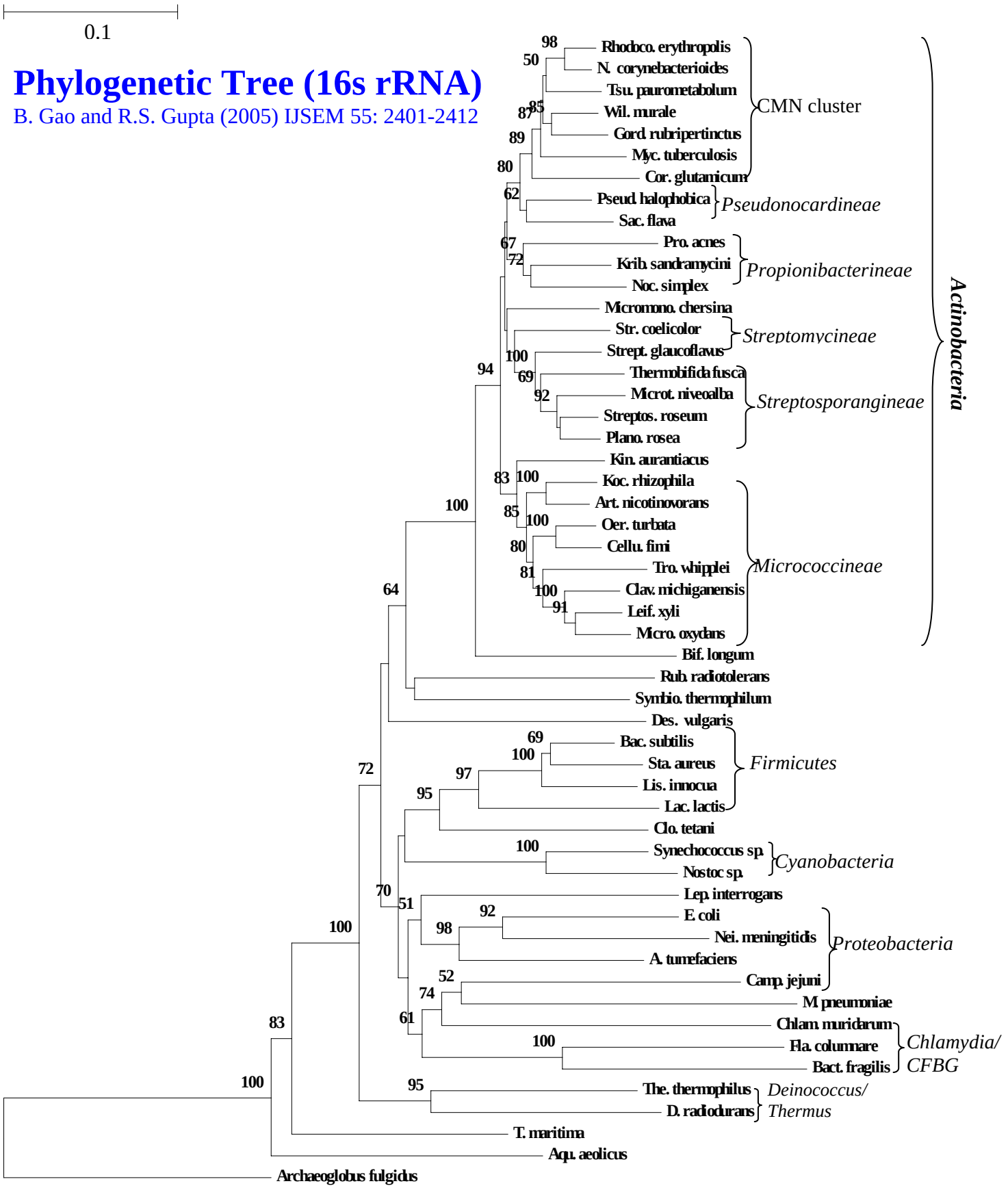

Fig. 5 Neighbor-joining distance tree based on 16s rRNA sequence. The tree is based on 1400 nucleotides, excluding all indels. The root of the tree has been assigned *Archaeoglobus fulgidus* from *Archaea*. The bootstrap scores for various nodes which were higher than 50 are indicated.