# Biodegradation of Oil by a Newly Isolated Strain *Acinetobacter junii* WCO-9 and Its Comparative Pan-Genome Analysis

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# Supplementary Materials

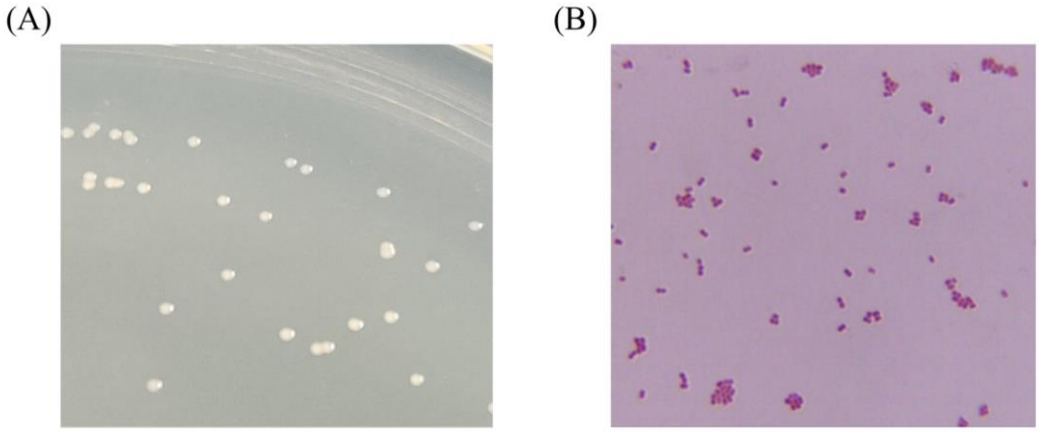


**Figure S1.** Colony morphology and Gram staining of strain WCO-9

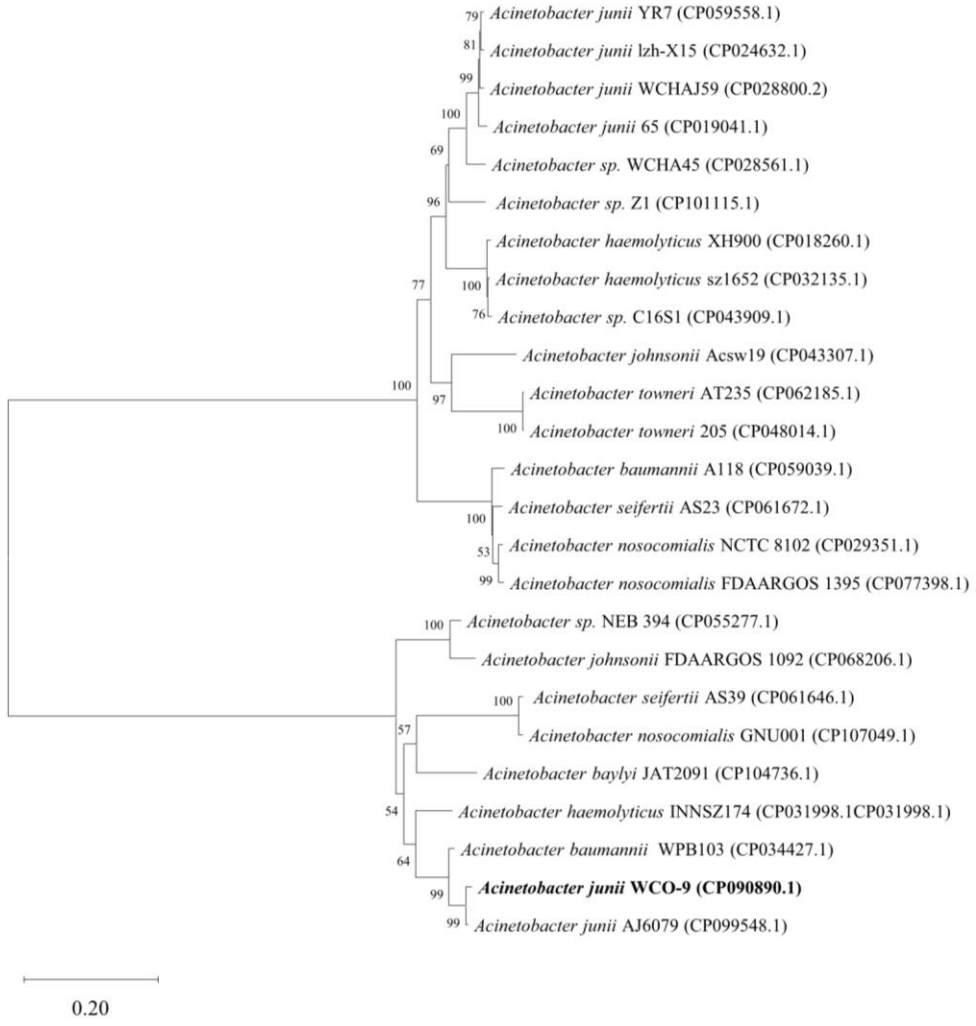


**Figure S2.** Phylogenetic tree based on the *ileS* gene sequence. This tree was constructed by the neighbor-joining method, and showed the phylogenetic relationship between strain WCO-9 and closely related species.

