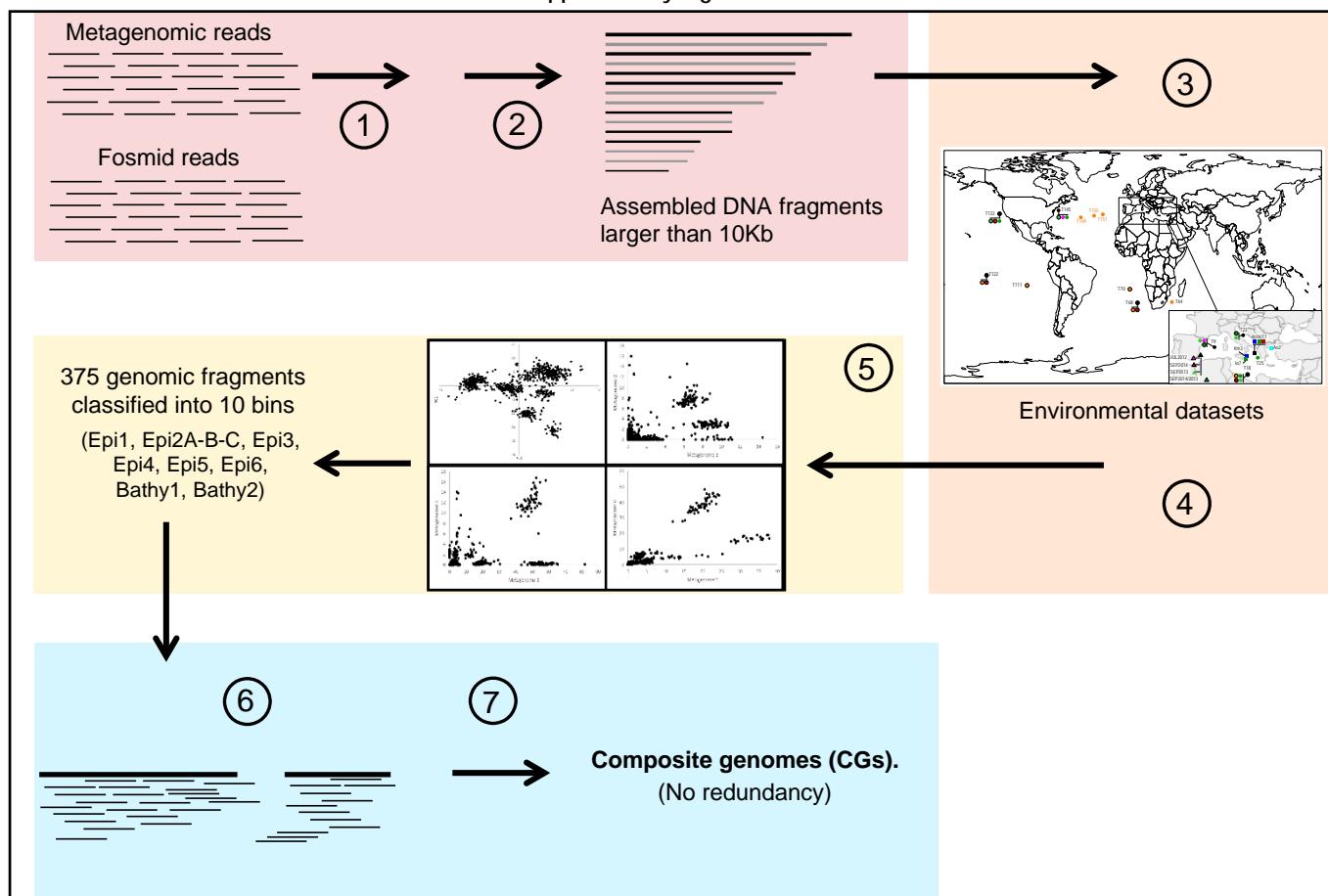


Supplementary Figure S1



Assembly and annotation of euryarchaeal sequences

- Assembly.** Metagenomic reads from MedDCM-JUL2012, MedDCM-SEP2014, Med-Io7-77mDCM, Med-Ae2-600mDeep and the fosmid reads from MedDCM-OCT2007, KM3 and AD1000 were independent and systematically assembled using IDBA_UD.
- Annotation.** Assembled DNA fragments were annotated using the NCBI nr-database, Pfam, COGs, arCOGs and TIGRFam. Only those fragments which have more than 50% of their ORFs annotated as Euryarchaeota were taken for further analysis (in black).

Retrieving of euryarchaeal sequences

- Recruitment.** Search of Euryarchaeota sequences within different marine databases. Only those metagenomes where our initial genomic fragments recruited more than 3 rpkg were selected.
- Assembly and annotation** of the selected metagenomes (as described previously). DNA fragments classified as euryarchaeota were collected.

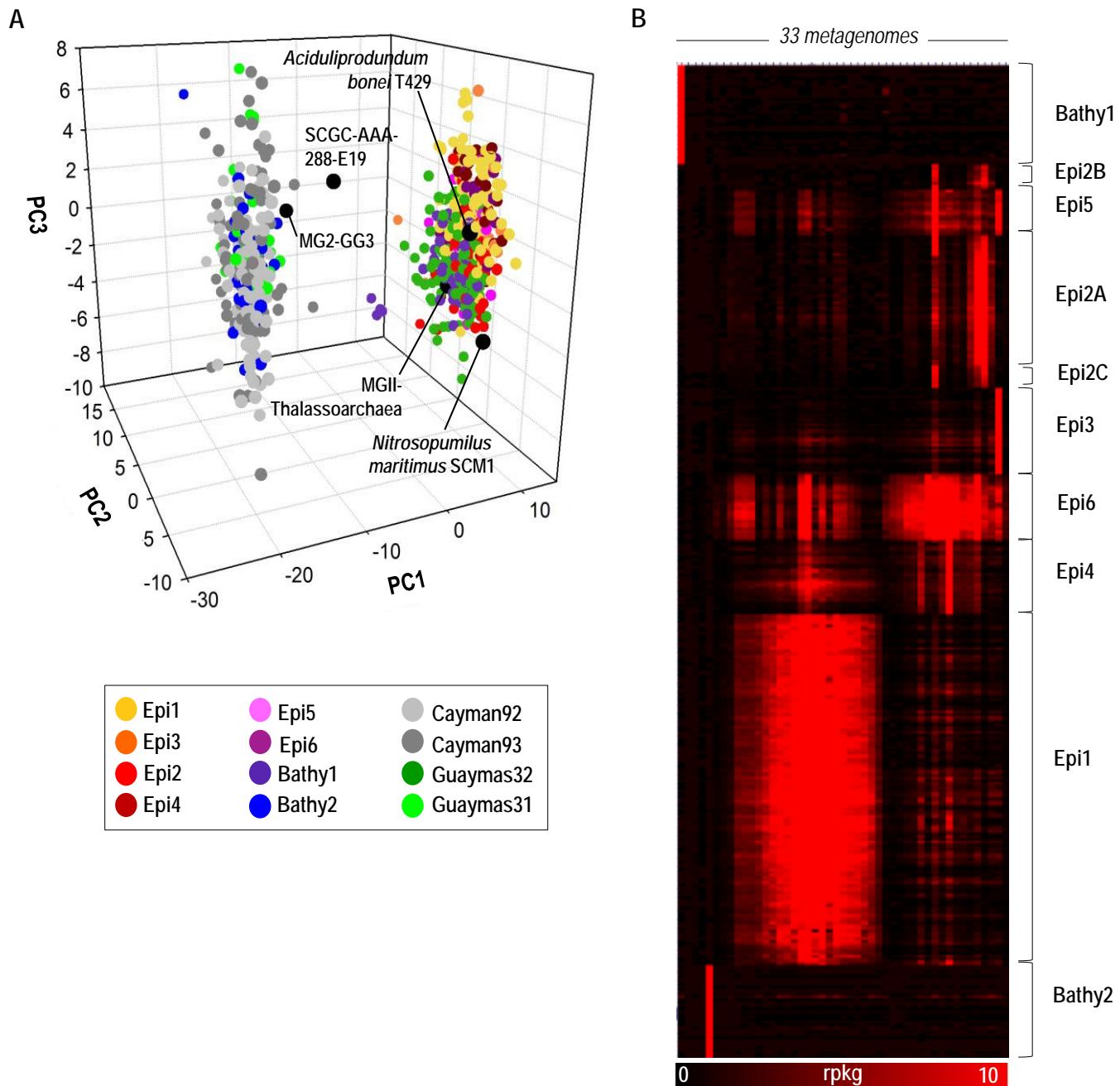
Binning of MG-III by tetranucleotide frequencies, %GC and differential coverage

- Binning** of the MG-III sequences by tetranucleotide frequencies, %GC and differential coverage among 33 metagenomes.

