

Supplemental Materials

Color	Order	Scientific name	Location	H	n	p_1	p_2	p_3	p_{10}	p_{14}	p_{18}
Red	Clionaida	<i>Cliona delitrix</i>	USA	1.10	3	0.333	0.333	0.333	0.0000	0.0000	0.0000
Yellow	Dictyoceratida	<i>Dysidea avara</i>	Spain	1.11	65	0.488	0.398	0.076	0.0004	0.0003	0.0003
Blue	Haplosclerida	<i>Amphimedon chloros</i>	Israel	1.11	1678	0.865	0.010	0.010	0.0032	0.0017	0.00012

Table S1: Three sponge microbiomes highlighted in Figure 6.

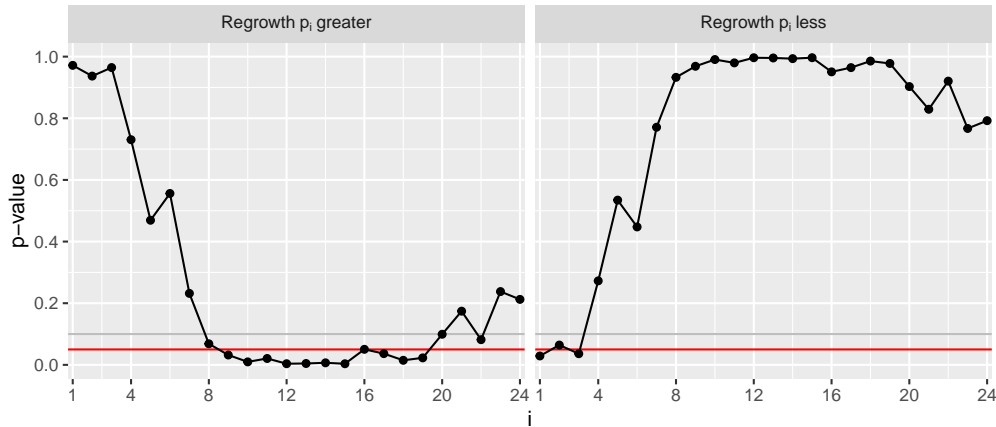


Figure S1: Wilcoxon rank sum P -values for comparisons of p_i between regrowth and non-regrowth reefs, for i from 1 to 24. The red line denotes the $P = 0.05$ significance threshold, and the gray line denotes $P = 0.10$. The left-hand panel presents P -values under the alternative hypothesis that p_i is greater for regrowth reefs than for non-regrowth reefs; the right-hand panel presents P -values under the alternative hypothesis that p_i is lower for regrowth reefs.

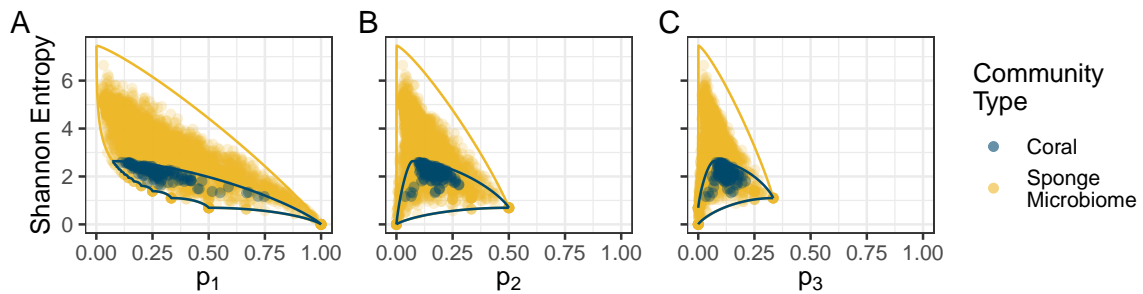


Figure S2: Bounds on Shannon entropy for truncated coral (navy) and sponge microbiome communities (orange), as functions of the abundance of the i th-most abundant taxon. (A) $i = 1$. (B) $i = 2$. (C) $i = 3$. The coral bounds assume $n = 14$, and the sponge bounds assume $n = 1,734$; vectors are truncated to these lengths then normalized to have sum equal to 1.