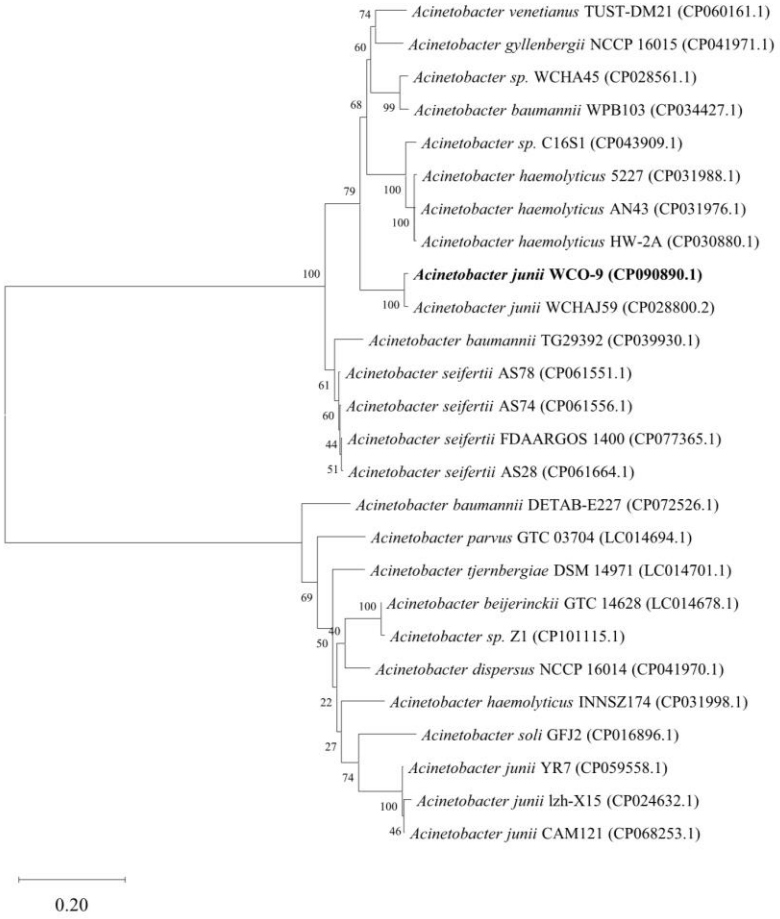


**Figure S3.** Phylogenetic tree based on the *recA* gene sequence. This tree was constructed by the neighbor-joining method, and showed the phylogenetic relationship between strain WCO-9 and closely related species.

