



**Additional Figure 3:**

Bayesian phylogeny of 89 unique *ATPSb-III* alleles. Numbers at branches correspond to posterior probability values from Bayesian analysis and Maximum Likelihood non-parametric bootstrap proportions, separated by a forward slash. Numbers with "A" prefix indicate allele number (compare Additional Table 1 and Fig. 4). Tree rooted with alleles from the Indian Ocean. Clades colored according to their main geographic distribution as in Figure 3.