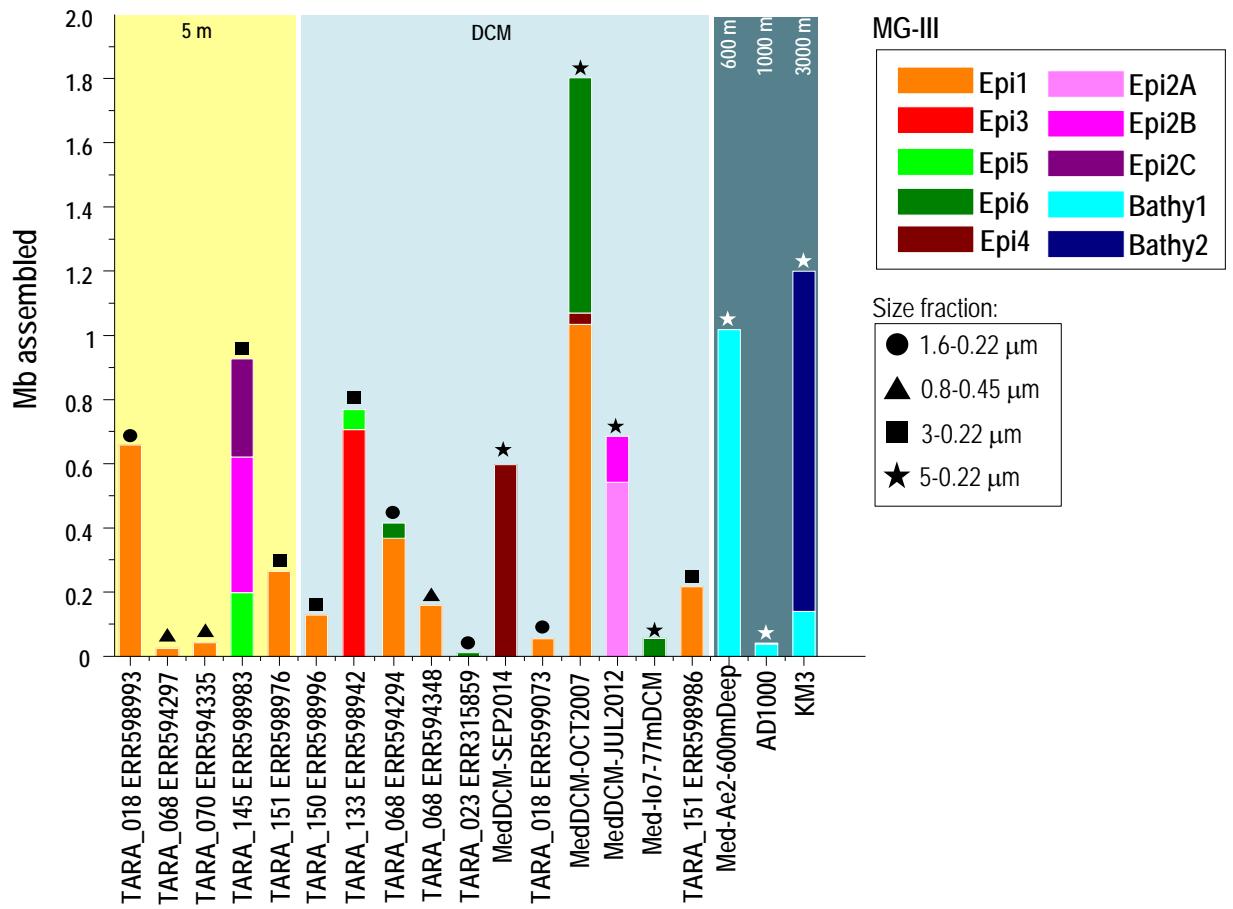
Re-assembly

- Alignment** of the sequences >10 Kb of one specific bin against the metagenomic reads where those contigs came from using BWA. The process was made for all the bins. The reads extracted were used in the following step.
- Assembly** of the reads with SPADES using the DNA fragments for scaffolding with the parameter –trusted_contigs. The re-assembly was done for all the bins and their products are called «composite genome» (CG).

