

Cowart DA, Matabos M, Brandt MI, Marticorena J and Sarrazin J (2019) *Exploring Environmental DNA (eDNA) to Assess Biodiversity of Hard Substratum Faunal Communities on the Lucky Strike Vent Field (Mid-Atlantic Ridge) and Investigate Recolonization Dynamics After an Induced Disturbance*. *Front. Mar. Sci.* 6:783. doi: 10.3389/fmars.2019.00783

Table S1. Sequencing depth values for all taxa and metazoan taxa only, for the four eDNA datasets and across the 18 collected samples.

Sequencing depth (all taxa)			
<i>Dataset</i>	<i>Mean number of reads per sample</i>	<i>Mean number of reads per OTU</i>	<i>Mean number of OTUs per sample</i>
18S-V1V2	38,933.33	257.66	394.16
18S-V4	34,354.67	189.16	517.94
18S-V9	51,124.50	128.79	1,486.28
COI	61,001.39	181.37	822.22
Sequencing depth (metazoan taxa)			
<i>Dataset</i>	<i>Mean number of reads per sample</i>	<i>Mean number of reads per OTU</i>	<i>Mean number of OTUs per sample</i>
18S-V1V2	22,785.44	990.67	112.66
18S-V4	13,310.17	788.10	77.55
18S-V9	31,001.72	1,917.63	96.16
COI	24,863.56	799.18	105.72