



Figure S5: RAxML phylogenetic tree of the elongation factor EF-1 α gene among the Coccidae family. Colors indicate the morphological subfamilies: Eulecaniinae, Filippiinae, Coccinae, Cardiococcinae, Ceroplastinae, Eriopeltinae, and Myzolecaniinae, to which *T. parvicornis* belongs. In red is the Italian isolate of *T. parvicornis*. The numbers in the branch labels indicate the bootstrap values.