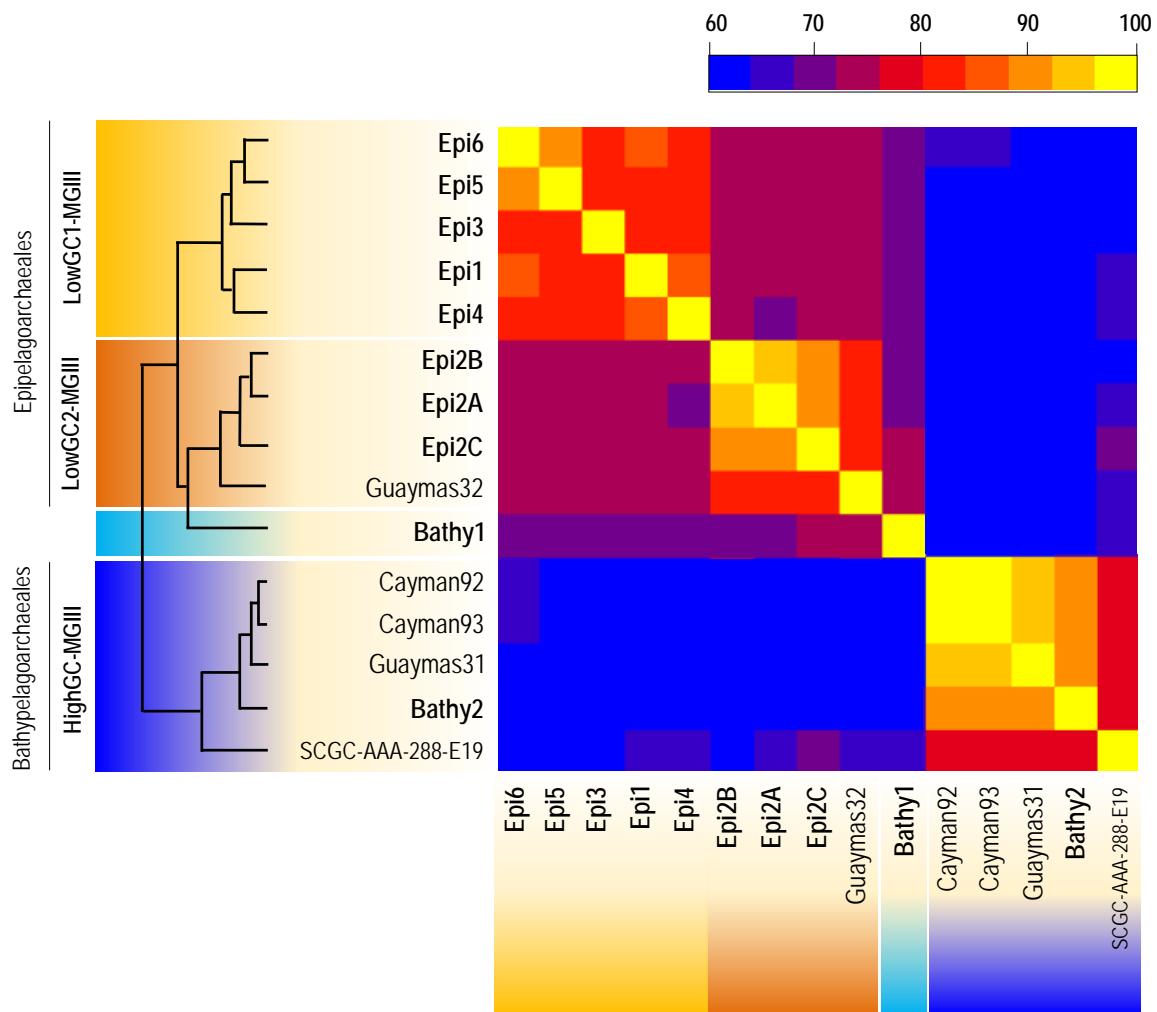
Supplementary Figure S2

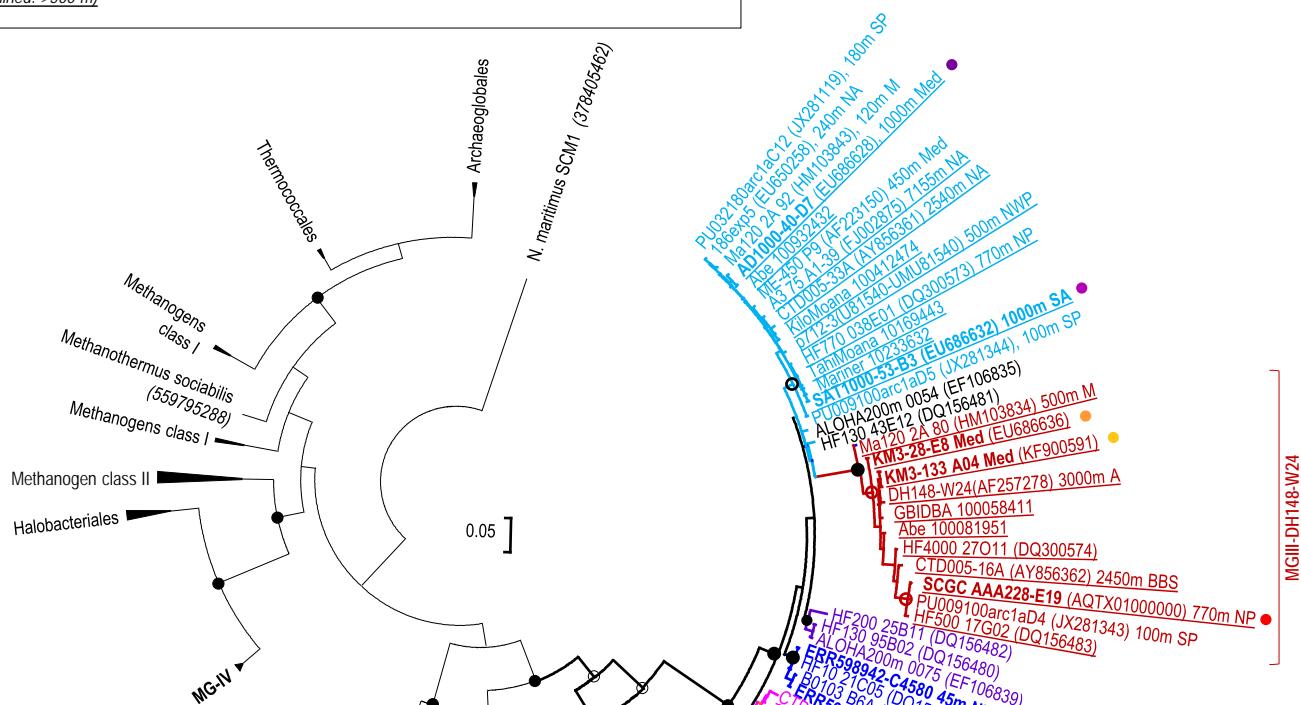
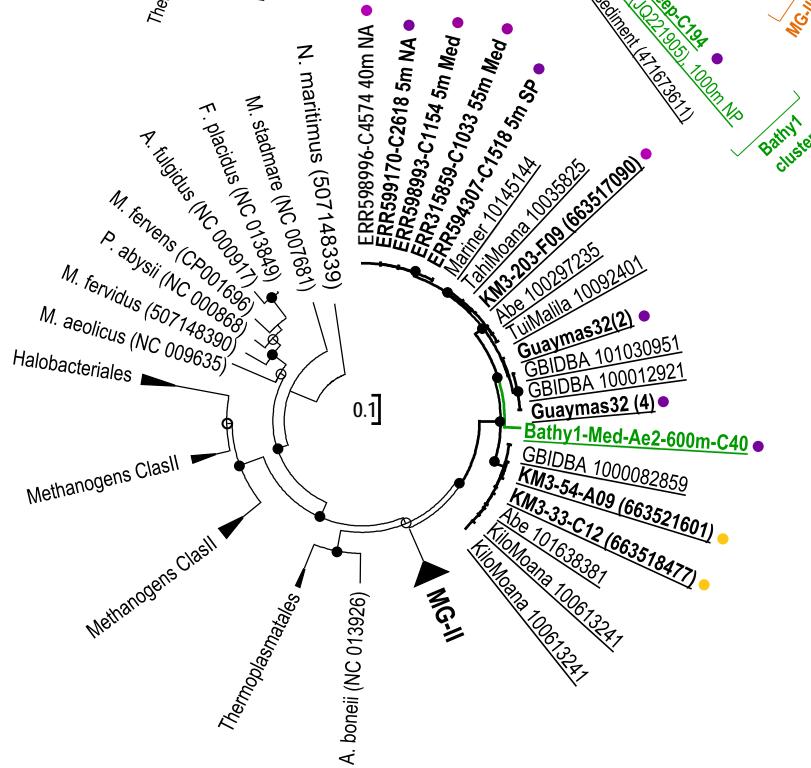


