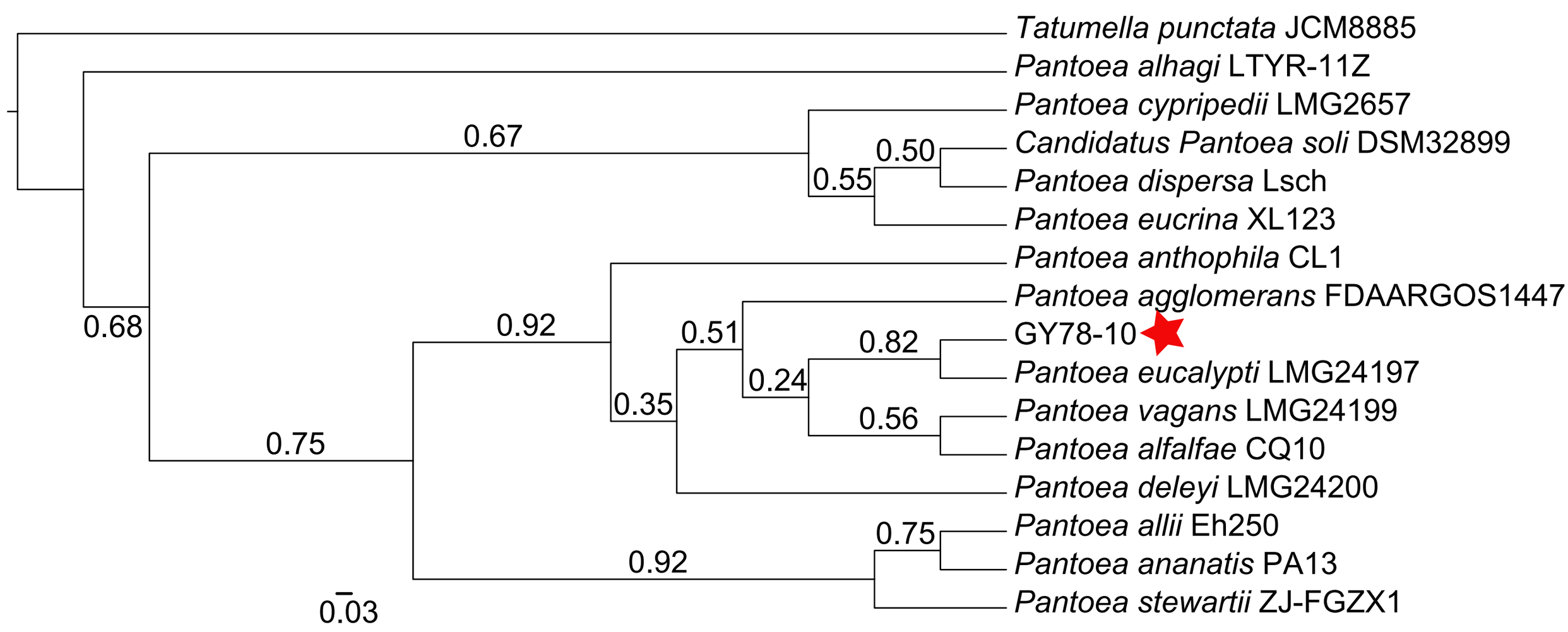


Figure S2. The whole-genome phylogenetic analysis of isolate GY78-10. The phylogenetic treewas constructed using the Orthofind program based on single copy orthologs of bacterial strains from the *Pantoea* species, with *Tatumella citrea* (Strain: DSM13699) designated as the outgroup. The red star indicates the phylogenetic position of isolate GY78-10.