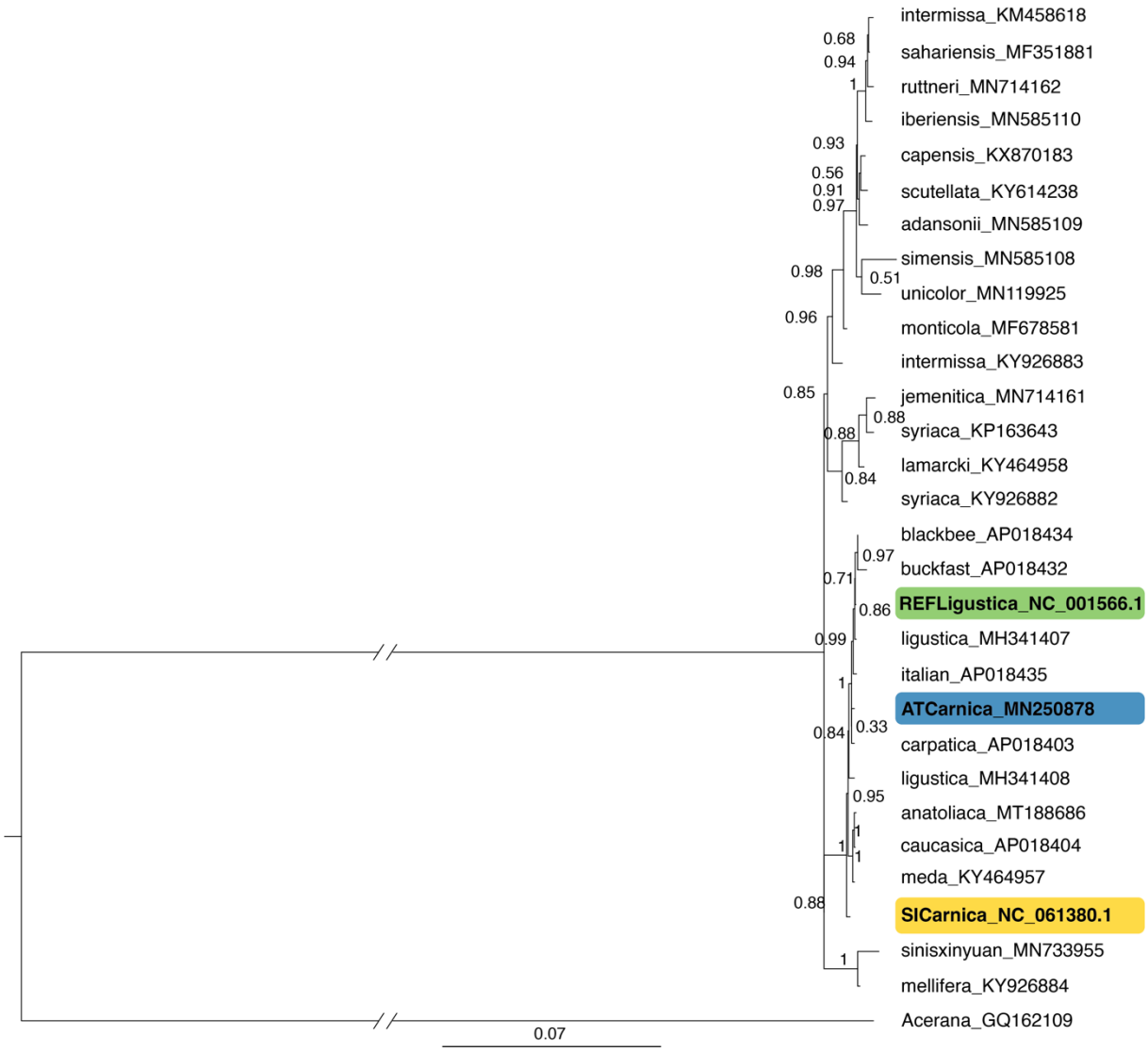

Figure S1. Maximum likelihood phylogenetic analysis of selected *A. mellifera* subspecies based on complete mitogenome sequences. Computed branch lengths are displayed. Bootstrap support values are presented on the nodes. Names of the samples include subspecies or strain name and GenBank accession number.