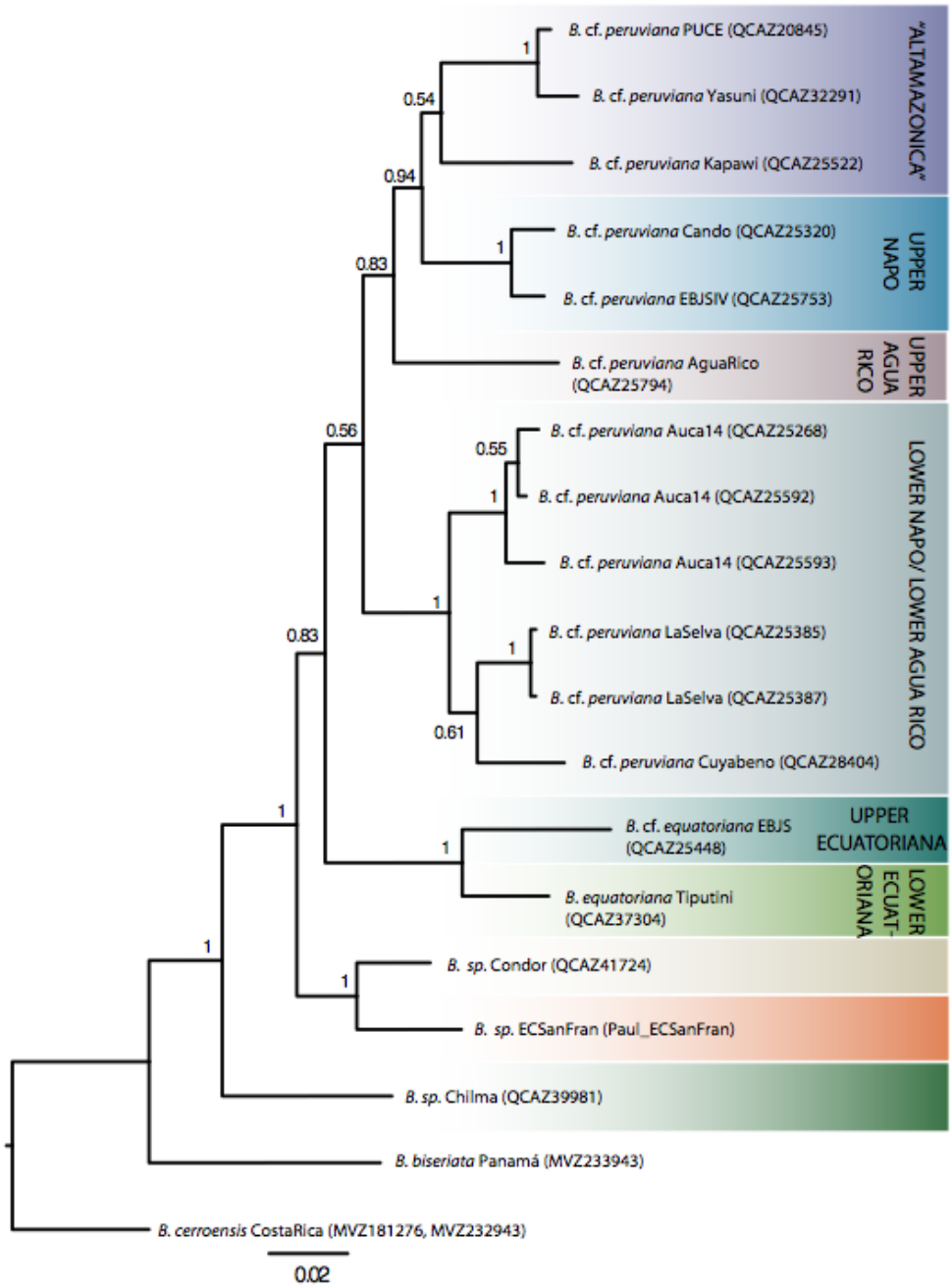
Additional file 5 - Multilocus combined phylogeny

The 50% majority-rule phylogenetic tree inferred from combined Bayesian analyses of *Cytb* and *Rag1* sequence data. Posterior probabilities are located at nodes or above branches. Clades referred to in the text are noted in vertical. See Table 1 and Figure 1 for more information on sample localities.