

Supplementary Materials

# Rooted tRNAomes and Evolution of the Genetic Code

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## Supplemental Figures

Figure S1. An evolutionary tree of the *Pyrococcus furiosus* DSM3638 (archaea) DNA tRNA<sup>ome</sup>. Trees are rooted to tRNA<sup>Pri</sup>. An error was corrected in tRNA<sup>Arg</sup> (anticodon GCG), in which the 5'- and 3'-acceptor stems did not fully pair.

Figure S2. An evolutionary tree of the *Pyrococcus abyssi* GE5 (archaea) tRNA<sup>ome</sup>.

Figure S3. An evolutionary tree of the *Pyrococcus horikoshii* OT3 (archaea) tRNA<sup>ome</sup>.

Figure S4. An evolutionary tree of the *Staphylothermus marinus* F1 (archaea) tRNA<sup>ome</sup>.

Figure S5. An evolutionary tree of the *Pyrobaculum aerophilium* str. IM2 (archaea) tRNA<sup>ome</sup>.

Figure S6. An evolutionary tree of the *Sulfolobus solfataricus* P2 (archaea) tRNA<sup>ome</sup>.

Figure S7. An evolutionary tree of the *Aeropyrum pernix* K1 (archaea) tRNA<sup>ome</sup>.

Figure S8. An evolutionary tree of the *Thermus thermophilus* HB27 (bacteria) tRNA<sup>ome</sup>.

Figure S9. Typical tRNA diagrams for *Pyrococcus furiosus*, *Thermus thermophilus* and *Escherichia coli*.

Diagrams were generated using the tRNA database.<sup>1</sup>

Figure S10. Modifications of tRNAs in archaea. Modifications are shown for *Haloferax volcanii* (archaea). In the *Pyrococcus* typical DNA diagram: cyan dots) anticodon bases; orange dots) loop bases 6-7 (loop bases 3-7 are tightly stacked as in a helix, filling the 7-mer loop); yellow dots) intercalation of D loop G19 lifts T loop base 5 to the position of base 7 in the anticodon loop, flipping bases 6 and 7 out of the T loop. U turns are indicated (red arrows). The discriminator base is indicated (cyan arrow).

## References:

1. Juhling F, Morl M, Hartmann RK, Sprinzl M, Stadler PF, Putz J. tRNADB 2009: compilation of tRNA sequences and tRNA genes. *Nucleic Acids Res* 2009; 37:D159-62.

