resistance. PLoS ONE, 2011, 6, e20212. | 2.5 | 80 |
| 4 | Temporal trends of molecular markers associated with artemether-lumefantrine tolerance/resistance in Bagamoyo district, Tanzania. Malaria Journal, 2013, 12, 103. | 2.3 | 62 |
| 5 | High similarity in the microbiota of cold-water sponges of the Genus <i>Mycale</i> from two different geographical areas. PeerJ, 2018, 6, e4935. | 2.0 | 62 |
| 6 | Efficacy and Effectiveness of Artemether-Lumefantrine after Initial and Repeated Treatment in Children < 5 Years of Age with Acute Uncomplicated Plasmodium falciparum Malaria in Rural Tanzania: A Randomized Trial. Clinical Infectious Diseases, 2011, 52, 873-882. | 5.8 | 58 |
| 7 | Effectiveness of artemether-lumefantrine provided by community health workers in under-five children with uncomplicated malaria in rural Tanzania: an open label prospective study. Malaria Journal, 2011, 10, 64. | 2.3 | 39 |
| 8 | <i>pfmdr1</i> Amplification Is Related to Increased Plasmodium falciparum In Vitro Sensitivity to the Bisquinoline Piperaquine. Antimicrobial Agents and Chemotherapy, 2012, 56, 3615-3619. | 3.2 | 34 |
| 9 | Serological and molecular study of Crimean-Congo Hemorrhagic Fever Virus in cattle from selected districts in Uganda. Journal of Virological Methods, 2021, 290, 114075. | 2.1 | 28 |
| 10 | Disentangling the Amyloid Pathways: A Mechanistic Approach to Etiology. Frontiers in Neuroscience, 2020, 14, 256. | 2.8 | 21 |
| 11 | Complete Genome Sequence of an African Swine Fever Virus Isolate from Sardinia, Italy. Genome Announcements, 2016, 4, . | 0.8 | 19 |
| 12 | Morphological and molecular identification of ixodid tick species (Acari: Ixodidae) infesting cattle in Uganda. Parasitology Research, 2020, 119, 2411-2420. | 1.6 | 18 |
| 13 | The evolution of African swine fever virus in Sardinia (1978 to 2014) as revealed by whole genome sequencing and comparative analysis. Transboundary and Emerging Diseases, 2020, 67, 1971. | 3.0 | 18 |
| 14 | General and Local Morphological Anomalies in Amblyomma lepidum (Acari: Ixodidae) and Rhipicephalus decoloratus Infesting Cattle in Uganda. Journal of Medical Entomology, 2019, 56, 873-877. | 1.8 | 16 |
| 15 | A review of congenital tremor type A-II in piglets. Animal Health Research Reviews, 2020, 21, 84-88. | 3.1 | 15 |
| 16 | Alternatively spliced transcripts and novel pseudogenes of the Plasmodium falciparum resistance-associated locus pfcrt detected in East African malaria patients. Journal of Antimicrobial Chemotherapy, 2015, 70, 116-123. | 3.0 | 14 |
| 17 | Sustained High Cure Rate of Artemether–Lumefantrine against Uncomplicated Plasmodium falciparum Malaria after 8 Years of Its Wide-Scale Use in Bagamoyo District, Tanzania. American Journal of Tropical Medicine and Hygiene, 2017, 97, 526-532. | 1.4 | 12 |
| 18 | Detection of atypical porcine pestivirus in Swedish piglets with congenital tremor type A-II. BMC Veterinary Research, 2020, 16, 260. | 1.9 | 11 |

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|----|---|-----|-----------|
| 19 | Clinical and Molecular Epidemiology of Crimean-Congo Hemorrhagic Fever in Humans in Uganda, 2013–2019. American Journal of Tropical Medicine and Hygiene, 2022, 106, 88-98. | 1.4 | 9 |
| 20 | Evolutionary genetics of canine respiratory coronavirus and recent introduction into Swedish dogs. Infection, Genetics and Evolution, 2020, 82, 104290. | 2.3 | 8 |
| 21 | Prevalence of and Risk Factors Associated with Polymerase Chain Reaction-Determined Plasmodium falciparum Positivity on Day 3 after Initiation of Artemether–Lumefantrine Treatment for Uncomplicated Malaria in Bagamoyo District, Tanzania. American Journal of Tropical Medicine and Hygiene. 2019. 100. 1179-1186. | 1.4 | 8 |
| 22 | The heart microbiome of insectivorous bats from Central and South Eastern Europe. Comparative Immunology, Microbiology and Infectious Diseases, 2021, 75, 101605. | 1.6 | 7 |
| 23 | Single nucleotide polymorphisms in Plasmodium falciparum V type H+ pyrophosphatase gene (pfvp2) and their associations with pfcrt and pfmdr1 polymorphisms. Infection, Genetics and Evolution, 2014, 24, 111-115. | 2.3 | 6 |
| 24 | The Genome of Setaria digitata: A Cattle Nematode Closely Related to Human Filarial Parasites. Genome Biology and Evolution, 2020, 12, 3971-3976. | 2.5 | 5 |
| 25 | Four novel picornaviruses detected in Magellanic Penguins (Spheniscus magellanicus) in Chile. Virology, 2021, 560, 116-123. | 2.4 | 5 |
| 26 | Atypical porcine pestivirus—A widespread virus in the Swedish wild boar population. Transboundary and Emerging Diseases, 2022, 69, 2349-2360. | 3.0 | 5 |
| 27 | Identification and molecular characterization of highly divergent RNA viruses in cattle, Uganda Virus Research, 2022, 313, 198739. | 2.2 | 4 |
| 28 | Phylogenomic analysis of the complete sequence of a gastroenteritis-associated cetacean adenovirus (bottlenose dolphin adenovirus 1) reveals a high degree of genetic divergence. Infection, Genetics and Evolution, 2017, 53, 47-55. | 2.3 | 2 |
| 29 | OCCURRENCE OF DAY 3 SUBMICROSCOPIC <i>PLASMODIUM FALCIPARUM</i> PARASITAEMIA BEFORE AND AFTER IMPLEMENTATION OF ARTEMETHER-LUMEFANTRINE TREATMENT POLICY IN TANZANIA. BMJ Global Health, 2017, 2, A16.3-A17. | 4.7 | 0 |