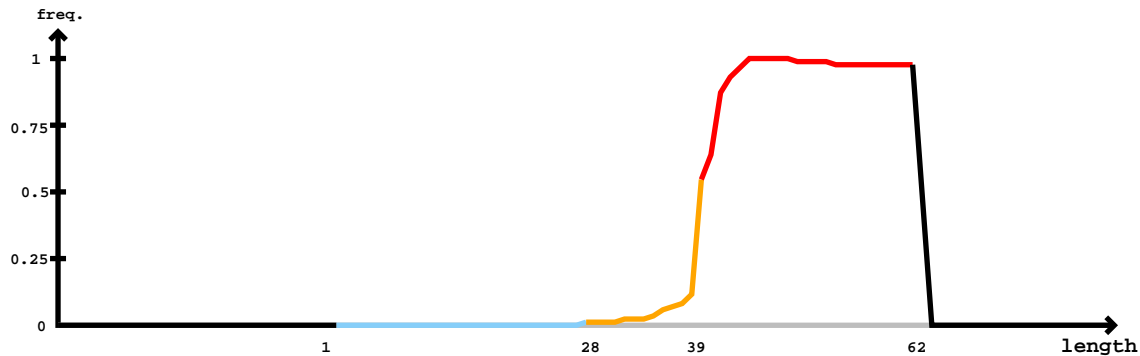
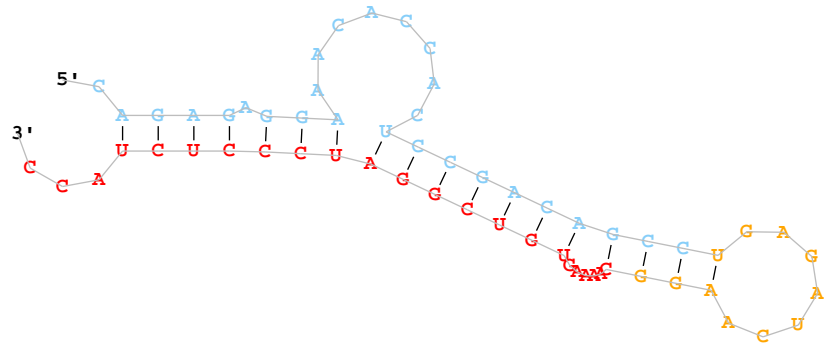


Provisional ID : NC_058077.1_Acropora_millepora_isolate_JS-1_chromosome_12_Amil_v2.1_whole_genome_shotgun_sequence_478802
 Score total : 3.1
 Score for star read(s) : -1.3
 Score for read counts : 0
 Score for mfe : 1.4
 Score for randfold : 0
 Score for cons. seed : 3
 Total read count : 80
 Mature read count : 80
 Loop read count : 0
 Star read count : 0



Star Mature

5'	agcgccugaaguacccccuagguacacagagaggaaacaccacucgcagagccugagaucaaggcaaaagugucggaucccucuaaccaggucggaauuauagccugc	-3'	exp	
..(((((((..((((.....))))).(((..(((.....((((((((.....)))).....))))))))))))))..))))..)).....	reads	mm	sample	
.....cugagaucaaggcaaaaguguc.....	1	0	seq	
.....gaucaaggcaaaagugucggau.....	1	0	seq	
.....caaggcaaaagugucggaucccucuaacc.....	1	0	seq	
.....aaggcaaaagugucggaucccucuaA.....	1	1	seq	
.....aaggcaaaagugucggaucccucuaacc.....	1	0	seq	
.....aggcaaaagugucggaucccucuaU.....	1	1	seq	
.....ggcaaaagugucggaucccucuaacca.....	1	0	seq	
.....gcaaaagugucggaucccucuaacc.....	2	0	seq	
.....gcaaaagugucggaucccucuaU.....	1	1	seq	
.....caaaagugucggaucccucuaA.....	9	1	seq	
.....caaaagugucggaucccucuaacc.....	19	0	seq	
.....caaaagugucggaucccucuaU.....	9	1	seq	
.....aaaagugucggaucccucuaacc.....	2	0	seq	
.....aaaagugucggaucccucuaU.....	2	1	seq	
.....aaaagugucggaucccucuaA.....	4	1	seq	
.....aaagugucggaucccucuaU.....	10	1	seq	
.....aaagugucggaucccucuaacc.....	6	0	seq	
.....aaagugucggaucccucuaacc.....	1	1	seq	
.....aaagugucggaucccucuaA.....	2	1	seq	
.....Gaagugucggaucccucuaacc.....	1	1	seq	
.....aagugucggaucccucuaacc.....	1	1	seq	
.....aagugucggaucccucuaacc.....	1	0	seq	
.....aagugucggaucccucuaU.....	3	1	seq	
.....agugucggaucccucuaacc.....	1	1	seq	
.....agugucggaucccucuaacc.....	1	0	seq	
.....agugucggaucccucuaU.....	1	1	seq	
.....gugucggaucccucuaA.....	1	1	seq	
.....gugucggaucccucuaacc.....	2	0	seq	