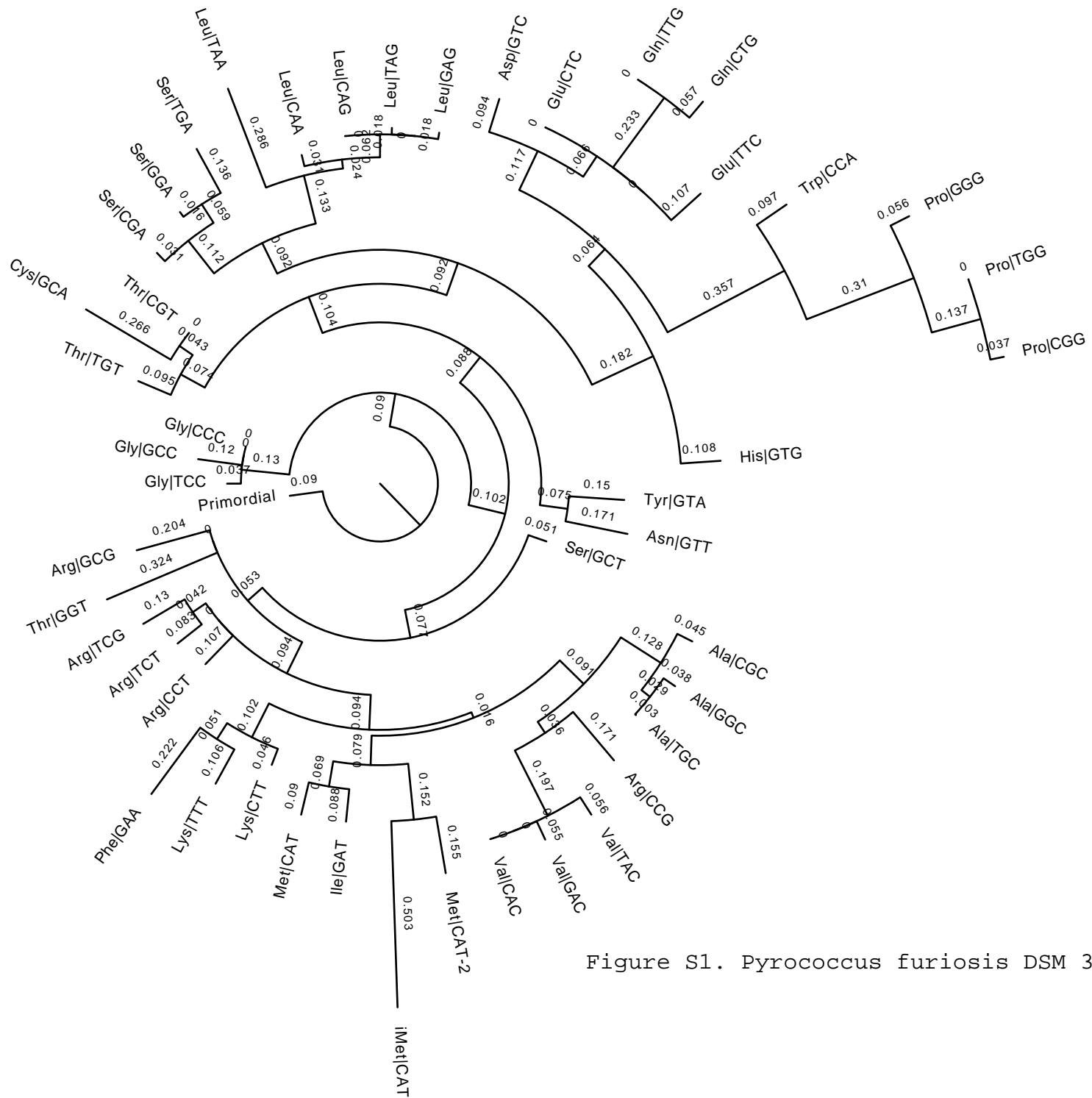


Figure S1. *Pyrococcus furiosus* DSM 3638.

