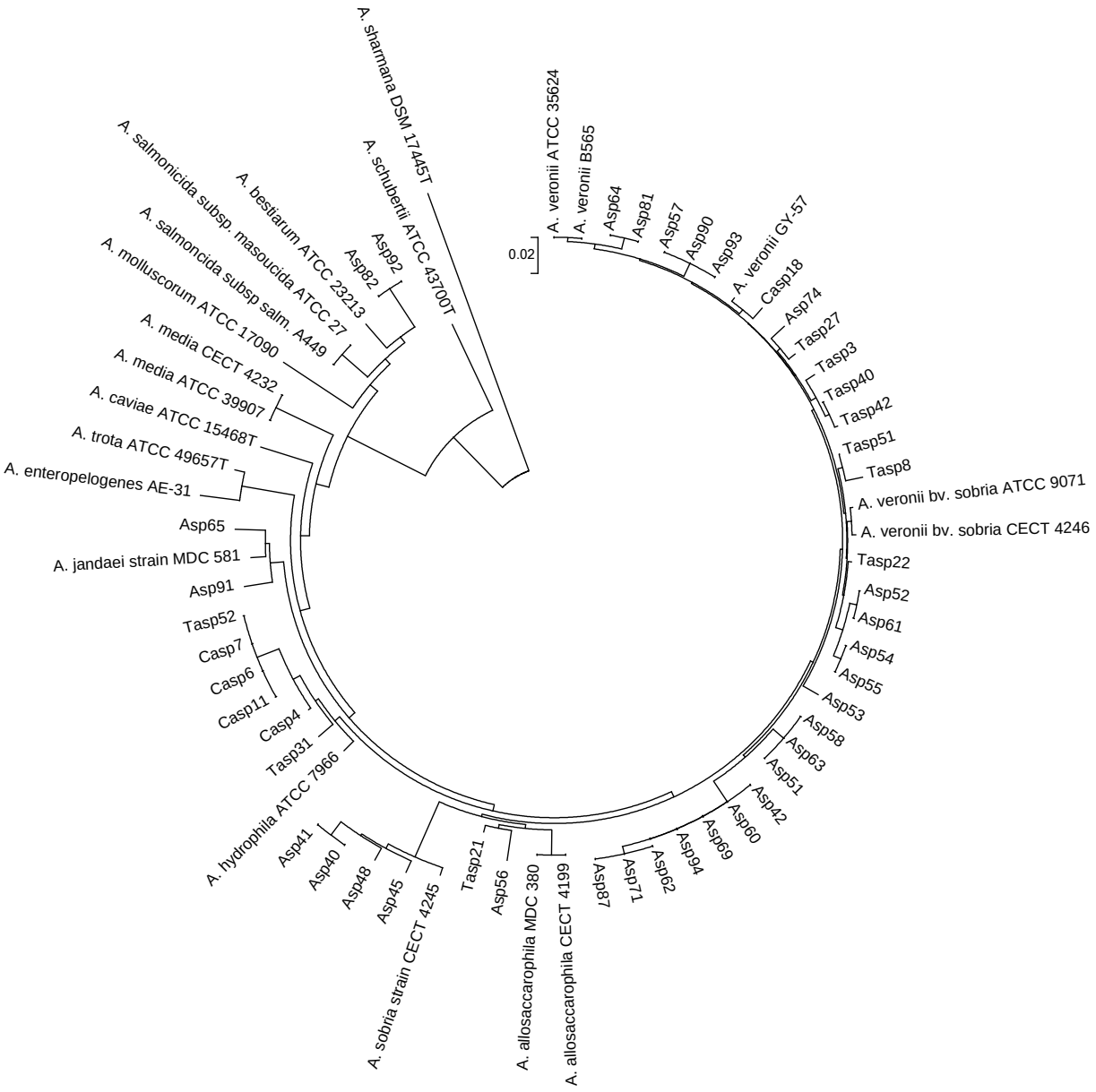

Supplemental Figure 1. Phylogenetic analysis of aquatic *Aeromonas* isolates from the Eastern Basin of Lake Erie. A 480 bps fragment of *gyrB* was sequenced and analyzed from 45 isolates using the Neighbor-Joining method in MEGA5.