



## CORRIGENDUM

# Corrigendum to: The genetic diversity and evolution of diatom-diazotroph associations highlights traits favoring symbiont integration

Please note in the original release of article fny297, the accession numbers for 2 sets of sequences (*rbcL*, *nifH*) generated in our study were incorrect. The correct accession numbers for all the sequences submitted to GenBank are as follows: MF536668-MF536674 (*rbcL*), MF527222-MF527230 (18S rRNA),

MF536663-MF536667 (*nifH*), and MF527112-MF527116 (16S rRNA). The supplementary file (SupplTables.MAY2019) is additionally updated with the correct reference to accession numbers.