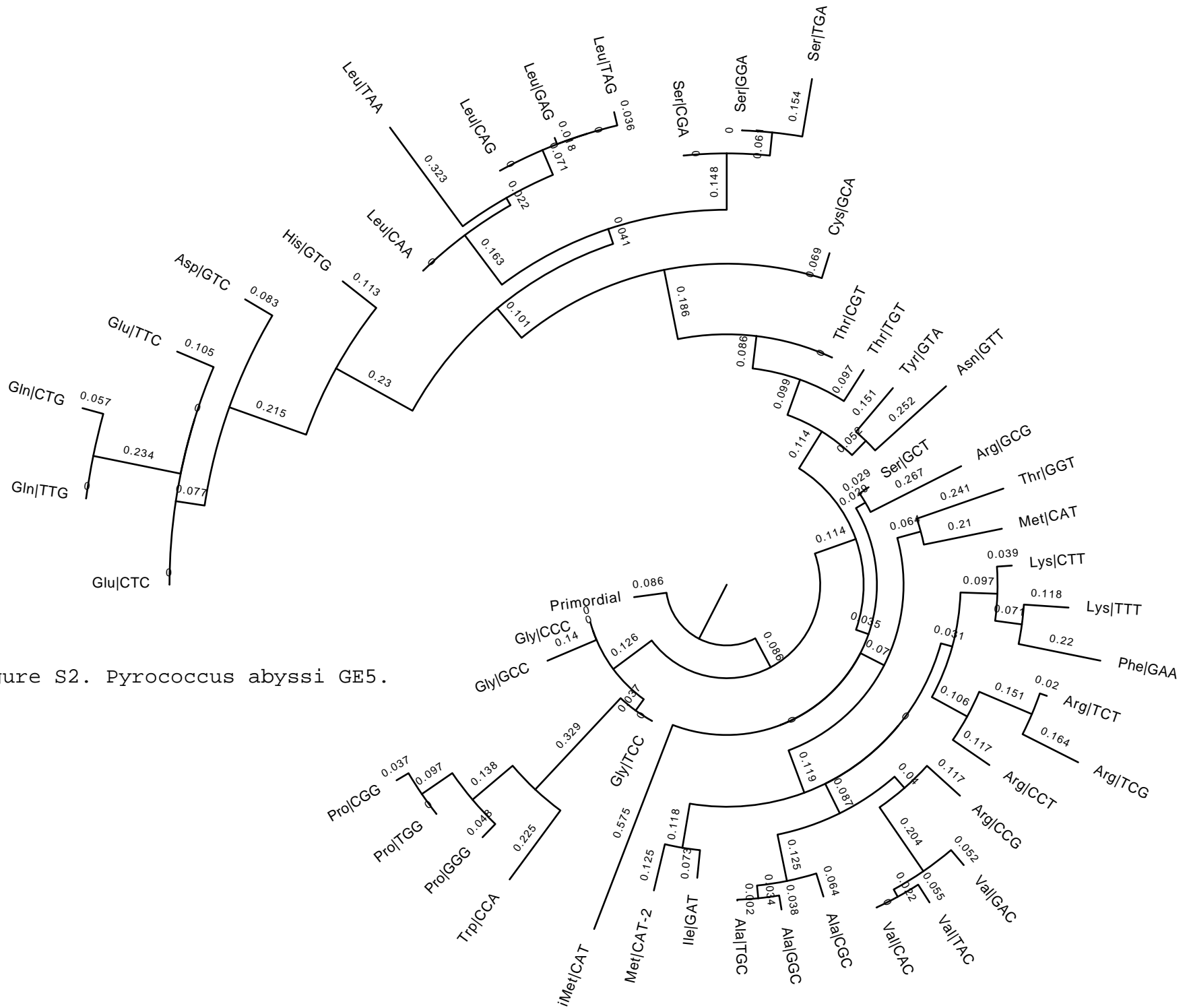


Figure S2. *Pyrococcus abyssi* GE5.

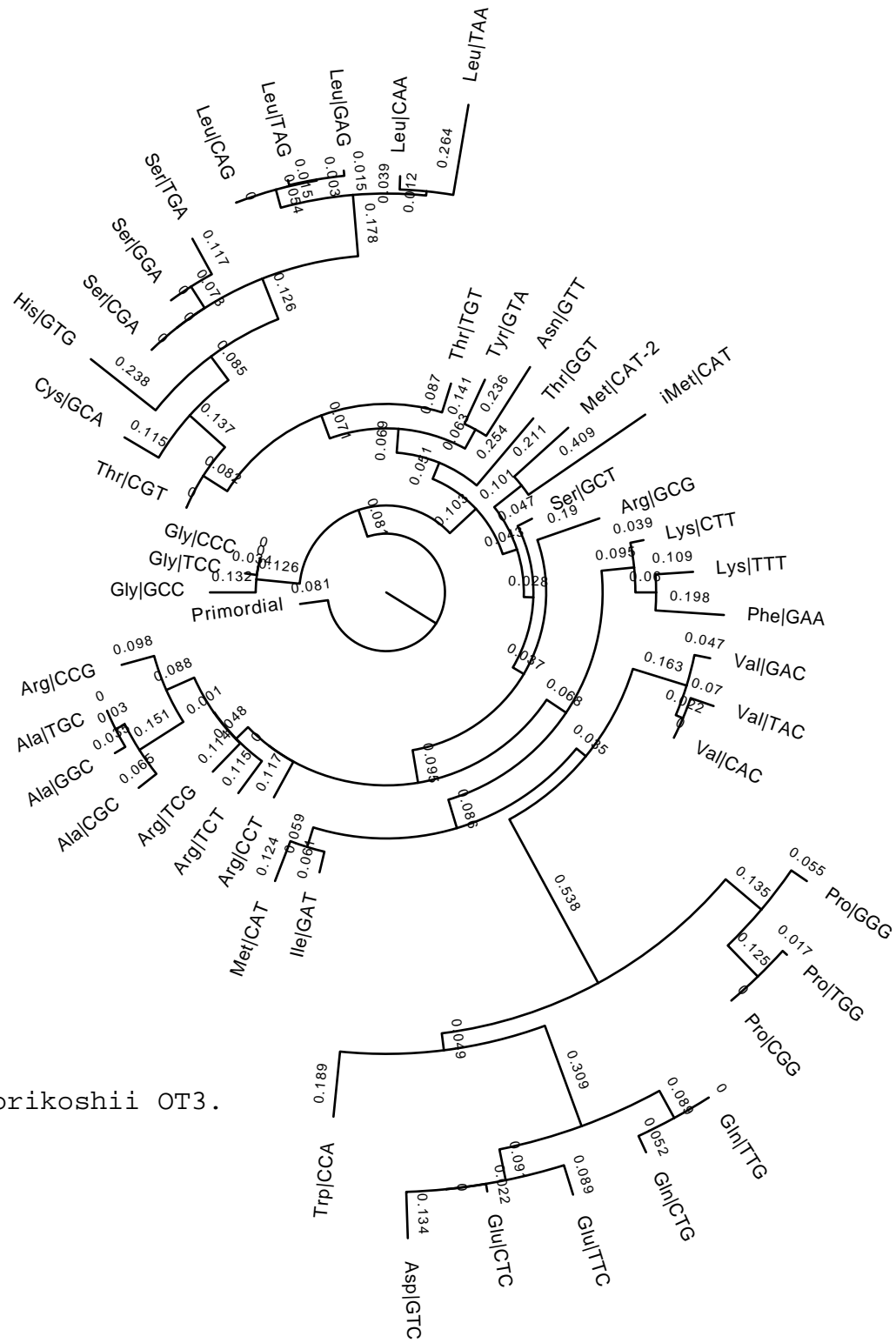


Figure S3. *Pyrococcus horikoshii* OT3.

