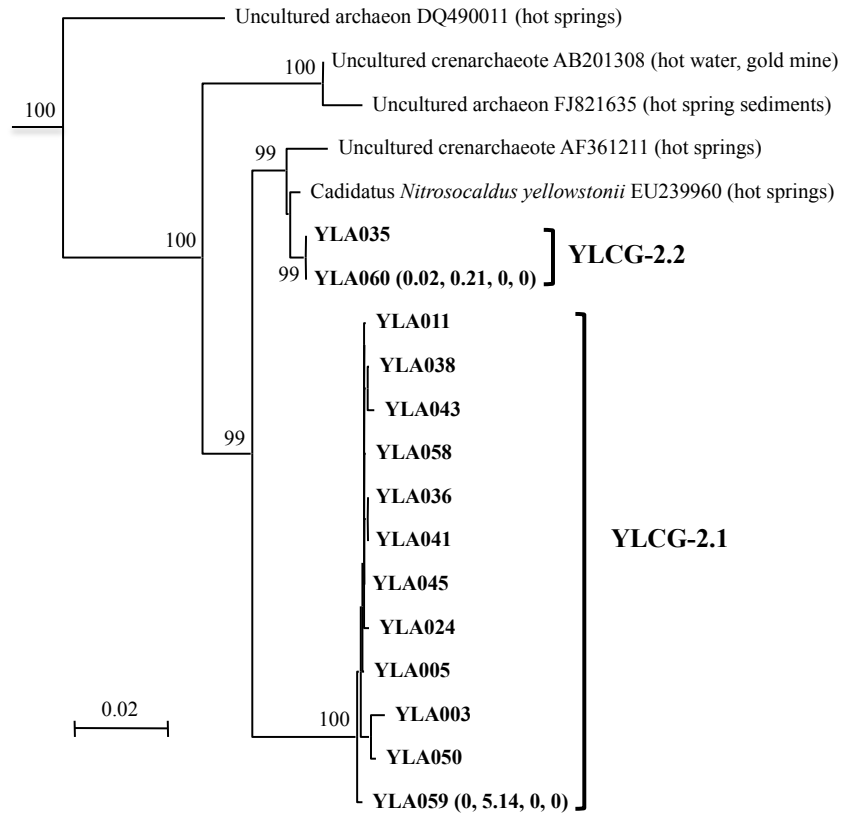


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Suppl. Fig. S5



Supplementary Figure S5. *Candidatus Nitrosocaldus* and YLCG-2.1 and YLCG-2.2. 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. Clone designators highlighted in grey boxes represent clones that represented at least 10% of the pyrosequencing clones in at least one location, whereas those lacking parenthetical data were not found in the pyrosequencing dataset.