Supplementary Figure S3

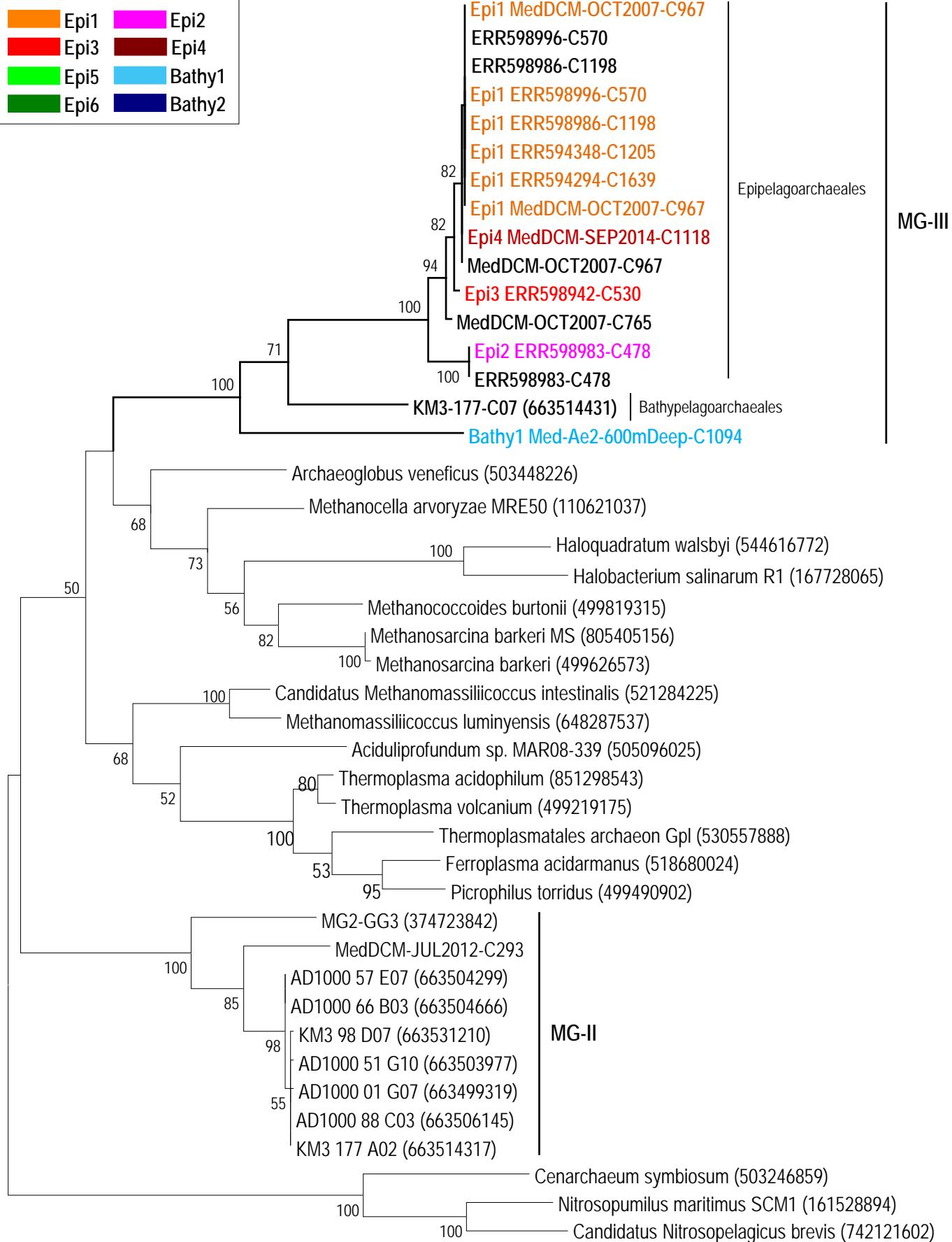


Supplementary Figure S4

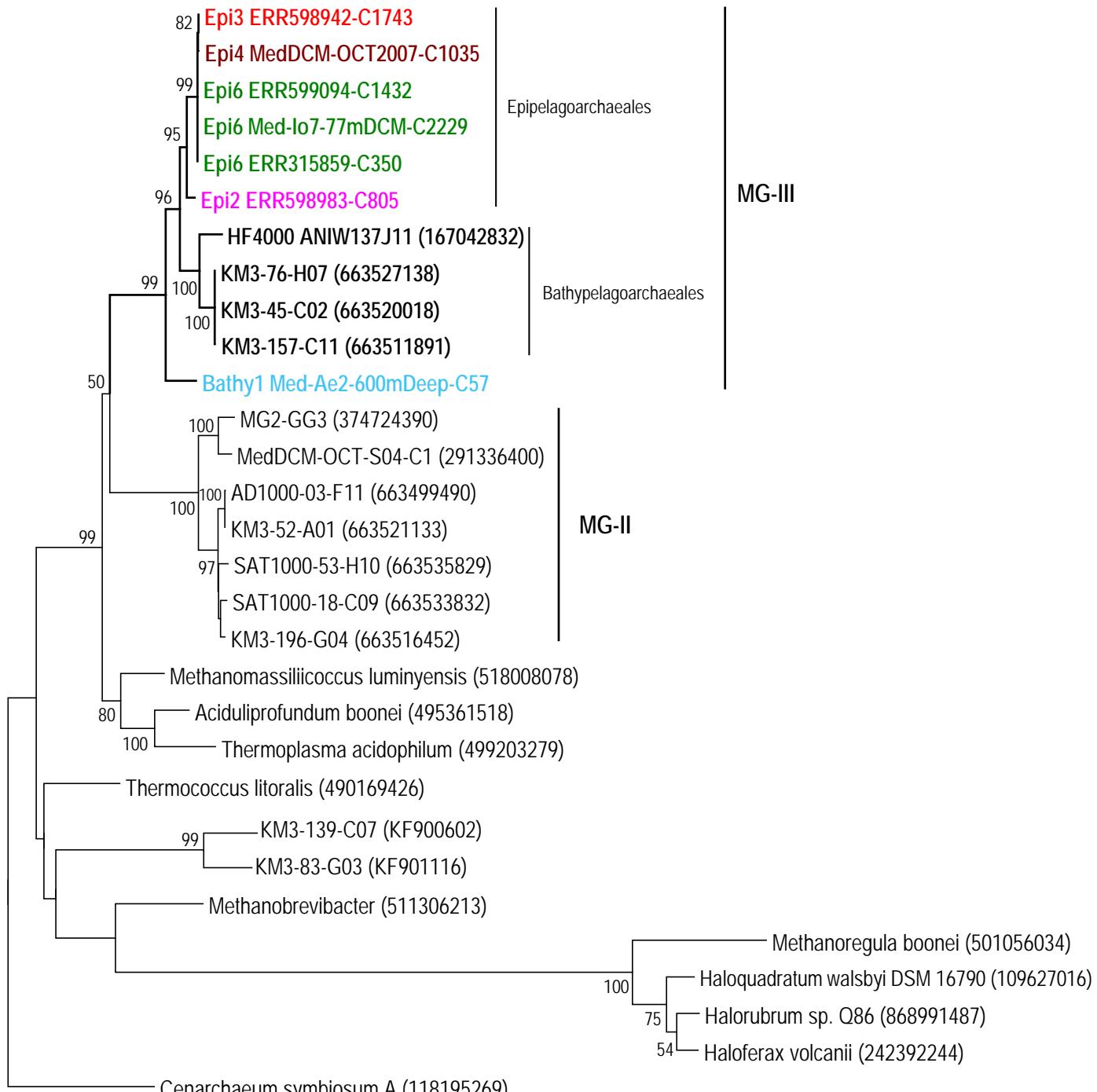


**A****B**

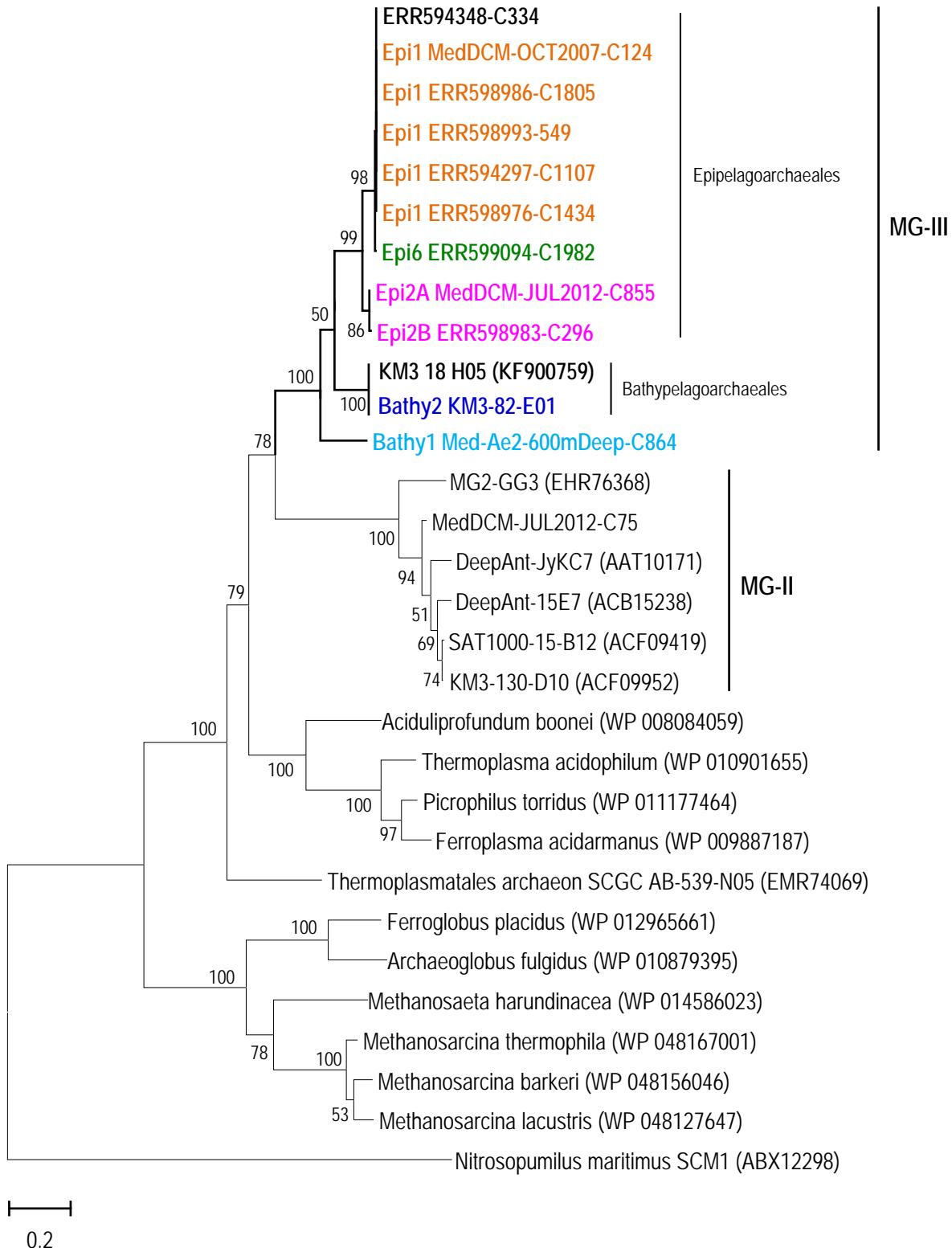