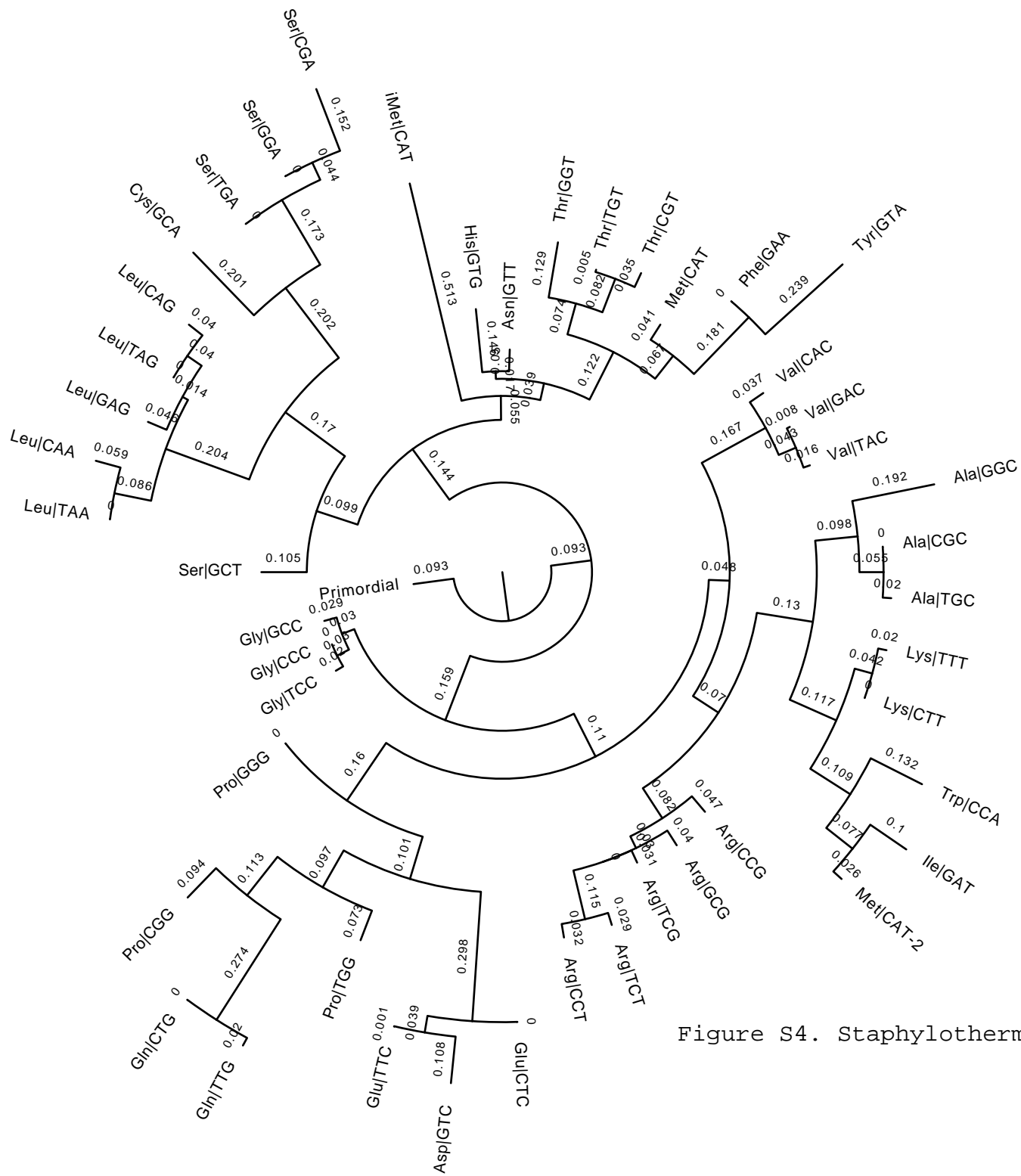


Figure S4. *Staphylothermus marinus* F1.

