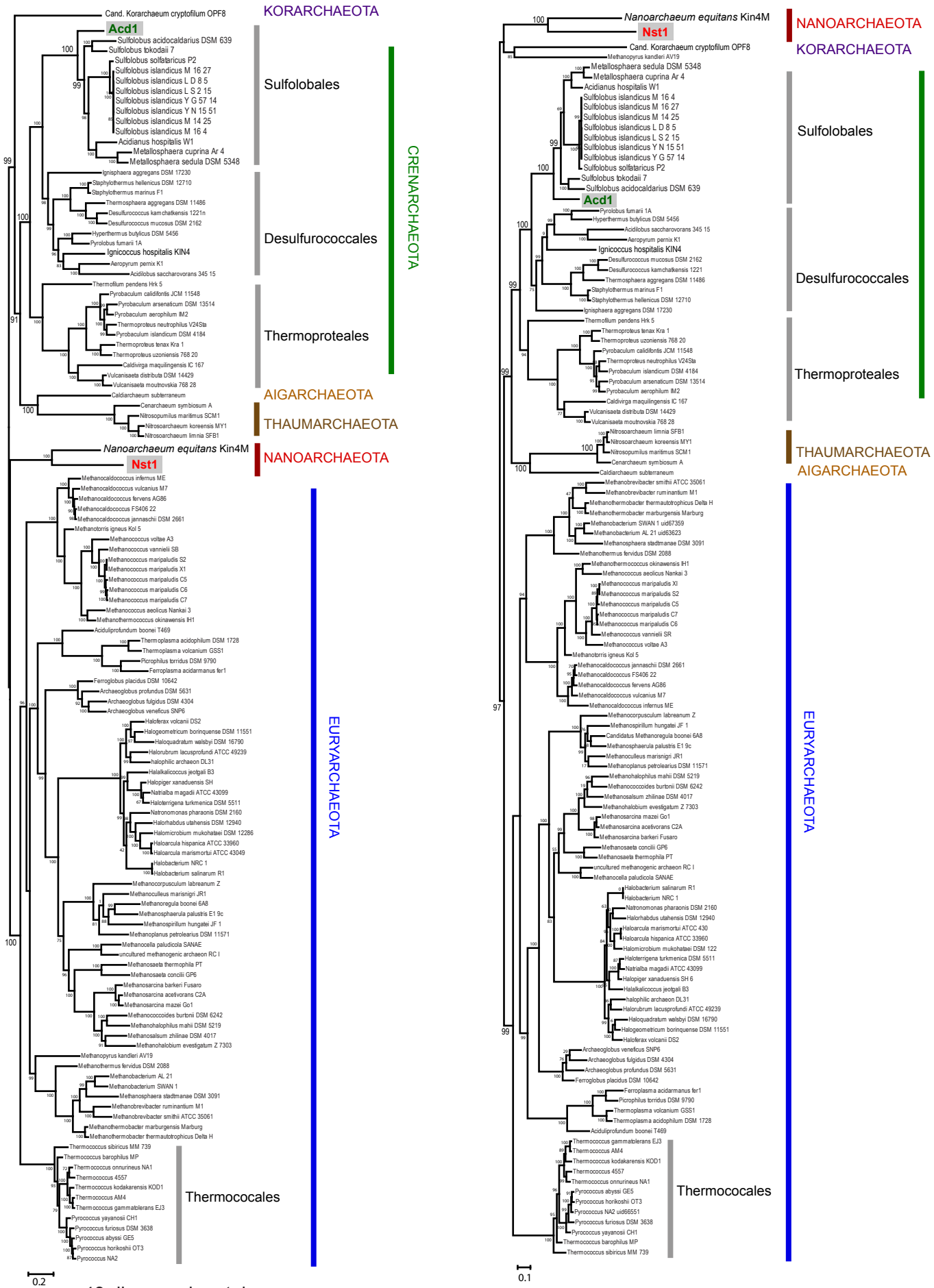

Additional file 3. Maximum likelihood phylogenies of the *Archaea* based on concatenated ribosomal protein gene sequences and RNA polymerase subunits.