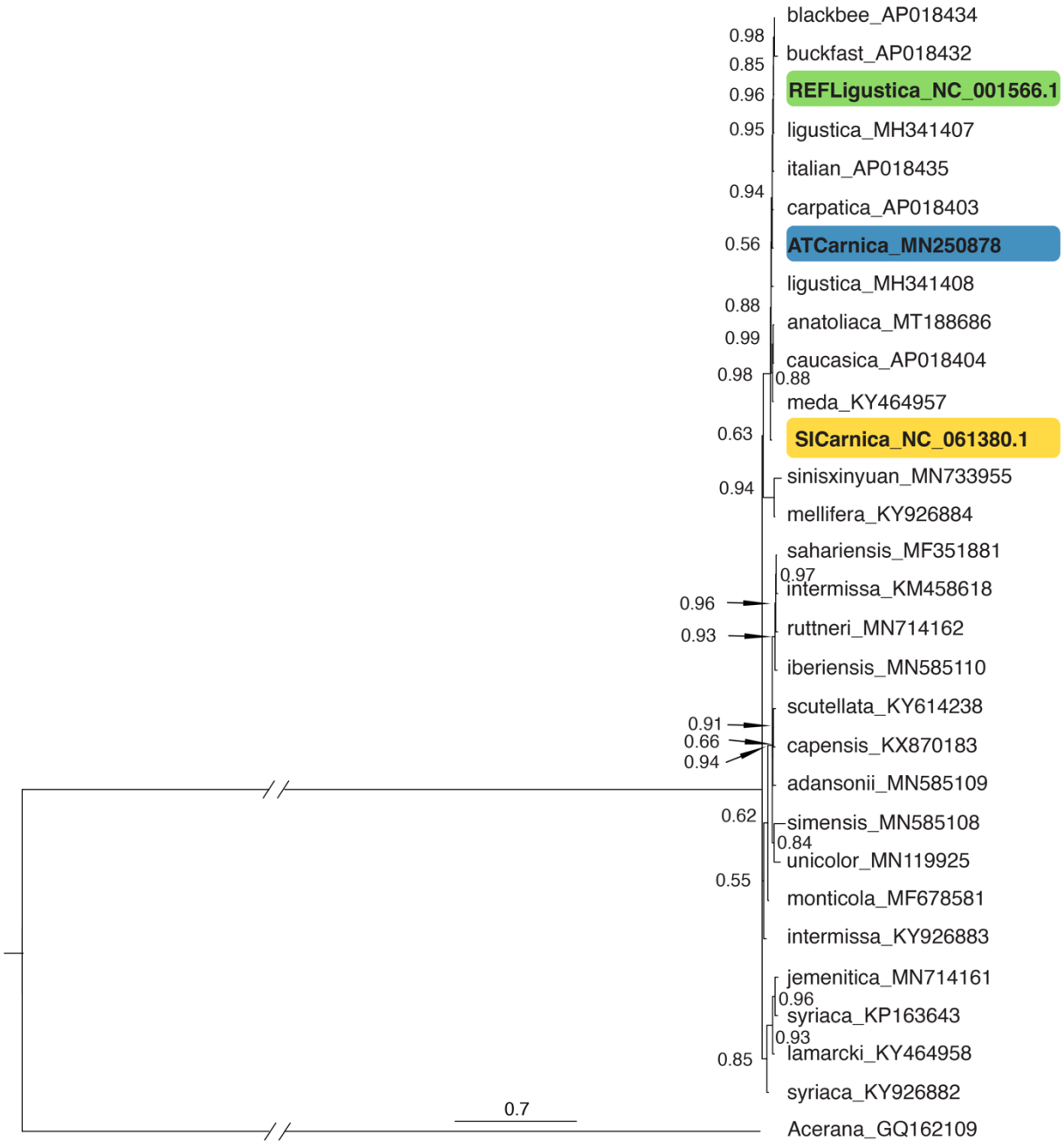

Figure S2. Bayesian phylogenetic analysis of selected *A. mellifera* subspecies using PhyloBayes software based on complete mitogenome sequences. Computed branch lengths are displayed. Posterior probabilities are presented on the nodes. Names of the samples include subspecies or strain name and GenBank accession number.