Supplementary Figure S6



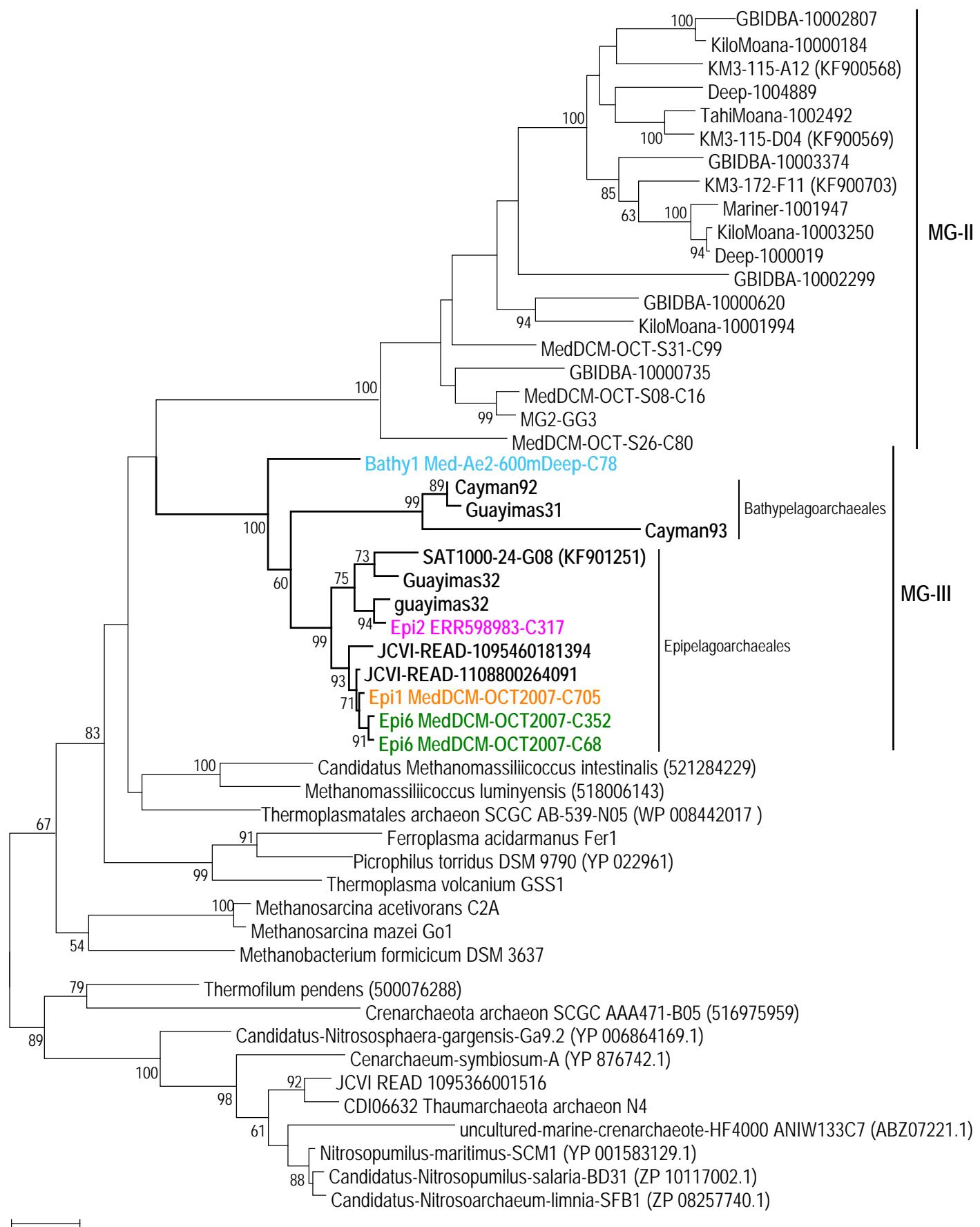
Supplementary Figure S7



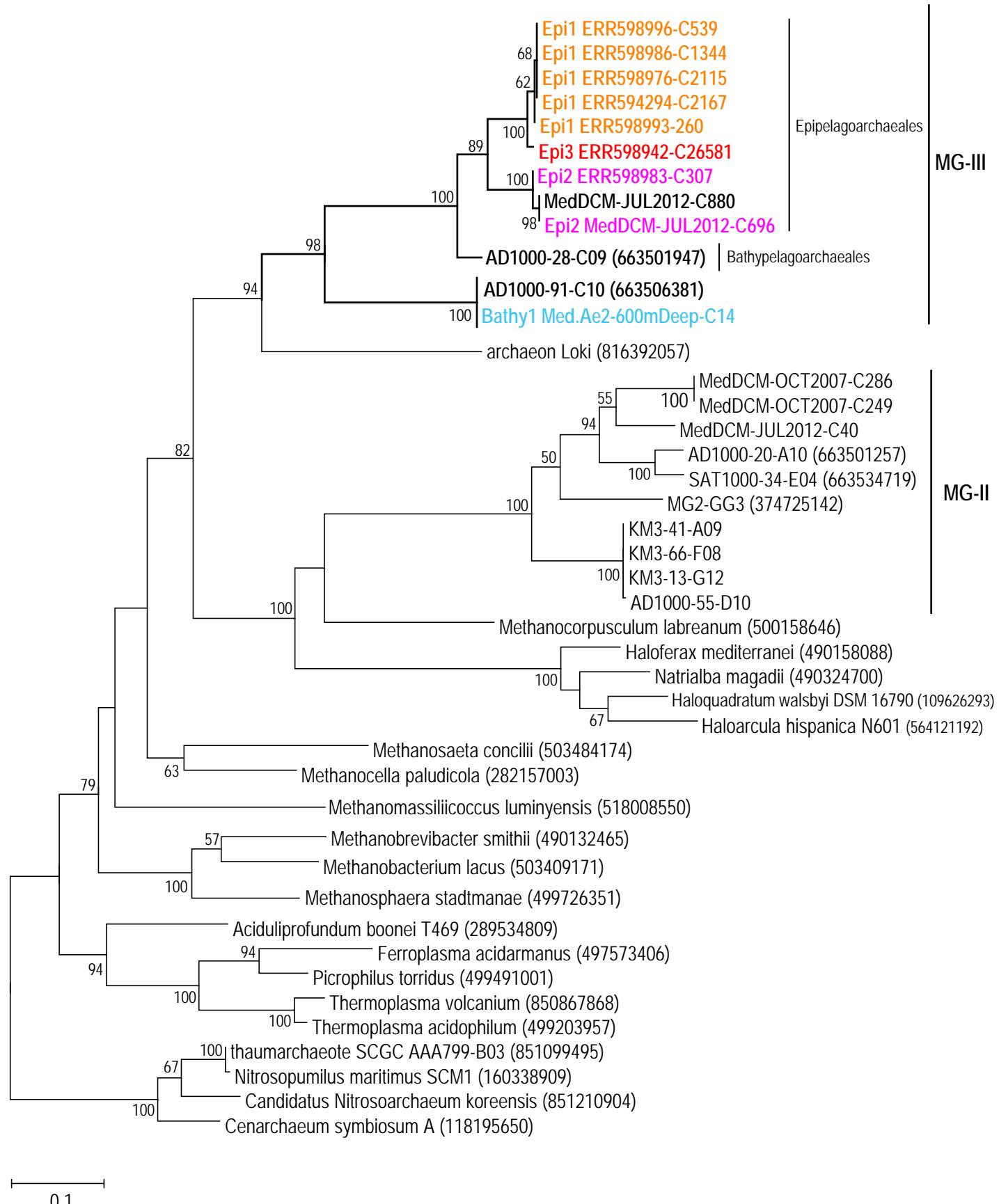
Supplementary Figure S8



