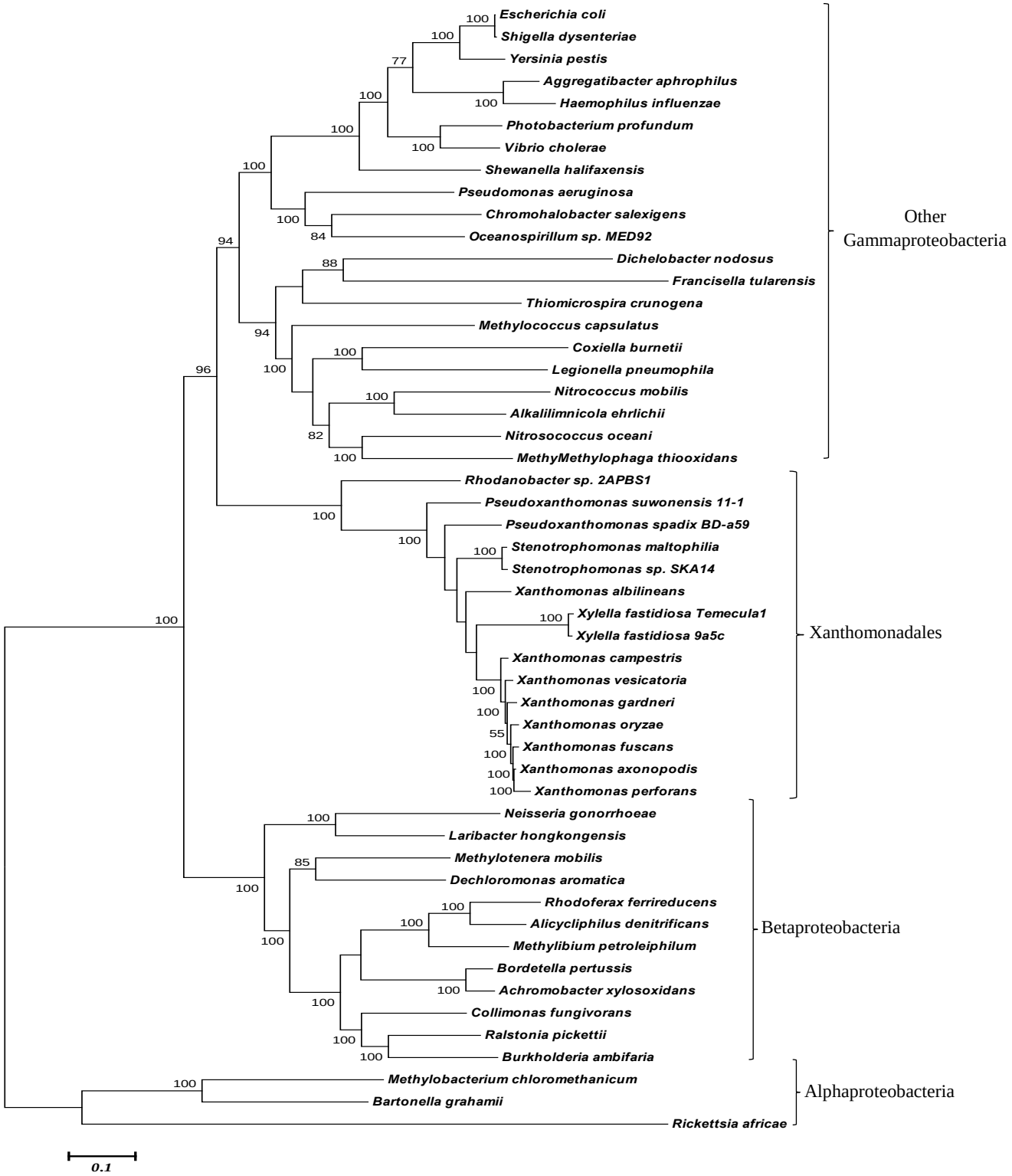

Figure S1

A maximum-likelihood (ML) tree based upon concatenated sequences for 28 conserved global proteins. The tree shows the branching of Xanthomonadales group with Gammaproteobacteria. The Tree the tree was rooted using Alphaproteobacteria. The numbers on the nodes indicate statistical support for the nodes.