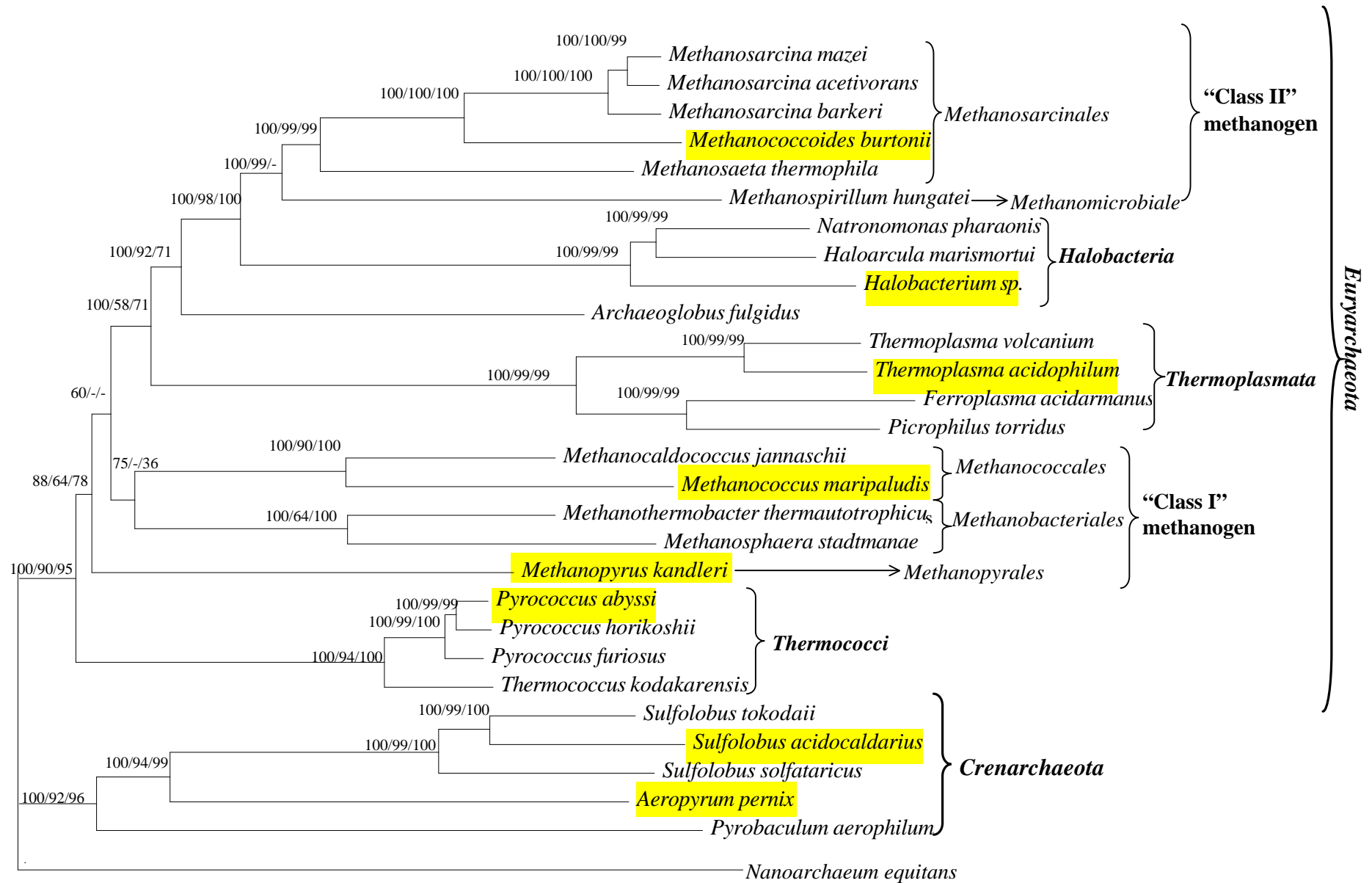


Phylogenetic Tree for Archaeal Species based on 51 conserved proteins

Gao, B. and Gupta, R.S. BMC Genomics (2007) 8, 86.



The tree shown is a neighbour-joining distance tree. The numbers on the nodes indicate bootstrap scores in NJ/ML/MP analyses.