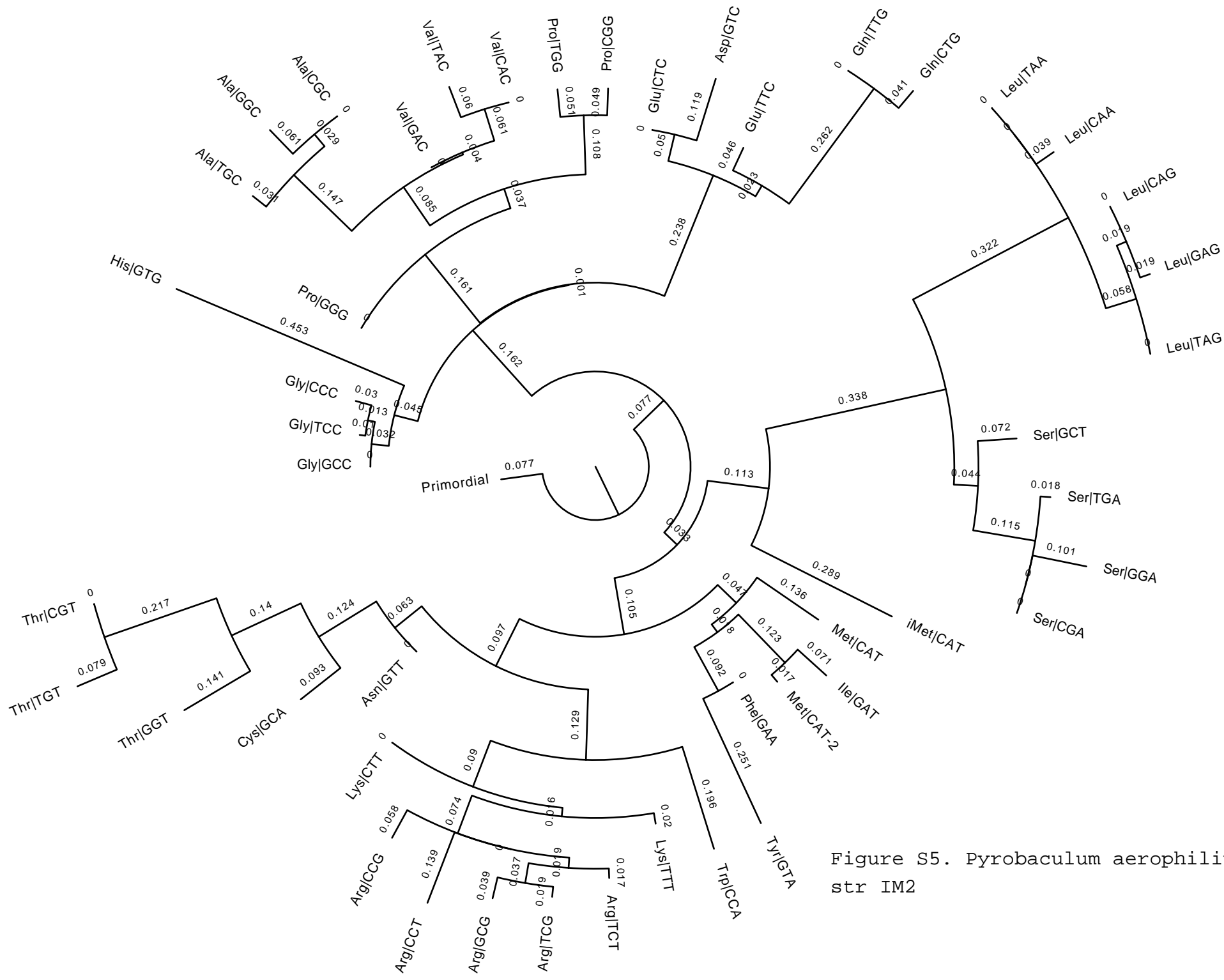


Figure S5. *Pyrobaculum aerophilum* str IM2

