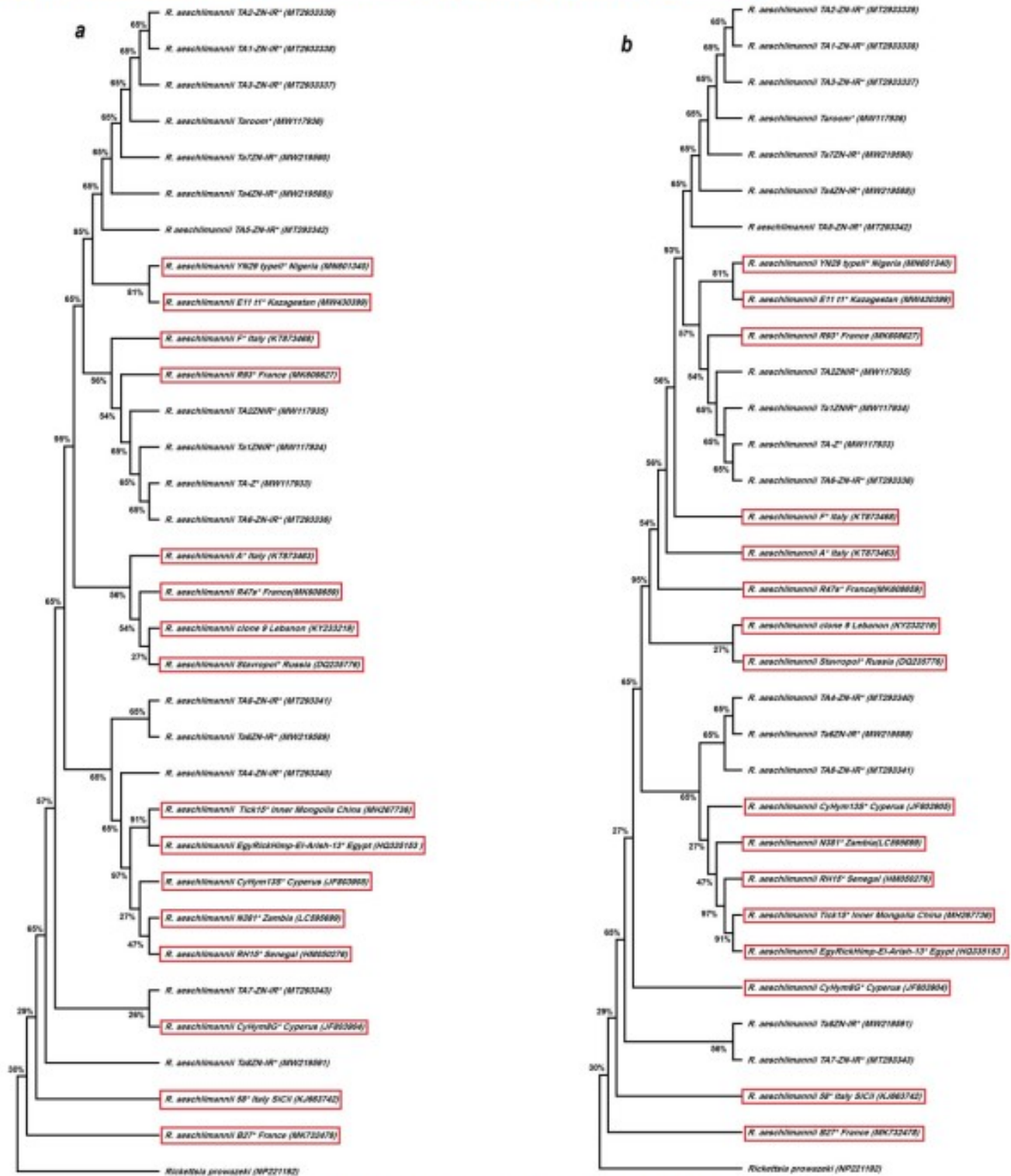


Figure 5 Phylogenetic tree based on the sequences of PCR amplicons of the *gltA* gene of *Rickettsia aeschlimannii* detected in *Hyalomma marginatum* with the Maximum Likelihood (a) and Neighbor-Joining (b) methods



Note: Percentages indicate bootstrap values based on 1000 replicates. The reference sequences in this study are represented in red box.