

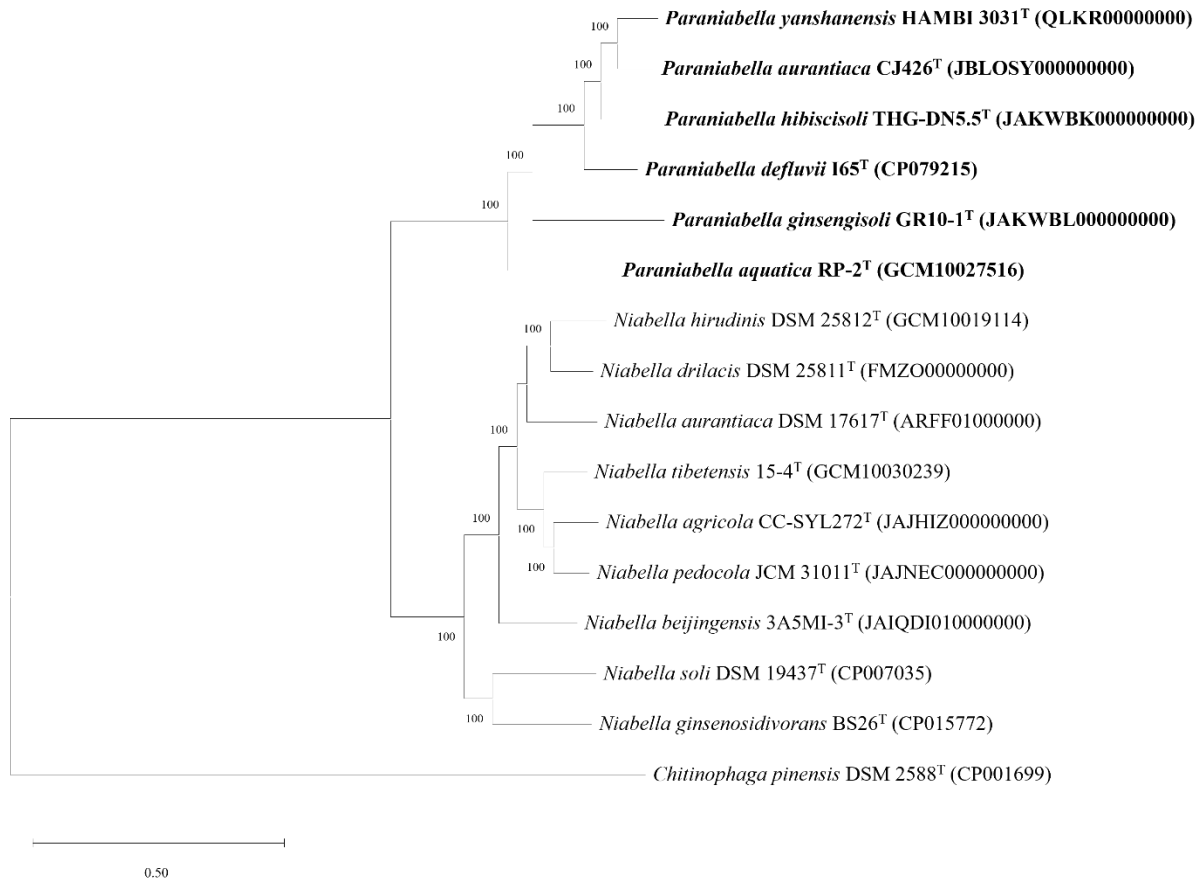


Fig. S4. Phylogenomic tree of strain CJ426^T and related type strains based on the concatenated sequences of 290 core genes. Bootstrap values are shown at branch points based on analyses of 1,000 replicated datasets. *Chitinophaga pinensis* DSM 2588^T was used as an outgroup.