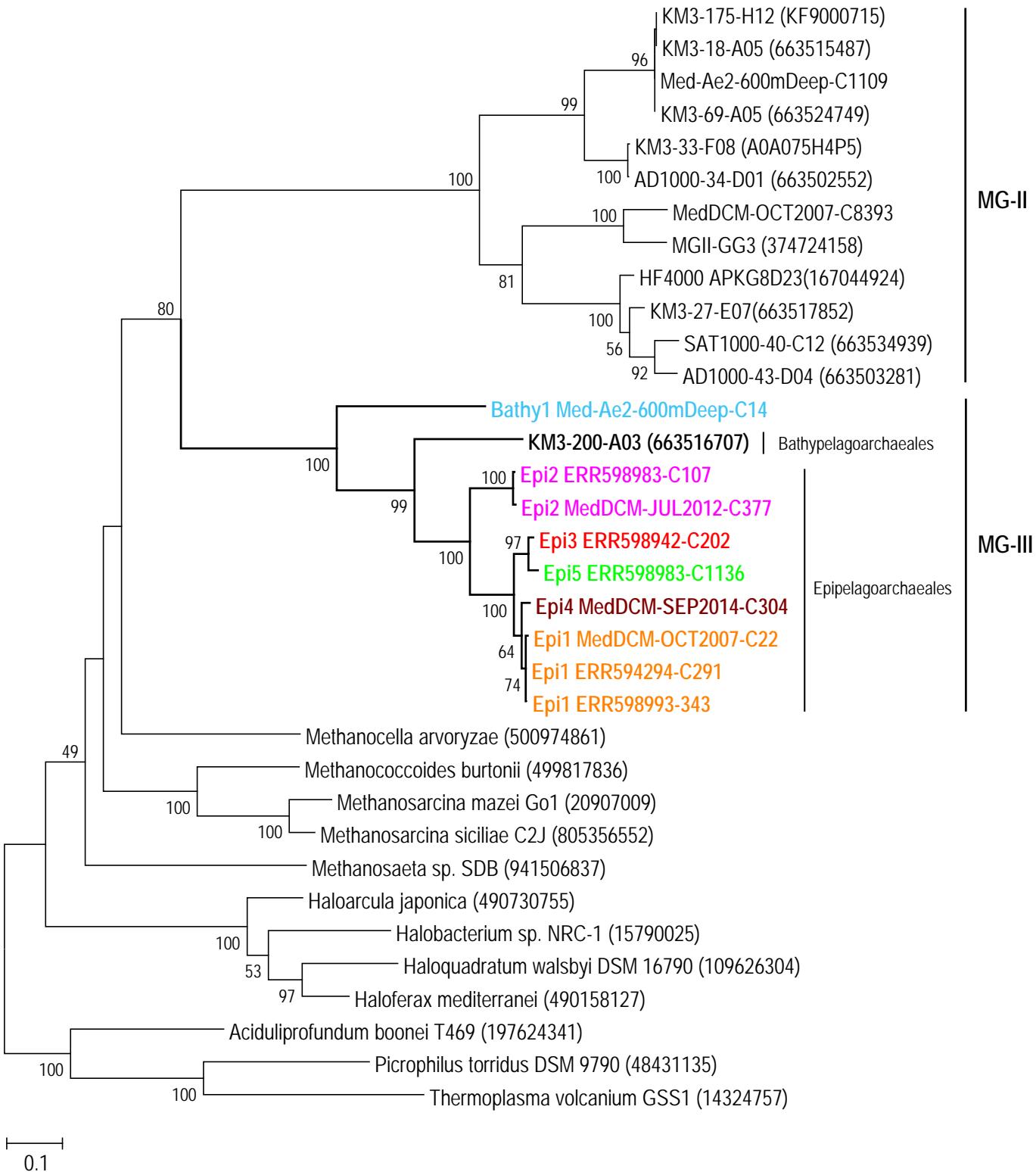
Supplementary Figure S9



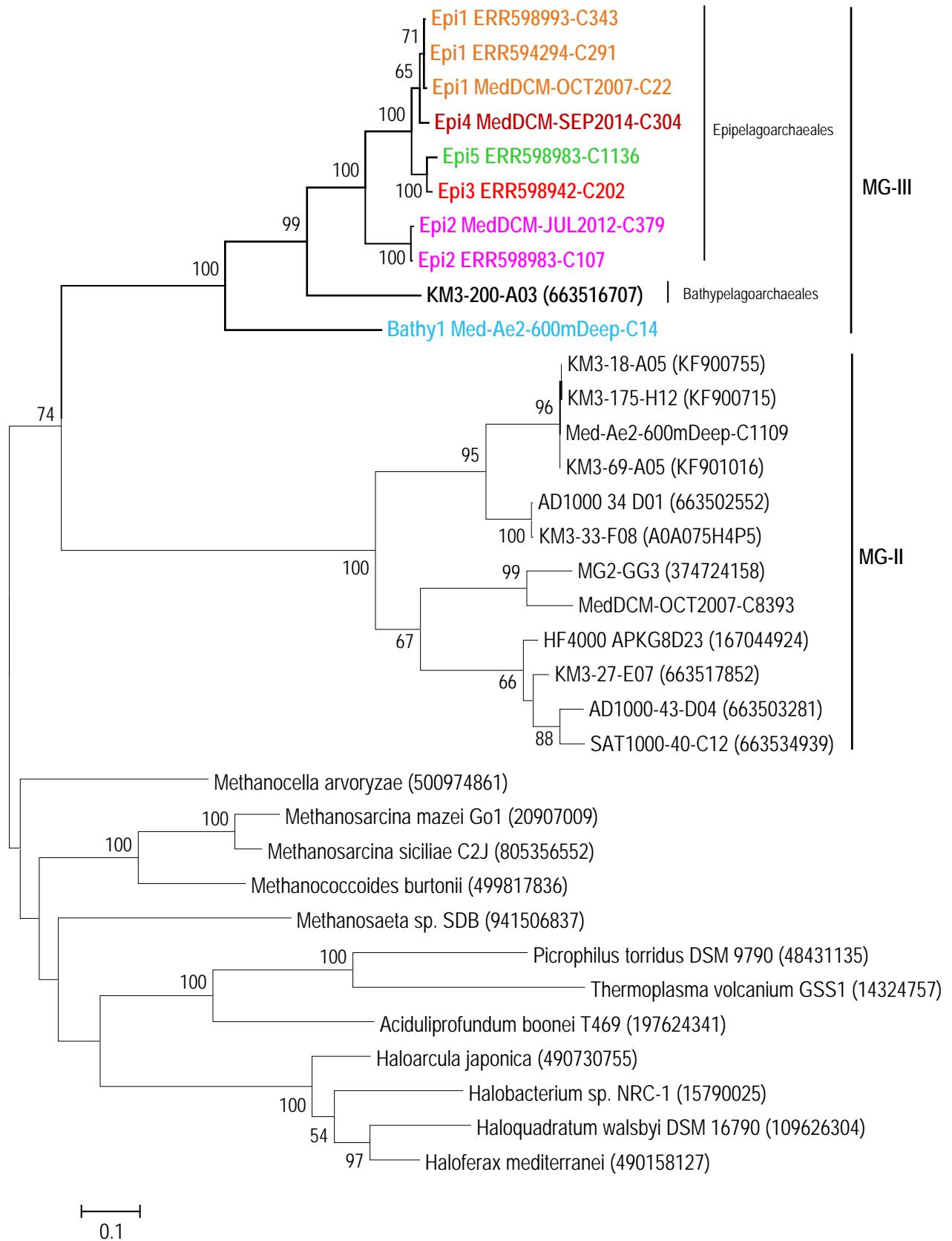
Supplementary Figure S10

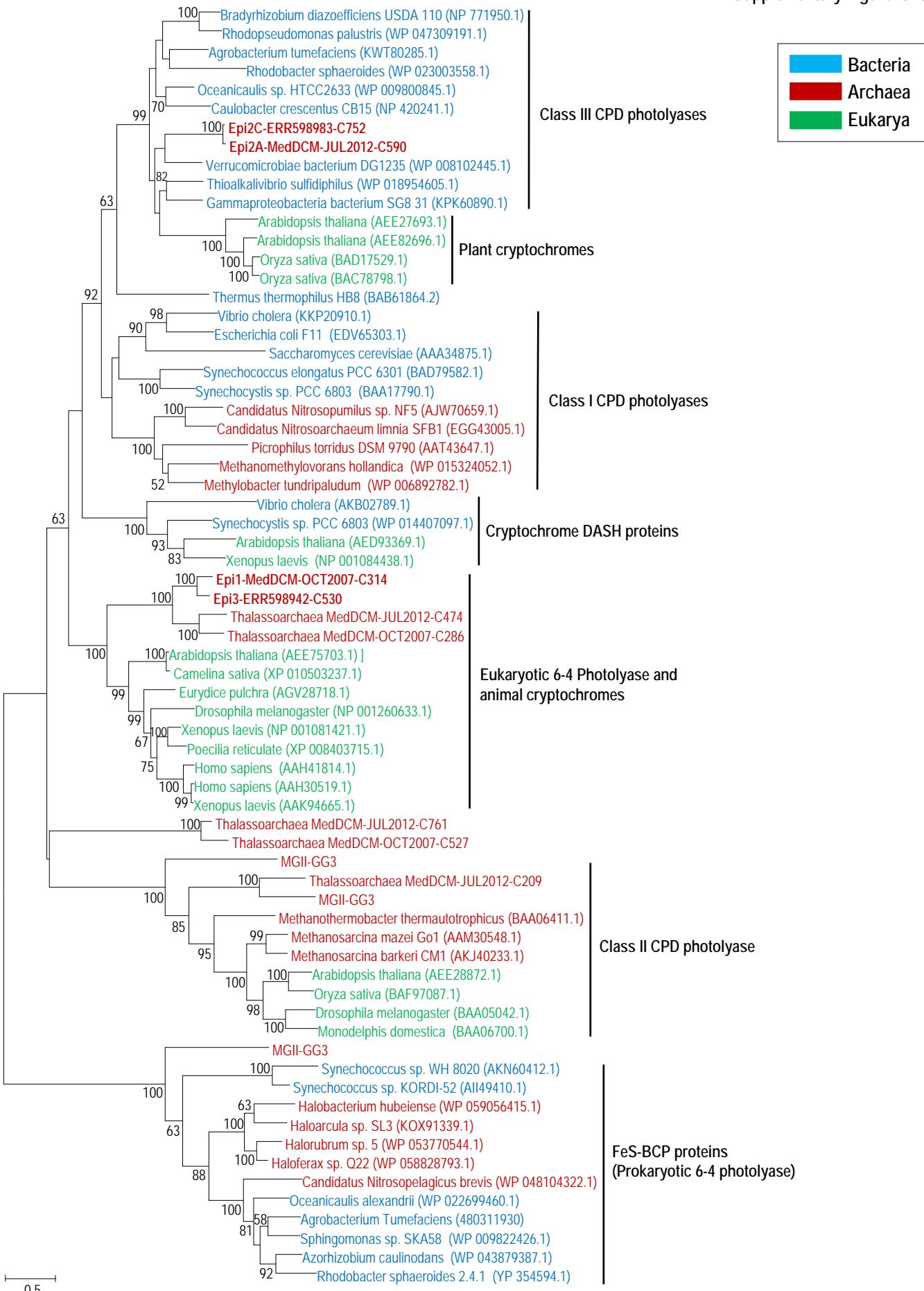


