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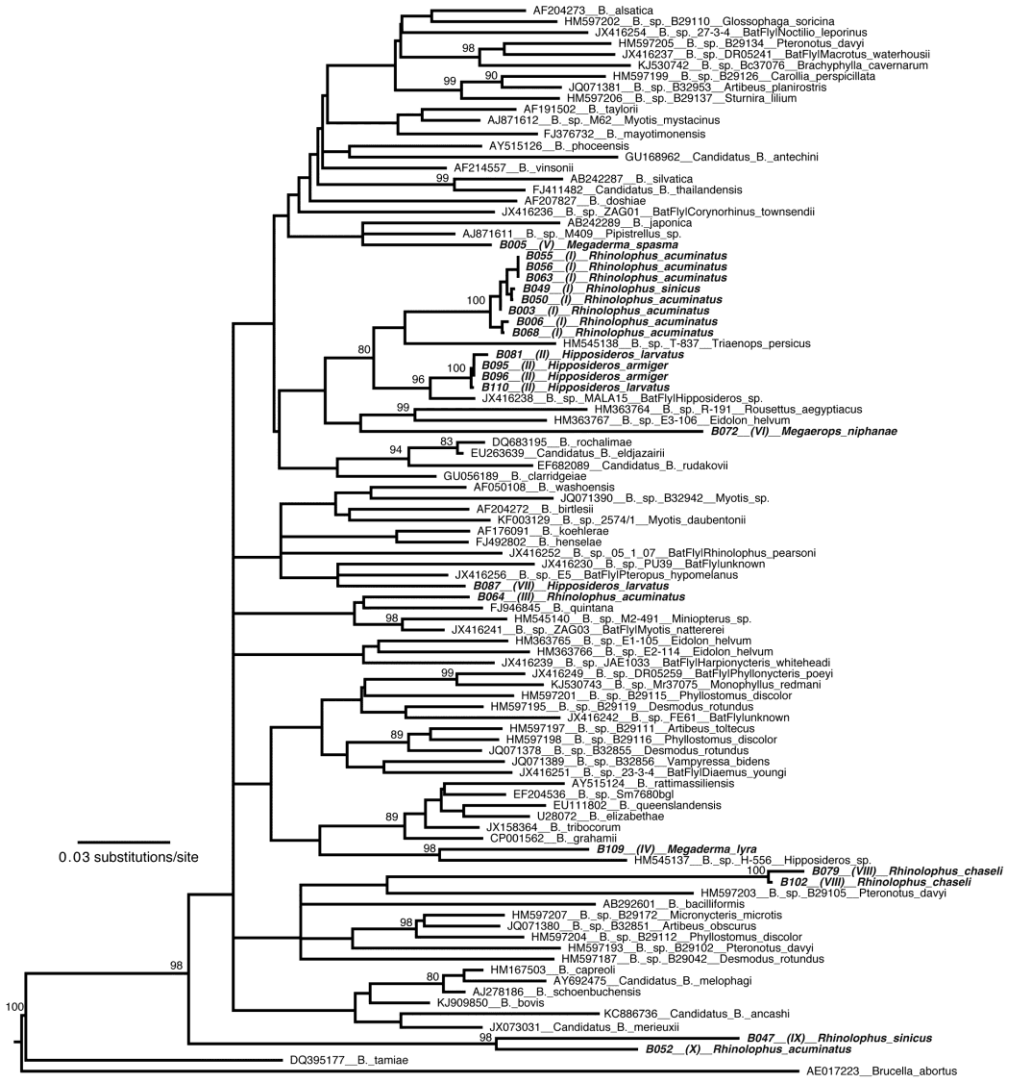
Diversity of *Bartonella* spp. in Bats, Southern Vietnam

Technical Appendix



Technical Appendix Figure. Neighbor-joining phylogeny of a 327-nt region of the citrate synthase A gene of 21 *Bartonella* spp. detected in bats in southern Vietnam and a sample of diverse *Bartonella* spp. causing infections in animals and humans globally. *Brucella abortus* (AE017223) is included as an outgroup. Bootstrap support values are shown for nodes with >80% support. Taxa names for sequences determined in this study are indicated in bold italics, phylogroups are indicated in parentheses (I–X), and bat species are indicated after phylogroups.