

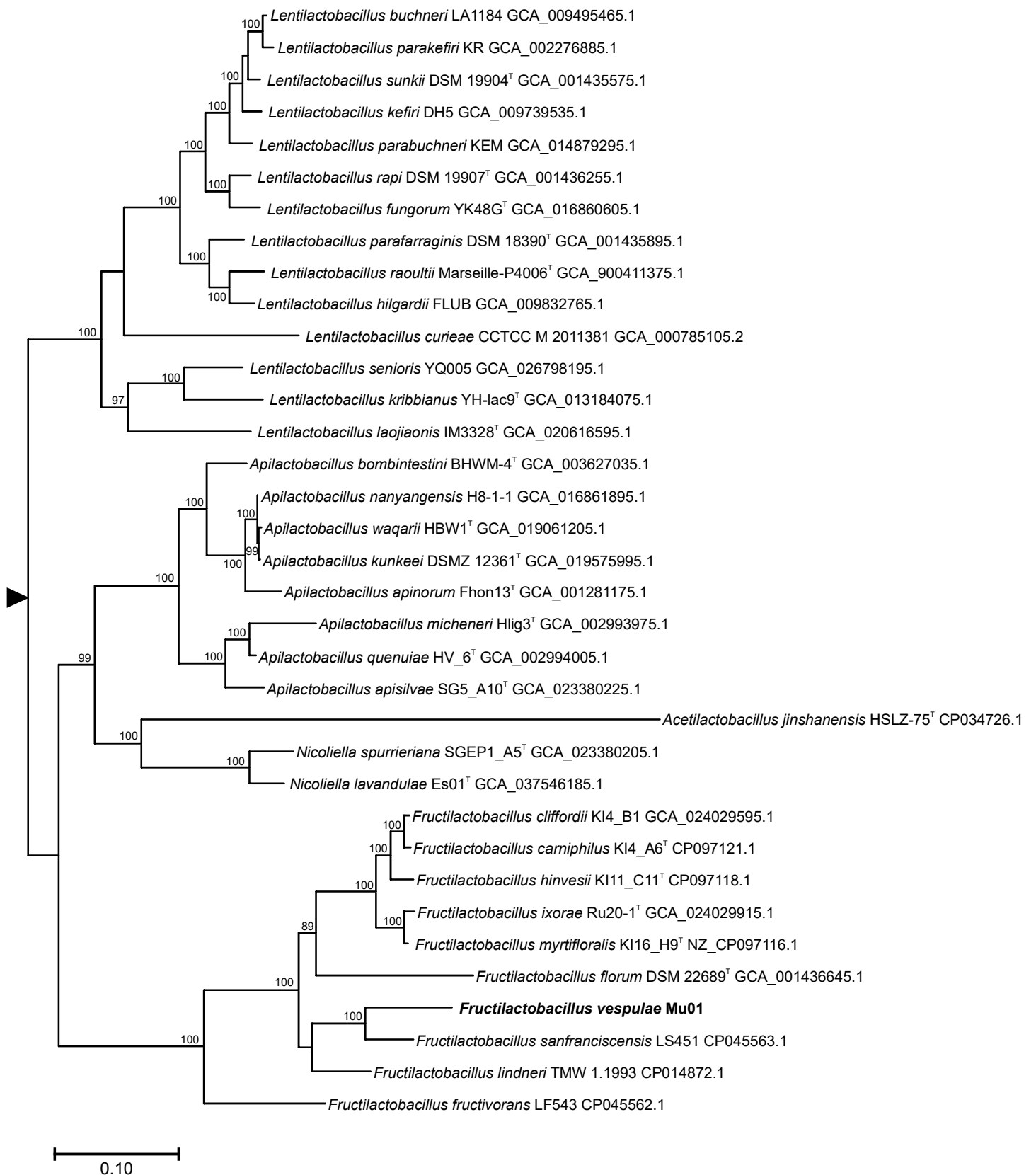


Figure S1. Maximum likelihood phylogenetic tree of core genes of species of *Fructilactobacillus* and closely related genera. Support values are given for those nodes with support higher than 75%. The tree has been arbitrarily rooted for ease of visualization.