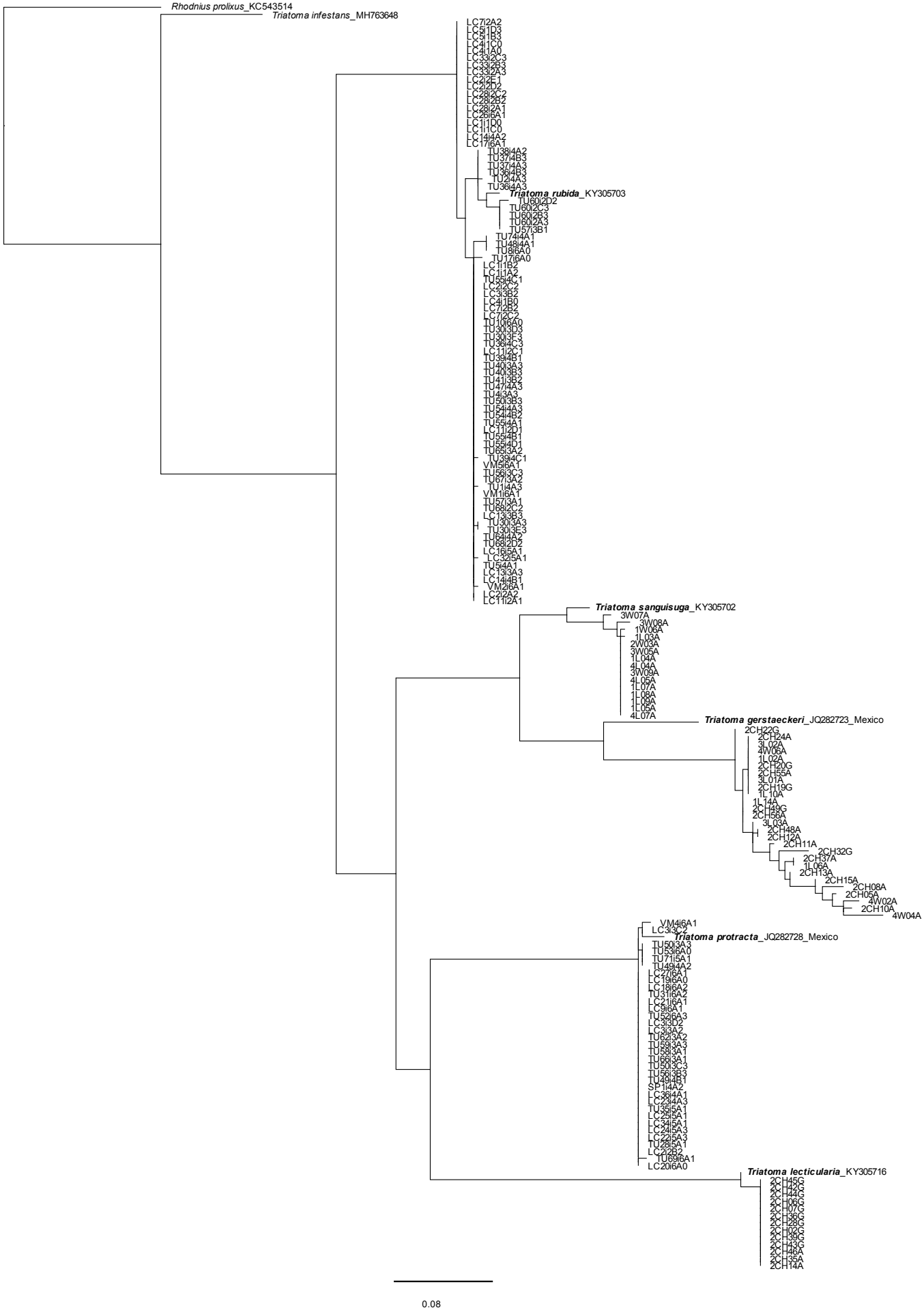

Additional File 4: ML based topology inferred from 351 bp long alignment of *coxB* sequences from 170 *T. cruzi* negative *Triatoma* individuals included in the "*ultraclean*" dataset. Sequences retrieved from GenBank are designated with the accession numbers.

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