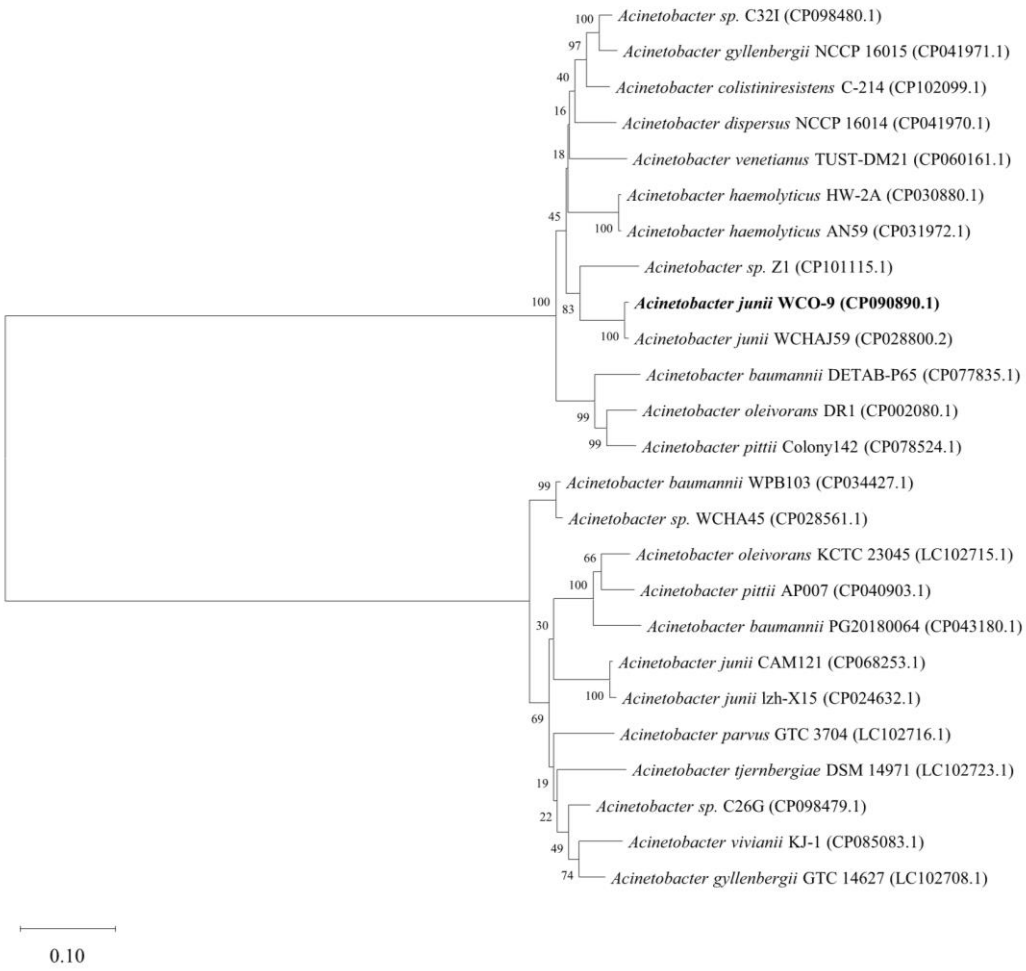


**Figure S4.** Phylogenetic tree based on the *rpoD* gene sequence. This tree was constructed by the neighbor-joining method, and showed the phylogenetic relationship between strain WCO-9 and closely related species.

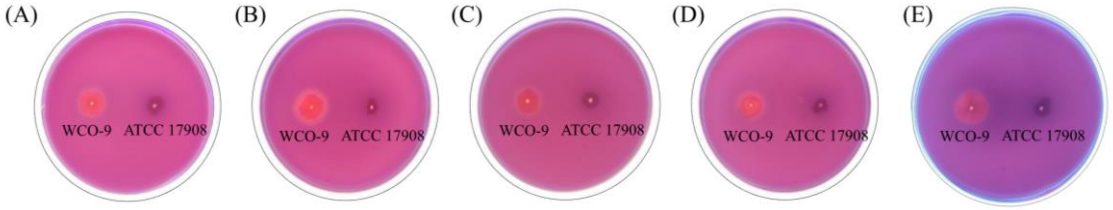


**Figure S5.** Degradation of five oils by strain WCO-9 and ATCC 17908. (A) Corn oil; (B) Peanut oil; (C) Canola oil; (D) Soybean oil; (E) Olive oil.

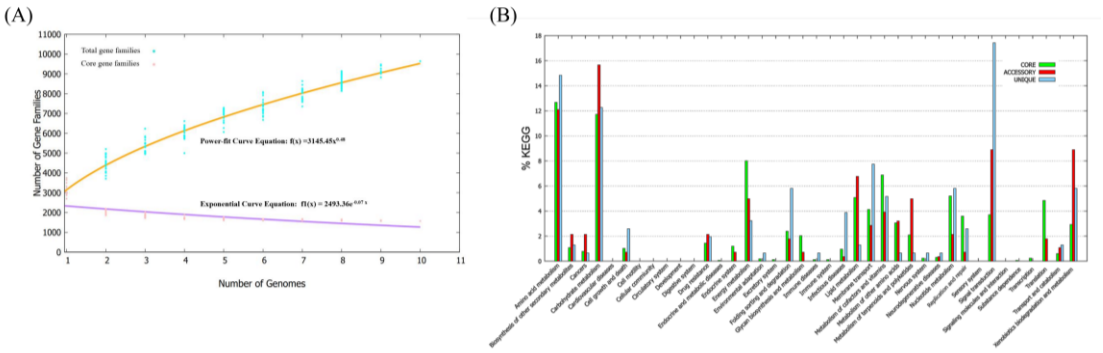


**Figure S6.** Pan-genome analysis of *A junii* strains. (A) Mathematical modeling of the pan-genome and core genome of *A. junii*. (B) Bar chart showing functional proportions (based on KEGG categories) of different parts of the *A. junii* pan-genome (i.e., core, accessory, unique).