



Supplementary Figure S4: **Phylogenetic tree of 27 ClpC genes.** Bayesian reconstruction under the $GTR_{nt} + \Gamma_4$ model. Edges with $PP < 0.9$ were collapsed, and $PP = 1$ are not shown. The ClpC phylogeny supports the monophyly of mammalian malaria parasites ($PP = 0.99$).