

Fig. S3A

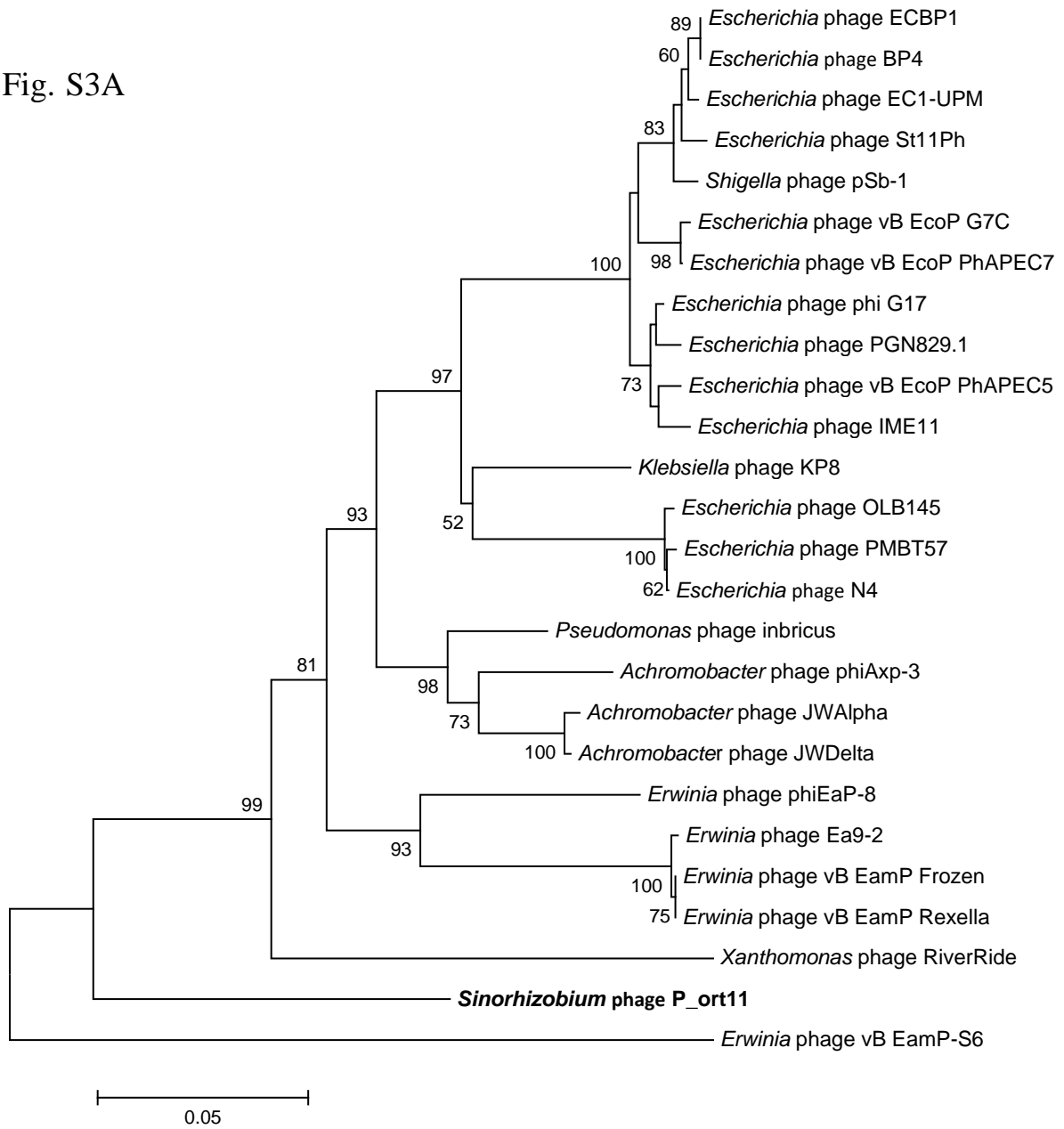


Fig. S3B

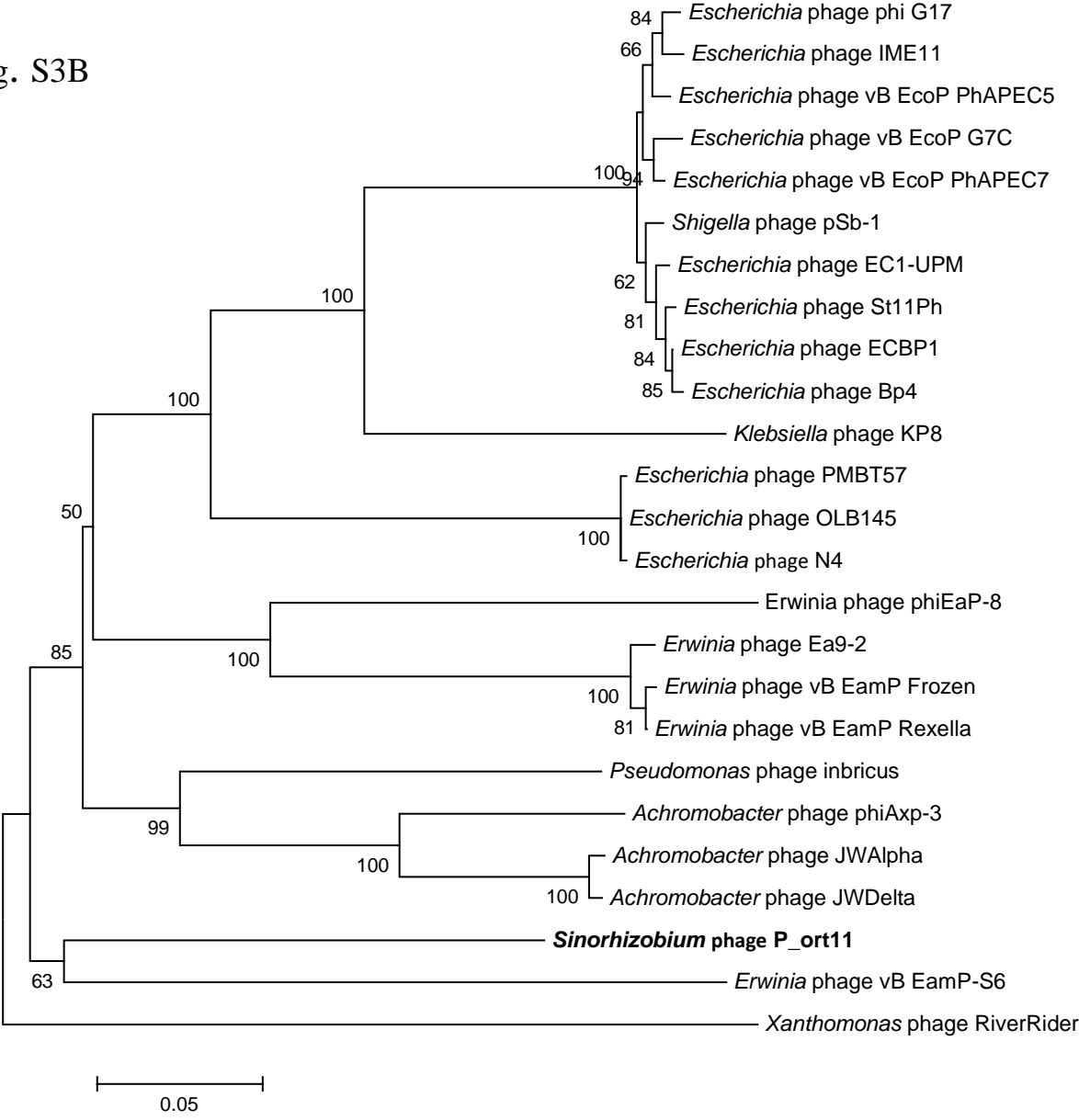
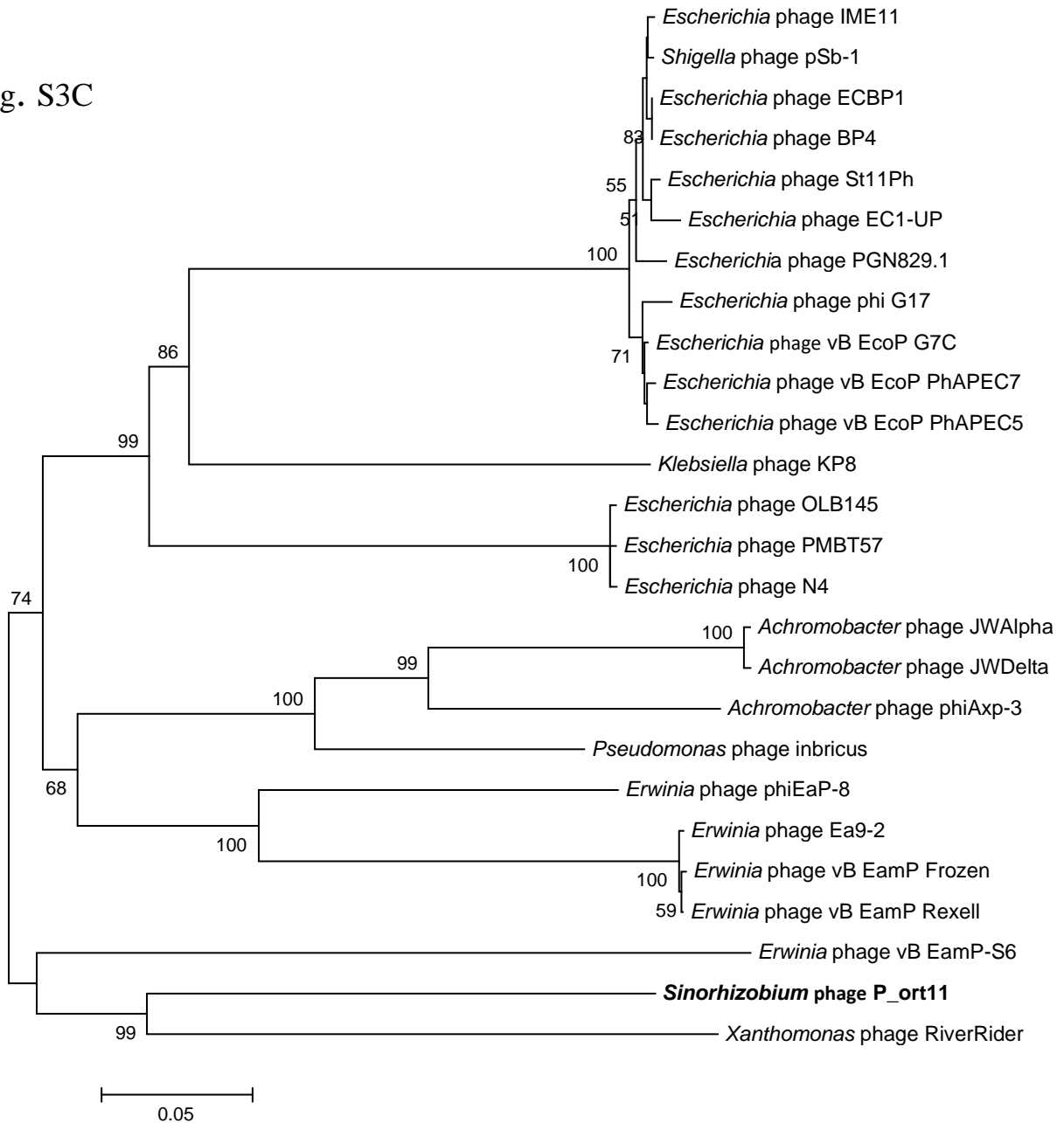


Fig. S3C



Supplementary Figure S3. Phylogenetic trees of the major capsid protein (S3A), portal protein (S3B), and DNA polymerase (S3C) of P_ort11 compared to other N4-like phages. Neighbour-joining tree was constructed based on ClustalW (MEGA5). Bootstrap values (>50%) are expressed as percentages of 1,000 replicated. Bar, 0.05 amino acid substitutions per site.