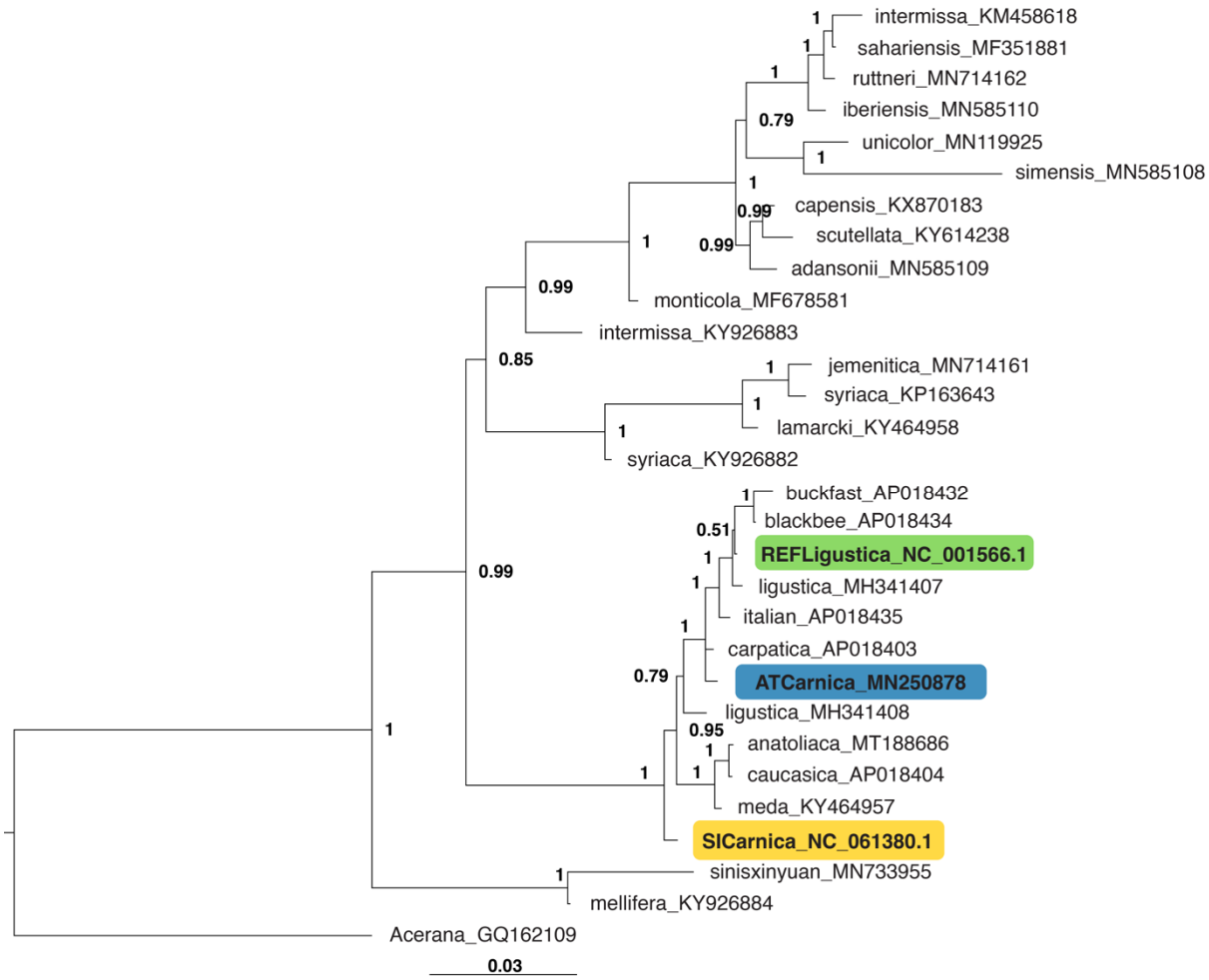

Figure S6. Bayesian phylogenetic analysis of selected *A. mellifera* subspecies using MrBayes software based on partial mitogenome sequences (Noisy dataset). Computed branch lengths are displayed. Posterior probabilities are presented on the nodes. Names of the samples include subspecies or strain name and GenBank accession number.