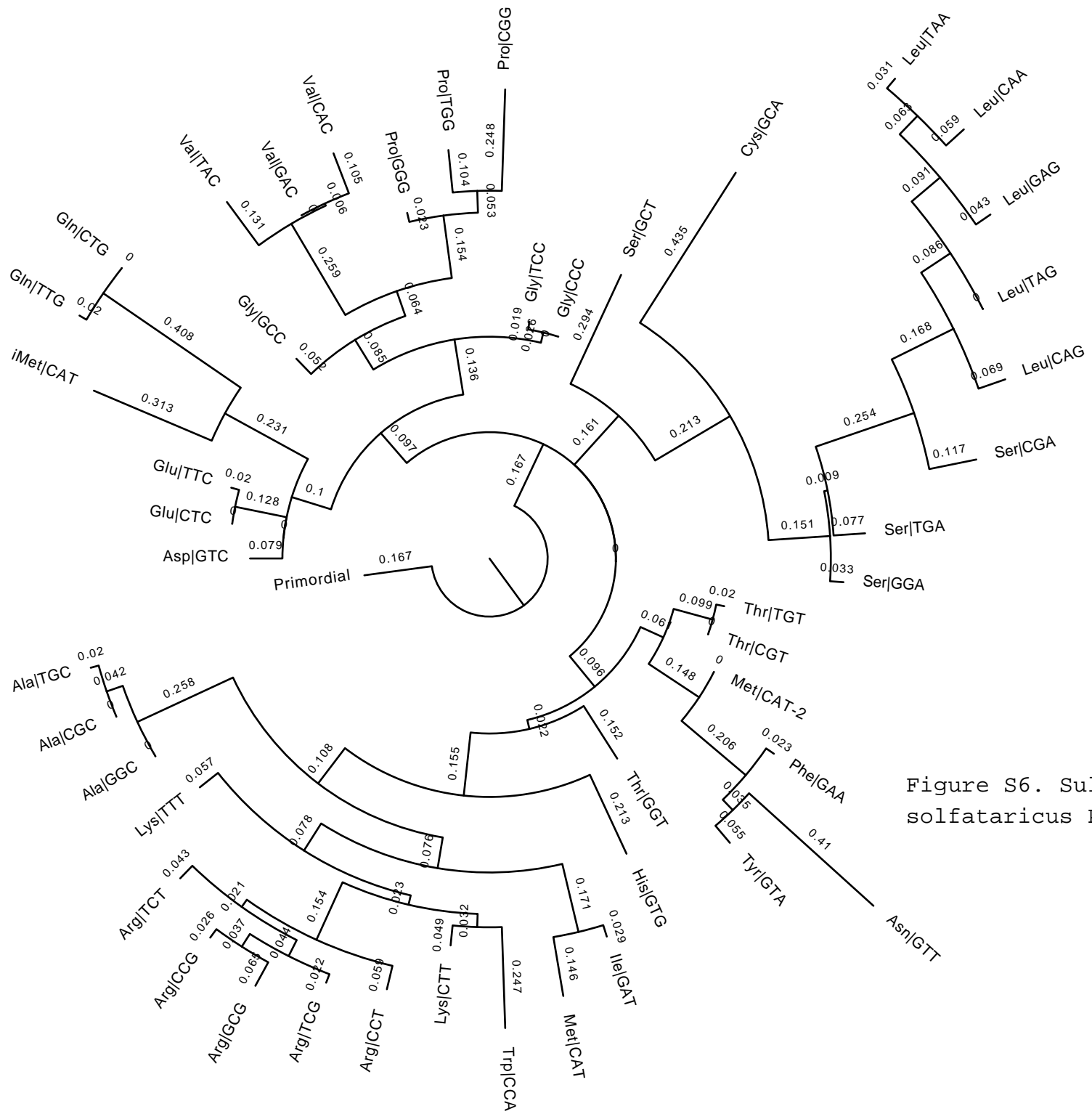


Figure S6. *Sulfolobus solfataricus* P2.



Figure S7. *Aeropyrum pernix* K1.