Supplementary Figure S11

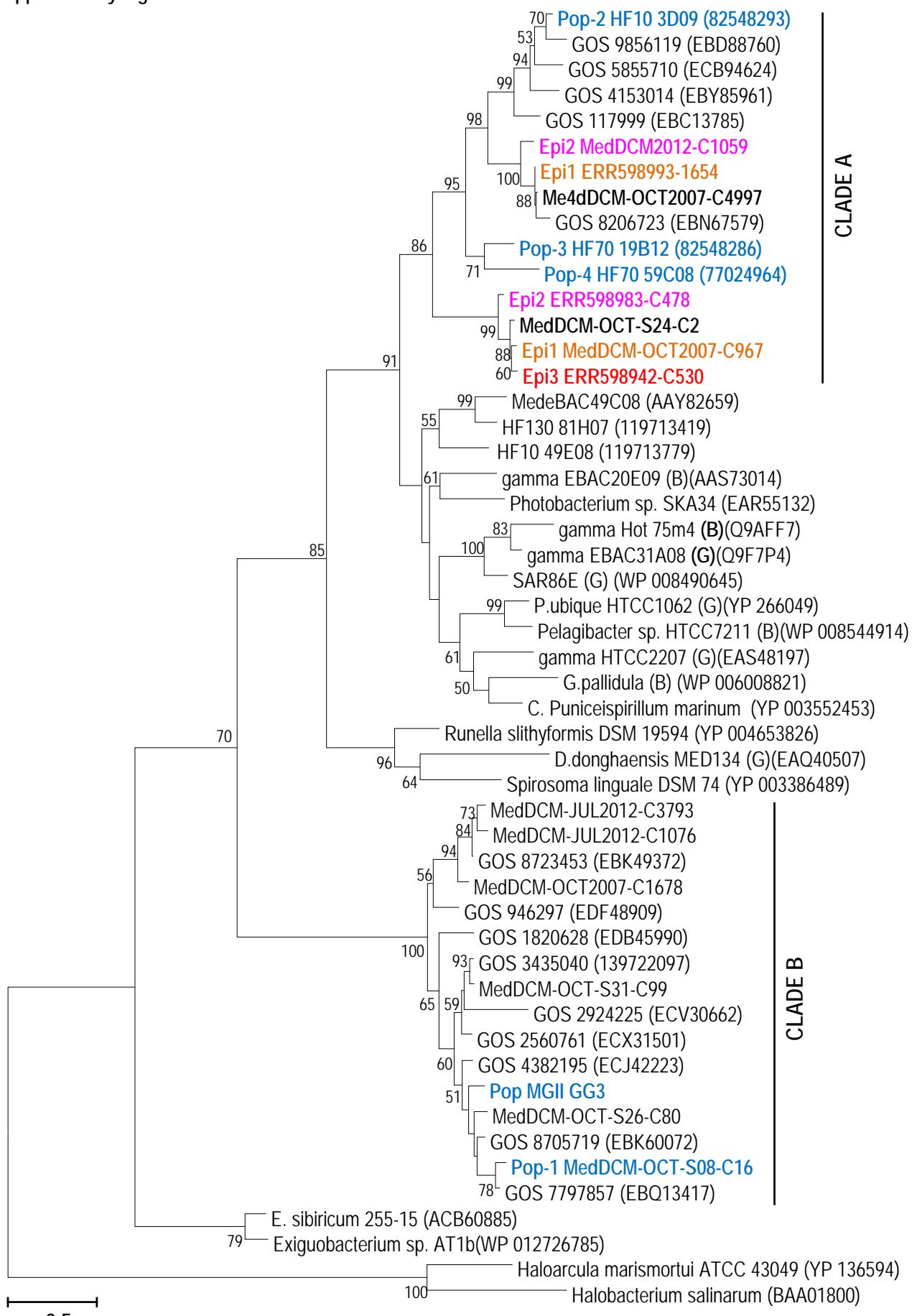


Supplementary Figure S12

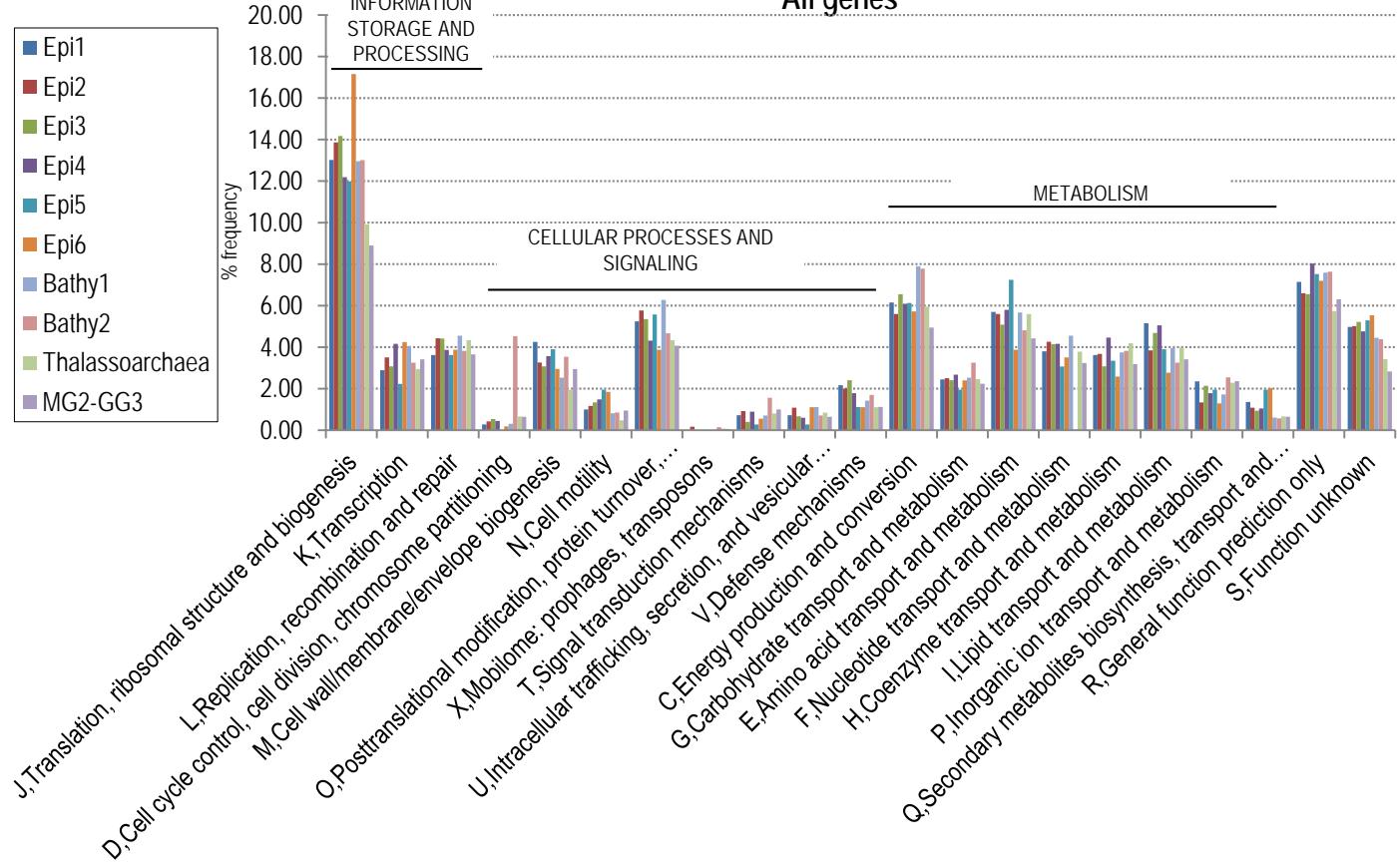
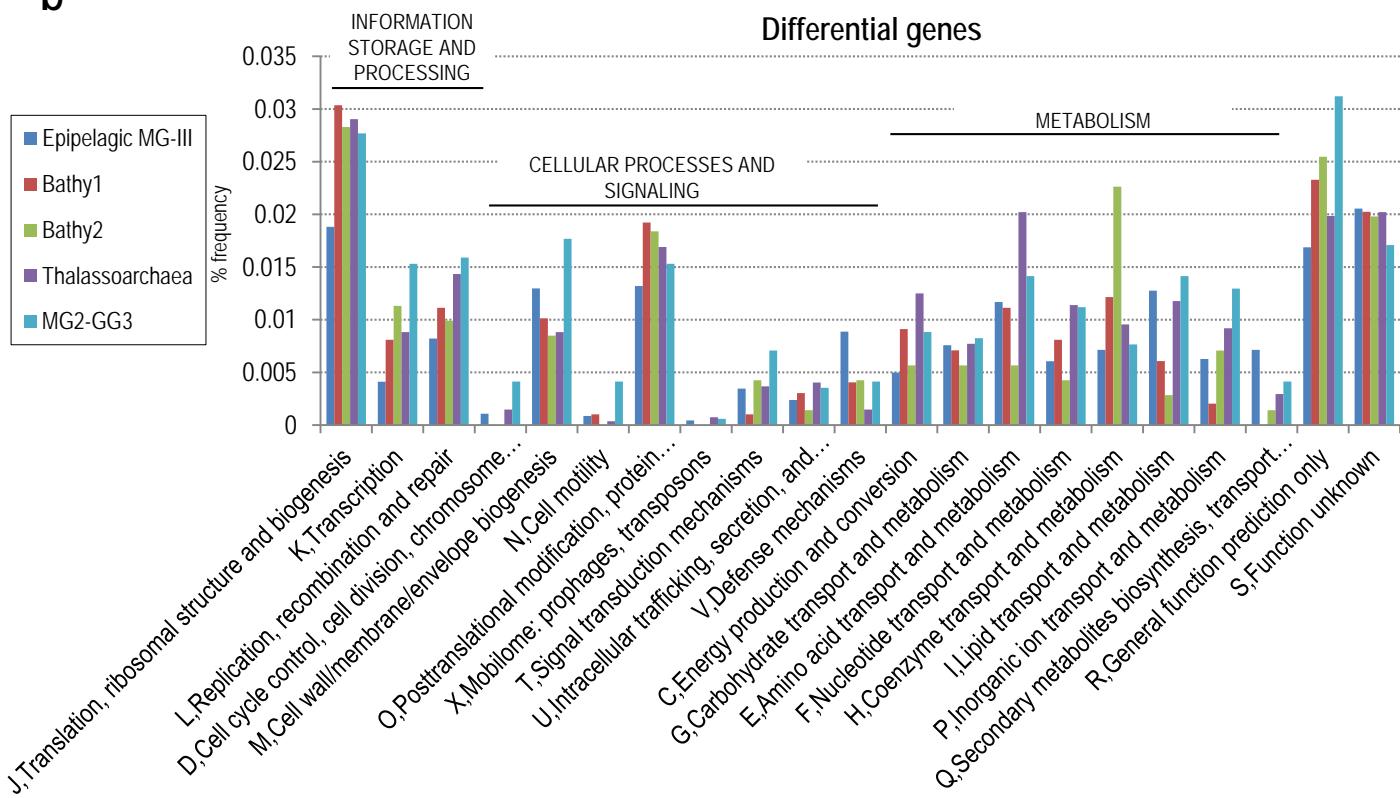




