Investigating bacterial endosymbiosis in *Leptogorgia virgulata* using a molecular and SEM approach

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The common sea-whip, *Leptogorgia virgulata* is reported to be asymbiotic in terms of endosymbiotic dinoflagellates. Though we tend to believe that *L. virgulata* uses possible bacterial symbionts as a mechanism to reduce acidic environments within the gorgonian tissue to form calcium carbonate spicules. This study investigates urease production by bacteria within *L.* virgulata to reduce acidity. The goal of the study is to characterize possible bacterial symbionts and using SEM determine locality within *L. virgulata* tissue. Possible microbe symbionts were isolated from *L. virgulata* tissue through a series of dilutions where bacteria colonies unique to surface bacteria were cultured and tested for urease production. Bacteria that were positive for urease production were then prepared for SEM analysis and a series of magnification SEM references for cultured bacteria were compared to freeze fractured *L. virgulata* SEM prepared samples to determine possible microbe symbiont location in the gorgonian tissue. Possible bacterial endosymbionts were then and using the gorgonian tissue. Possible bacterial bacteria were the prepared for SEM analysis and a series of magnification SEM references for cultured bacteria were compared to freeze fractured *L. virgulata* SEM prepared samples to determine possible microbe symbiont location in the gorgonian tissue. Possible bacterial endosymbionts were characterized using univ518r primers typical for identifying universal bacteria. Preliminary molecular data suggests there are unique microbes inhabiting *L. virgulata* tissue, though their location is still being investigated.