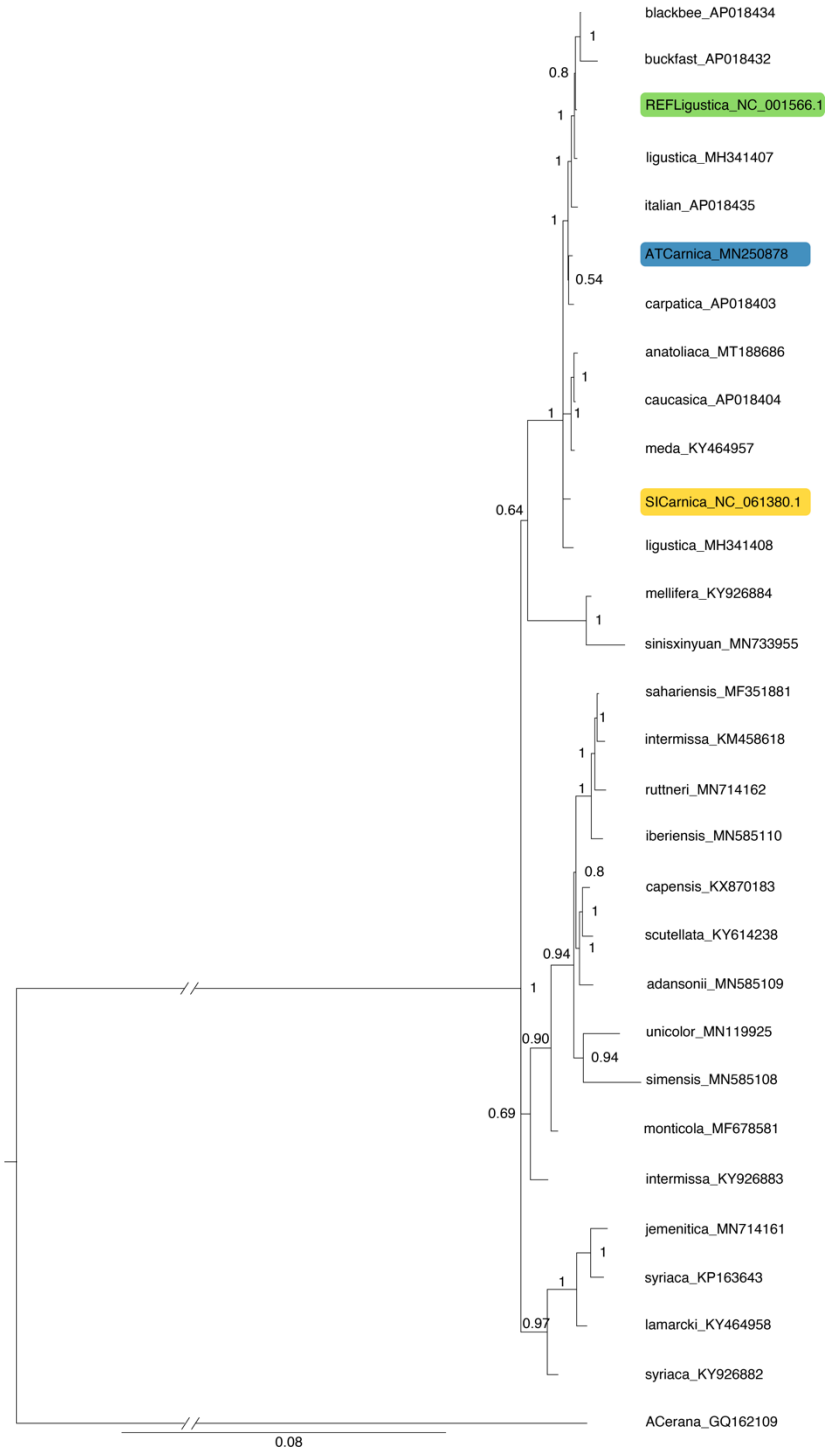

Figure S5. Bayesian phylogenetic analysis of selected *A. mellifera* subspecies using MrBayes software based on partial mitogenome sequences (GBLOCKS dataset). Computed branch lengths are displayed. Posterior probabilities are presented on the nodes. Names of the samples include subspecies or strain name and GenBank accession number.