

Supplementary file

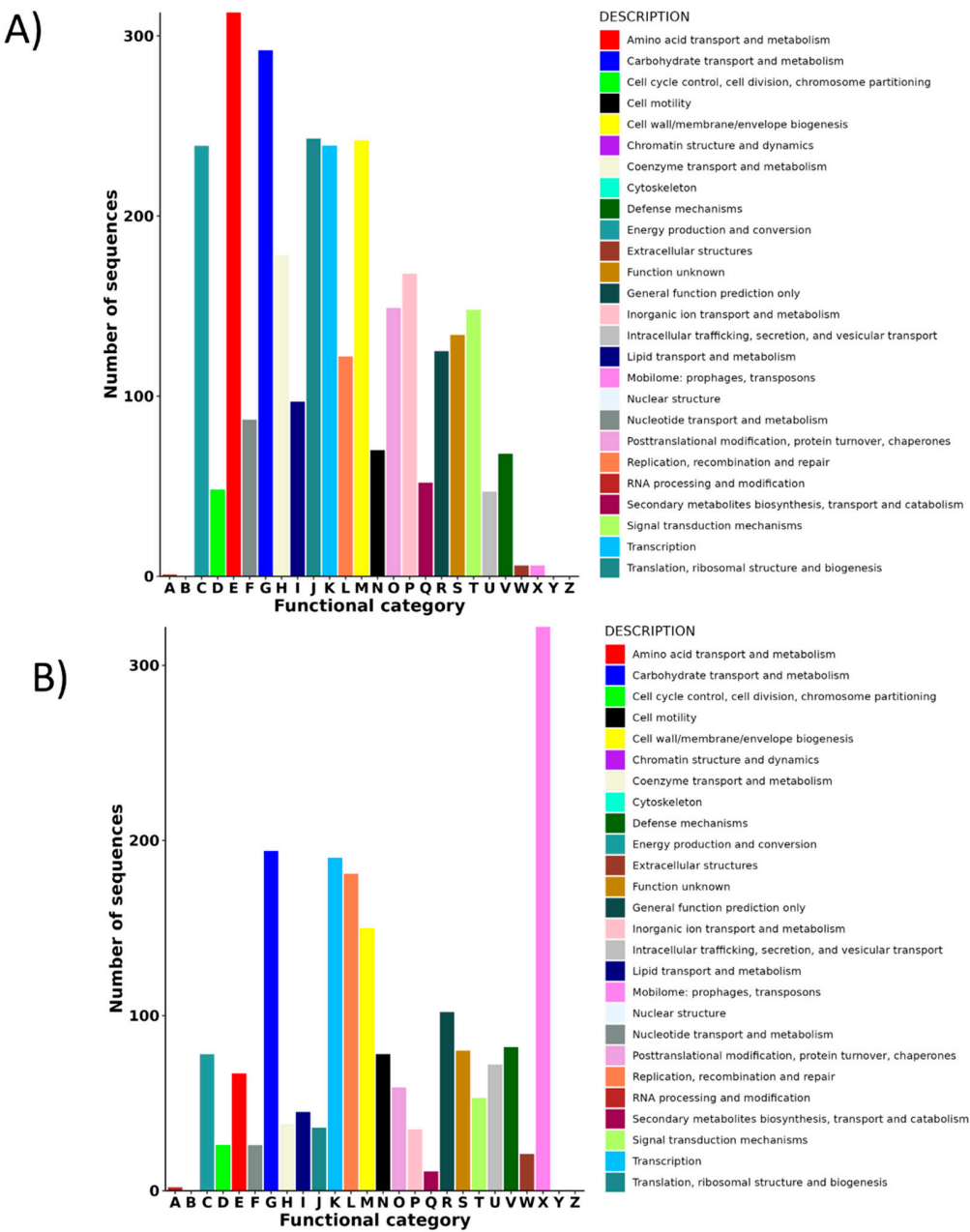


Figure S1. Functional annotation of the core(A) and the accessory(B) genes using the cluster of orthologous genes (COG).

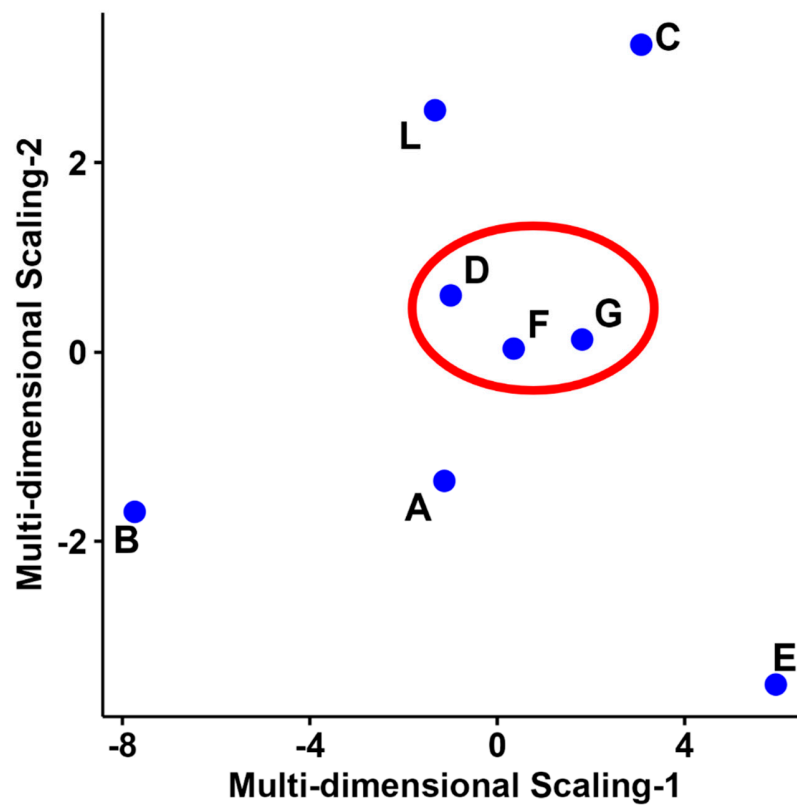


Figure S2. Multi-dimensional scaling (MDS) plot constructed using the geographical location of the farms (8) in R. The distance matrix was computed using each farm's latitude and longitude coordinates. Out of 8 farms, 3 Farms (D, F, and G) were located closer to each other.