



FIGURE S2 Maximum likelihood phylogenetic tree of 16 concatenated ribosomal proteins from Acidobacteria with Candidatus Aminicenantes as the outgroup, with 500 bootstrap replicates. The genomes identified in this study are indicated in bold. The numbers on each node represent the bootstrap support. Colored monophyletic clades indicate a specific Acidobacteria subdivision as determined by the inclusion of a known member. Green = subdivision 3, Purple = subdivision 1, Orange = subdivision 13, Maroon = subdivision 4, Pink = subdivision 8. All grey clades are undetermined subdivisions.