

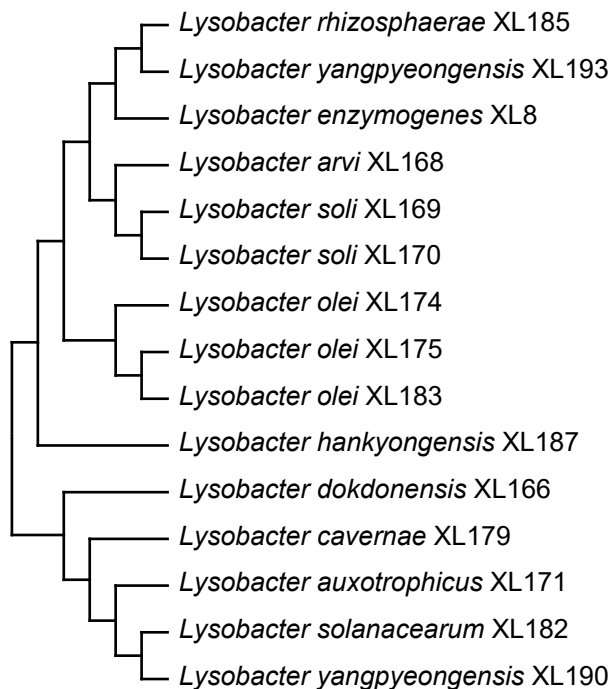


Figure S2. Taxonomy and phylogenetic tree of *Lysobacter* spp. isolates used in this study. The evolutionary history was inferred using the Neighbor-Joining method.