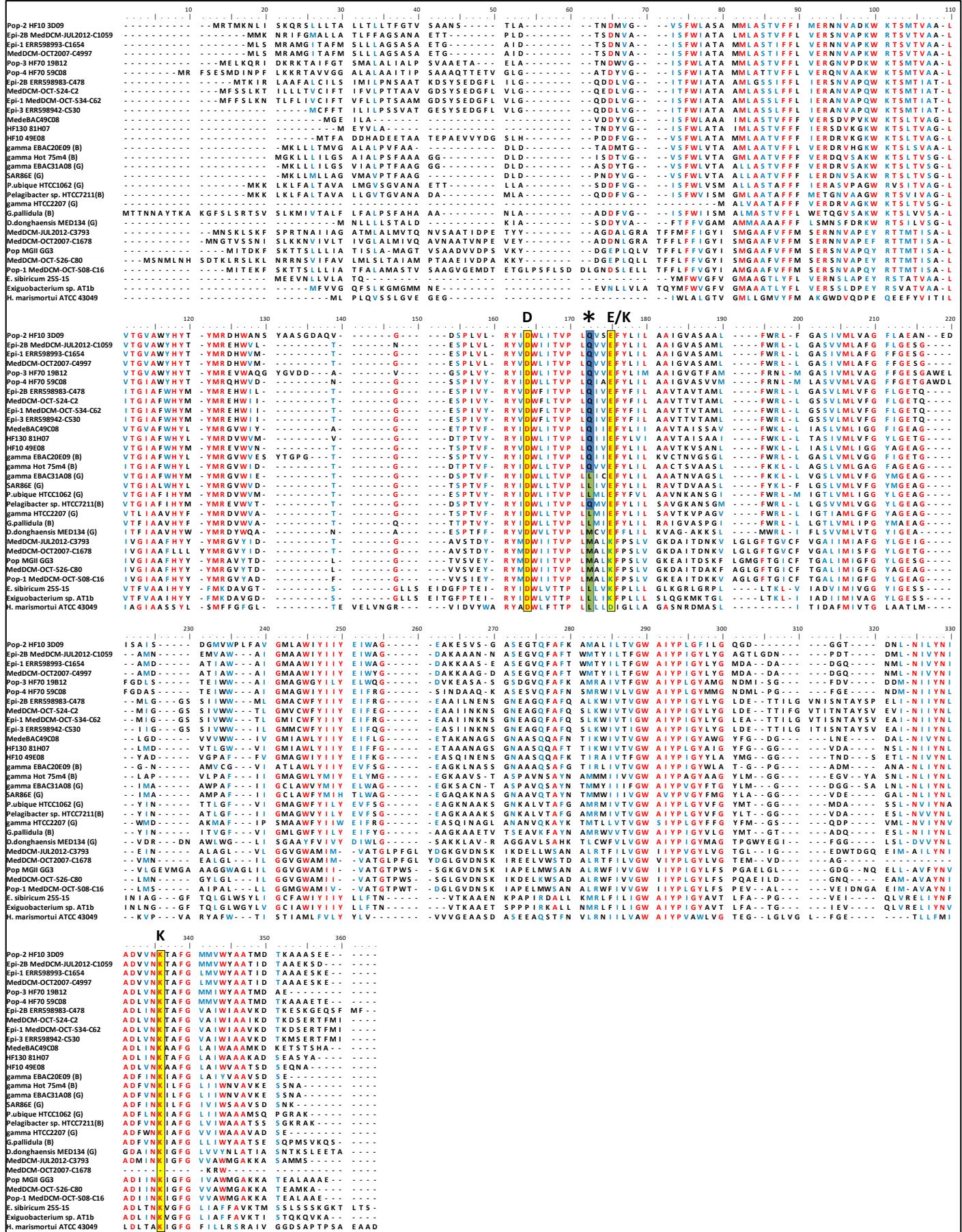
Supplementary Figure S14



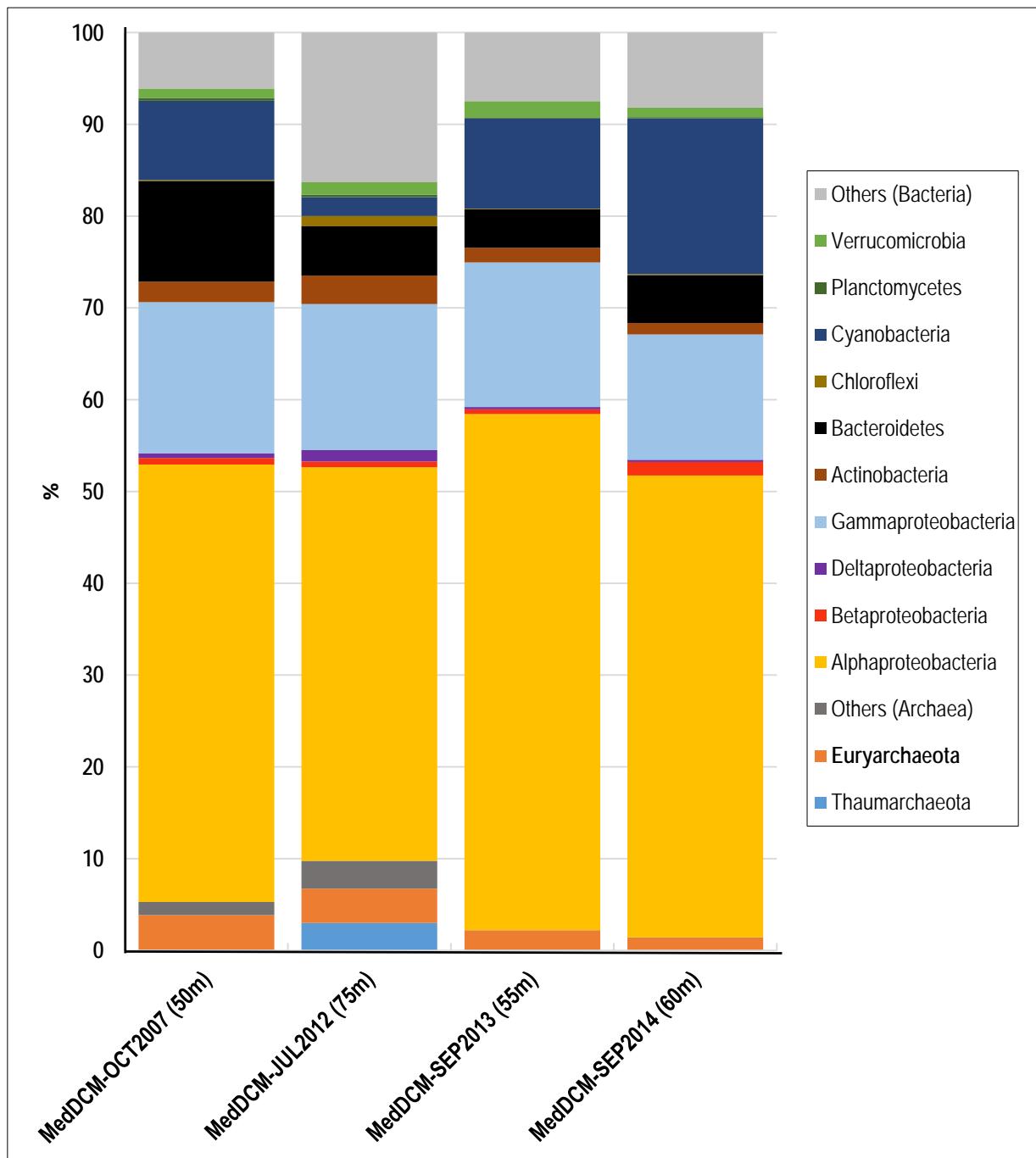
Supplementary Figure S15

a**b**

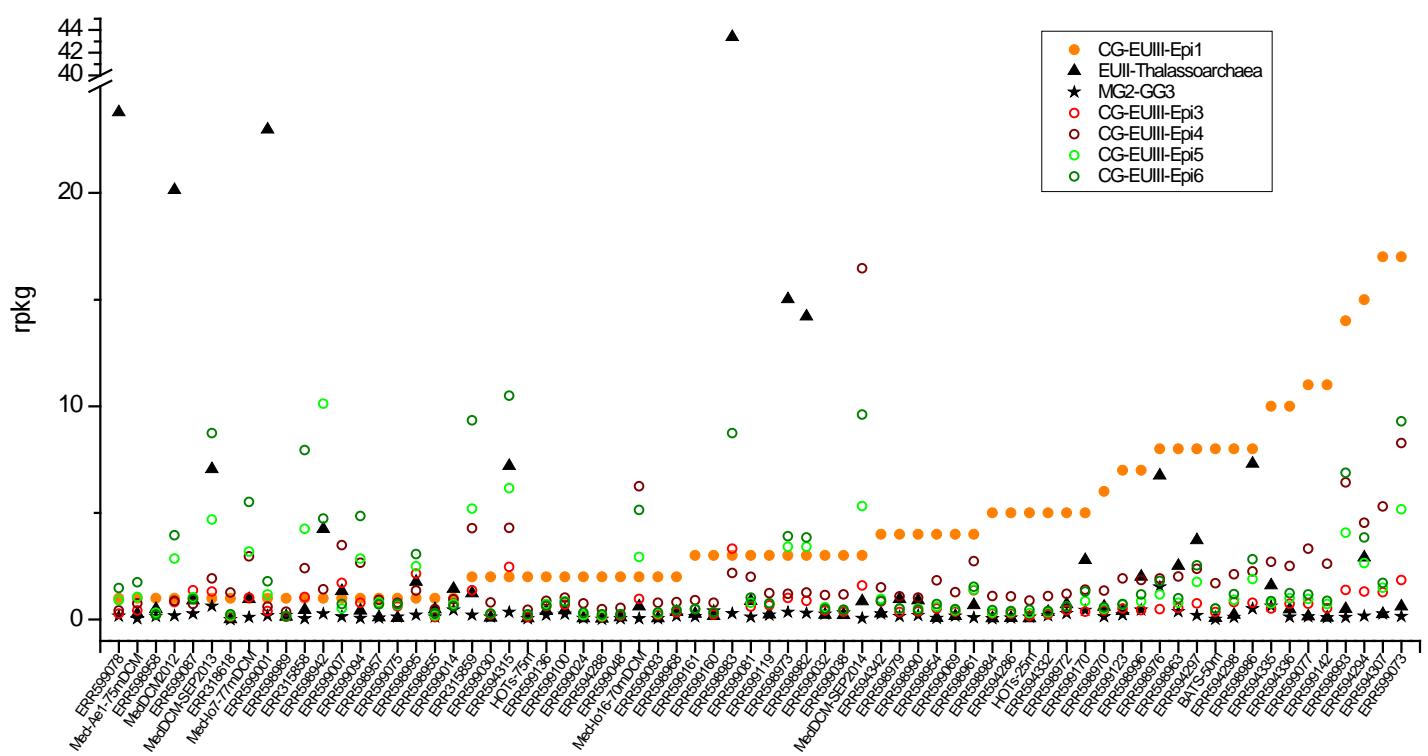
Supplementary Figure S16



Supplementary Figure S17



Supplementary Figure S18



Supplementary Figure S19