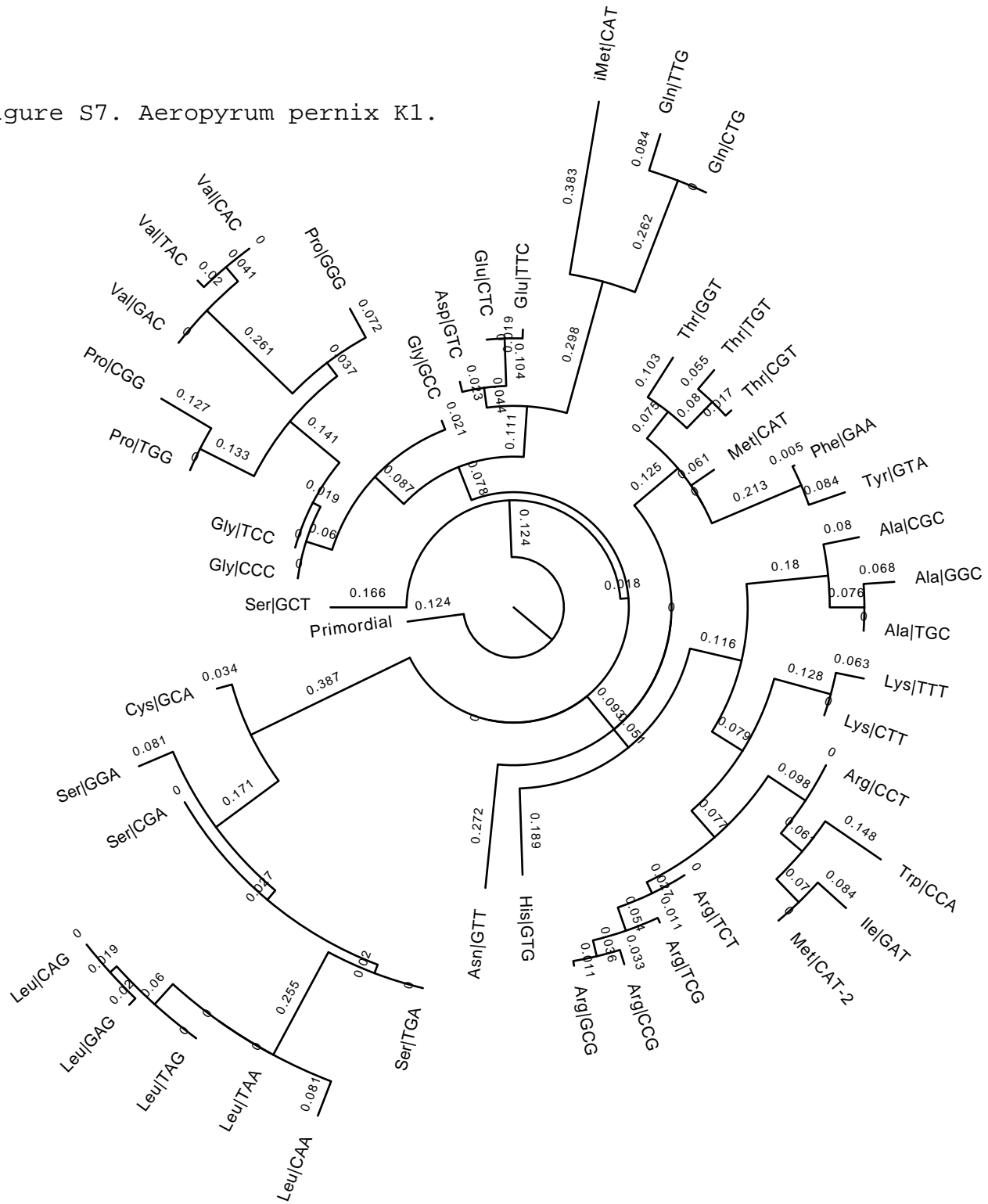
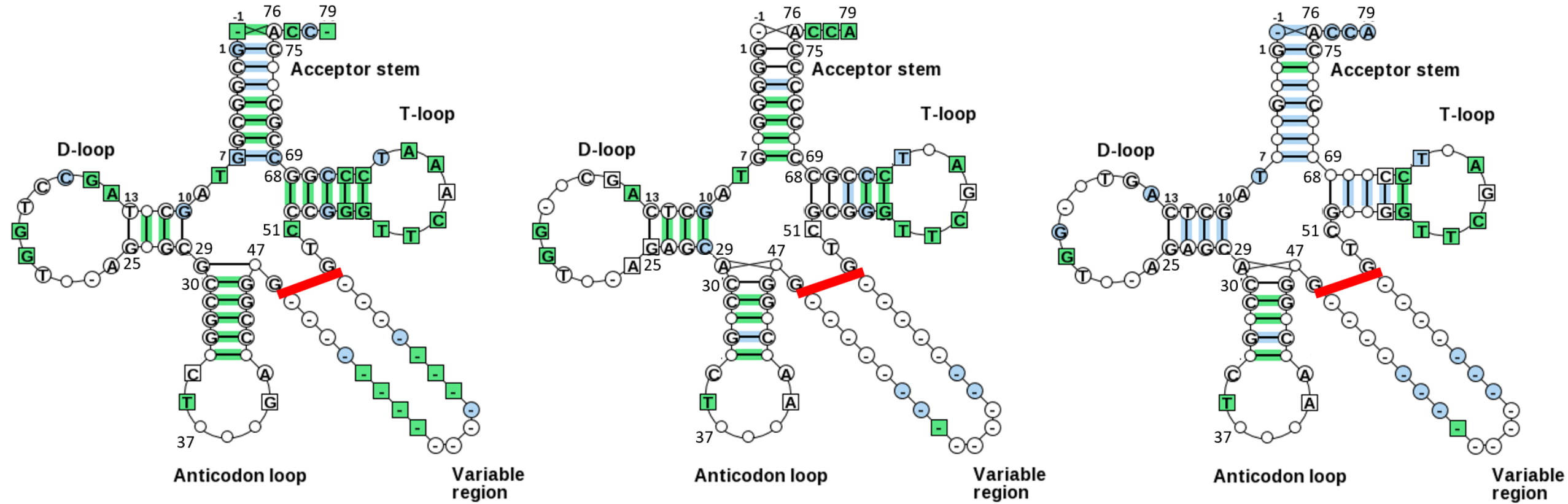




Figure S9. Typical tRNAs.

