SM Table 1. Read abundances assigned to each DNA source interpretation category for each siphonophore species.

SM Table 2. Read abundances assigned to each DNA source interpretation category for each siphonophore species by barcode.

SM Table 3. Read abundances assigned to each DNA source interpretation category for each siphonophore specimen.

SM Table 4. Read abundances assigned to each DNA source interpretation category for each siphonophore specimen by barcode.

SM Table 5. Read abundances assigned to each OTU broad taxon for each siphonophore species.

SM Table 6. Read abundances assigned to each OTU broad taxon for each siphonophore species by barcode.

SM Table 7. Read abundances assigned to each OTU broad taxon for each siphonophore specimen.

SM Table 8. Read abundances assigned to each OTU broad taxon for each siphonophore specimen by barcode.

SM Table 9. Read abundances assigned to each prey OTU broad taxon for each siphonophore species.

SM Table 10. Read abundances assigned to each prey OTU broad taxon for each siphonophore specimen.

SM Table 11. Read abundances assigned to each prey OTU broad taxon for each siphonophore species by barcode.

SM Table 12. Read abundances assigned to each prey OTU broad taxon for each siphonophore specimen by barcode.

SM Table 13. Number of unique sequences assigned to each barcode in each DNA source interpretation category.

SM Table 14. Number of unique sequences assigned to each OTU broad taxon.