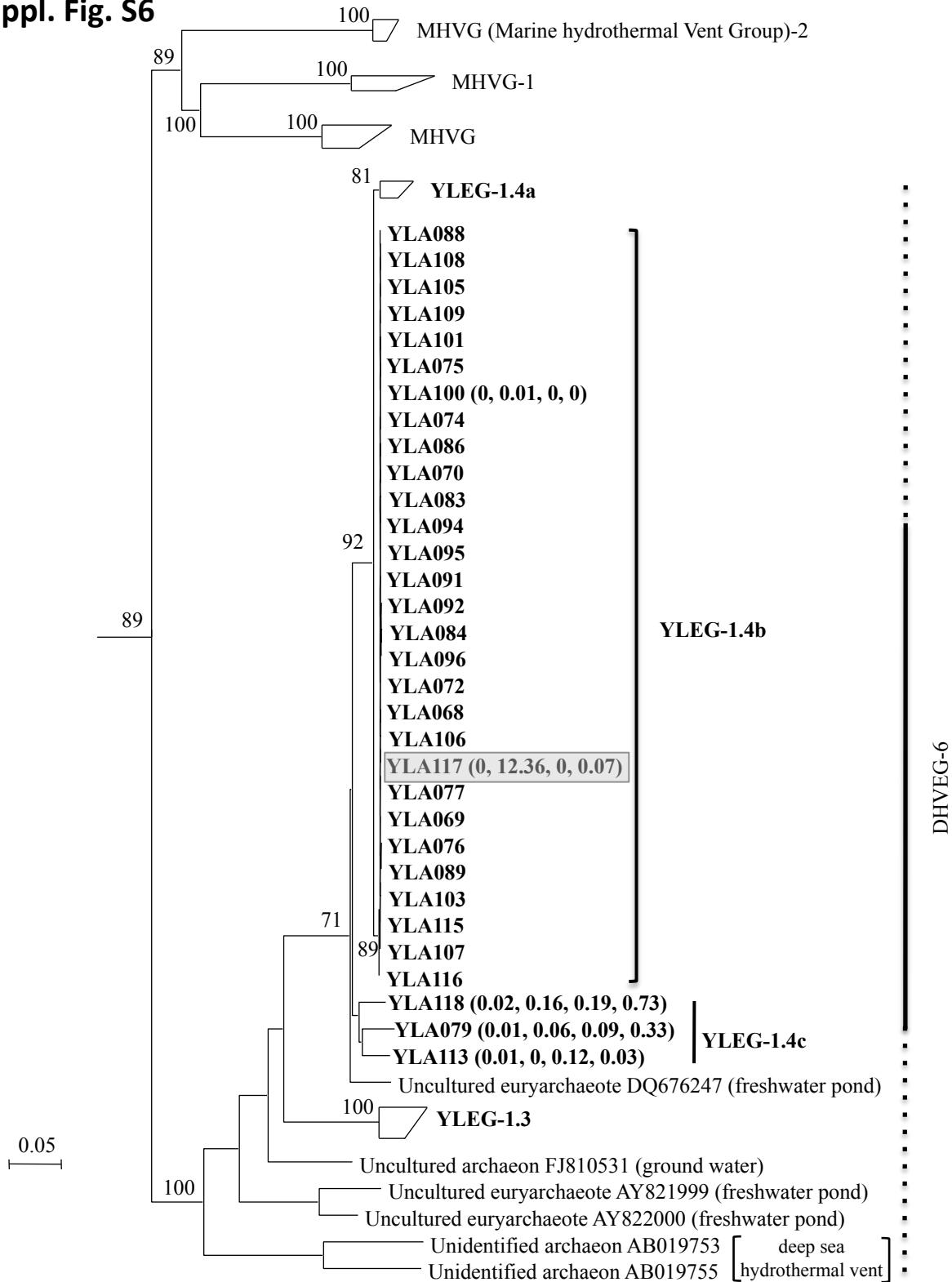


Kan et al.
Suppl. Fig. S6



Supplementary Figure S6. DHVEG-6 (Deep Hydrothermal Vent Euryarchaeal Group-6) and YLEG-1.4b. Clone distribution across the Lake is shown in parentheses as the % of 454 sequences retrieved from the West Thumb Deep vent, Otter vent, Inflated Plain Photic, and Southeast Arm Photic zones. Clones highlighted in grey boxes represent clones that represented at least 10% of the pyrosequencing clones in at least one location, whereas clone designators lacking parenthetical data were not found in the pyrosequencing dataset.