# Supplementary Tables

**Supplementary Table 1.** Location and metadata of stations sampled in this study.

**Supplementary Table 2.** Primer sequences and references used in this study

**Supplementary Table 3.** Isolates obtained in this study with clustering to OTUs

**Supplementary Table 4.** Representative isolates with accession numbers, isolate ID, Rfam accession and pairwise similarity to GenBank sequences.

**Supplementary Table 5.** Reference sequences with accession used in alignment.