S4 Fig. Relative sequence abundances of Archaeal and Bacterial phyla or class based on the 833,468 unique OTU reads (out of 5,141,055 total reads) obtained from separate reactions in triplicate for each depth (10 m, 25 m, 100 m, 130 m, and 250 m) in the Namibian Upwelling seawater at Station 116. The Thaumarchaeota, which consist of all AOA, constitutes at least 25% of the microbiota at depth 100 m and below.