

“Investigation of bacterial communities within the digestive organs of the hydrothermal vent shrimp *Rimicaris exoculata* provide insights into holobiont geographic clustering” - Dominique A. Cowart, Lucile Durand, Marie-Anne Cambon-Bonavita, Sophie Arnaud-Haond
 (corresponding author : sophie.arnaud@ifremer.fr)

Sample	Sample Name	Unclassified bacteria (count)	Total assigned	'Bacteria' (%)
1	41_RexRTD22	146	2090	6.99
2	43_RexRTD21	208	1067	19.49
3	47_RexRTD10	638	3109	20.52
4	49_RexRTD9	320	1037	30.86
5	51_RexRTD16	287	2084	13.77
6	53_RexRTD18	121	3280	3.69
7	61_RexRTD7	94	2279	4.12
8	65_RexRTD19	577	1076	53.62
9	67_RexRTD12	1027	4268	24.06
10	87_RexRTD14	259	2608	9.93
11	69_RexTTD1	909	2381	38.18
12	70_RexTE1	1112	2096	53.05
13	72_RexTTE4	464	2161	21.47
14	73_RexTTD2	1139	4808	23.69
15	75_RexTTD12	299	4317	6.93
16	76_RexTE12	263	1263	20.82
17	77_RexTTD11	670	14583	4.59
18	78_RexTE11	1188	5377	22.09
19	79_RexTTD9	381	888	42.91
20	15_RexLTD13	179	956	18.72
21	17_RexLTD14	341	4537	7.52
22	19_RexLTD15	311	3709	8.39
23	21_RexLTD16	183	1349	13.57
24	22_RexLE16	503	4173	12.05
25	24_RexLE13	513	2212	23.19
26	28_RexLE7	192	2282	8.41
27	30_RexLE9	882	2303	38.30
28	32_RexLE10	633	1180	53.64
29	82_RexLW16	668	2734	24.43
30	84_RexLW2	446	3831	11.64
31	85_RexLW11	346	3134	11.04