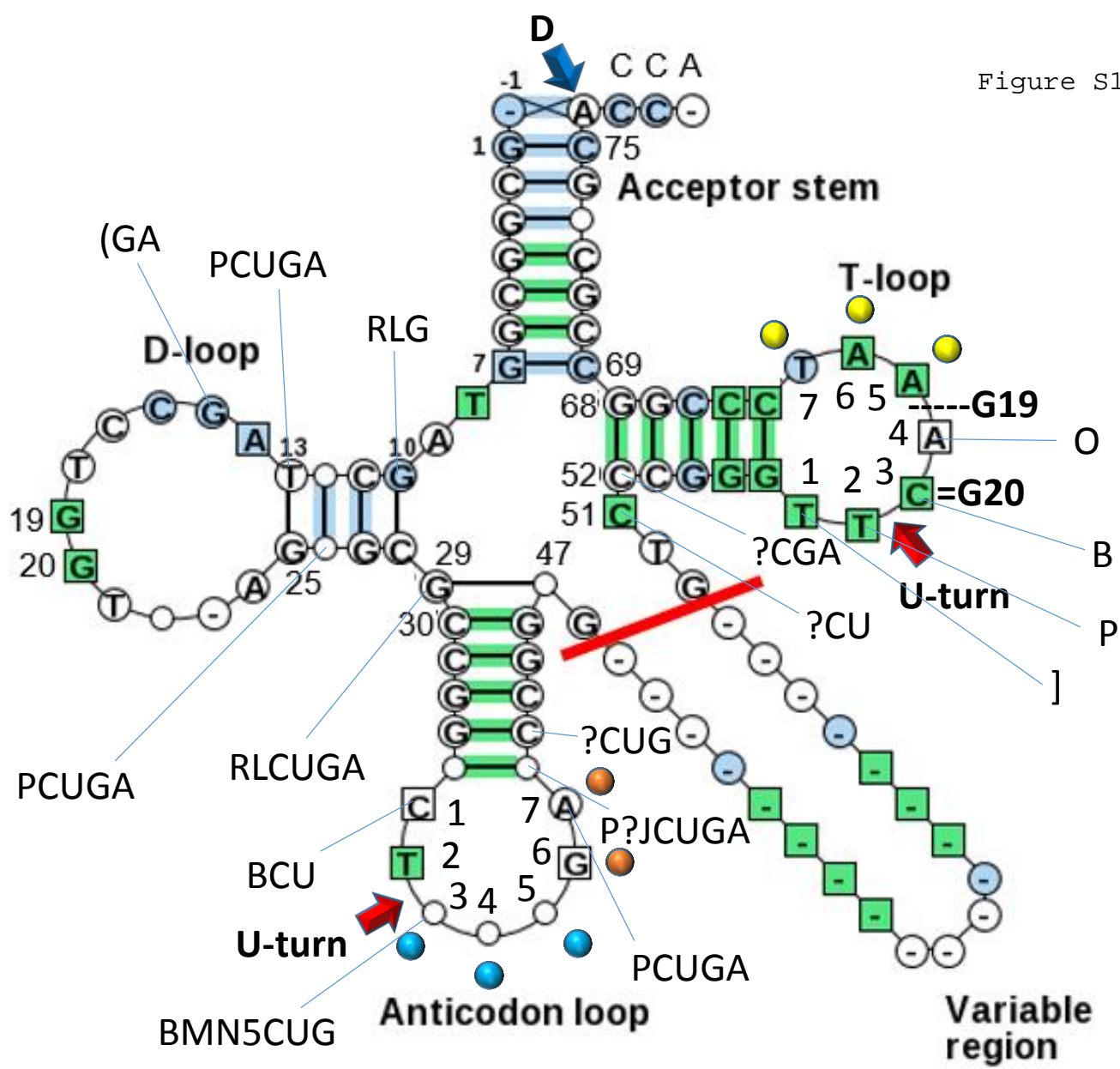


*Pyrococcus furiosus* DSM 3638

*Thermus thermophilus* HB27

*Escherichia coli* K12

Figure S10. tRNA modifications in archaea.



Key:

- R N2-N2-dimethylguanosine
- L N2-methylguanosine
- P pseudouridine
- ( archaeosine
- B 2'-O-methylcytidine
- M N4-acetylcytidine
- N unknown modified uridine
- 5 5-methoxyuridine
- ? 5-methylcytidine
- J 2'-O-methyluridine
- ] 1-methylpseudouridine
- O 1-methylinosine

Modifications from *Haloferax volcanii*