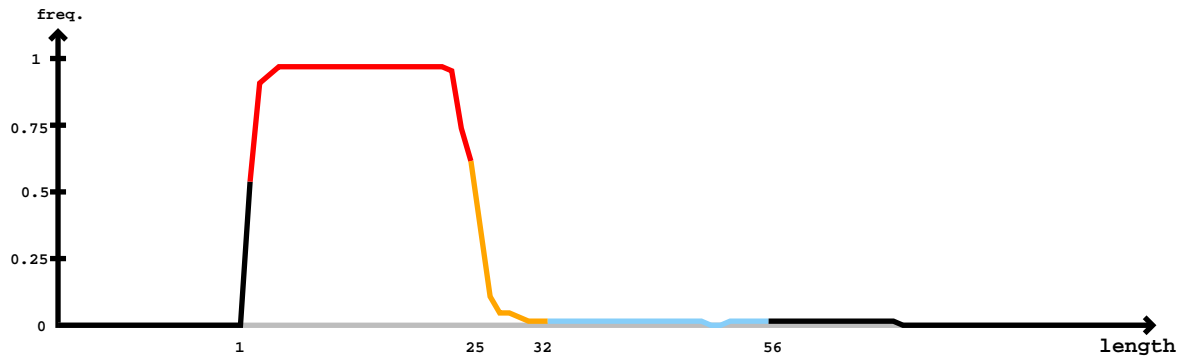
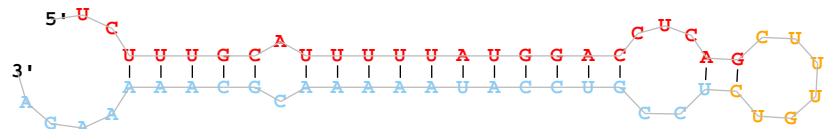


Provisional ID : NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_63339
Score total : 3.4
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 1.7
Score for randfold : 0
Score for cons. seed : 3
Total read count : 63
Mature read count : 63
Loop read count : 0
Star read count : 0



Mature		Star					
5'				-3'	exp		
cacacuggccuugcucuc	ucuuugcauuuuuauaggaccucag	cuuuguc	uccguccauaaaaacgc	caaaaag	gaacuuggccagauaccagcuaucuuugaccucgcgcug	reads	mm
...	((((((.....)))))).....					sample	
.....	ucuuugcauuuuuauaggaccu				1	0	seq
.....	ucuuugcauuuuuauaggaccuc				6	0	seq
.....	ucuuugcauuuuuauaggaccuA				1	1	seq
.....	ucuuugcauuuuuauaggaccuU				2	1	seq
.....	ucuuugcauuuuuauaggaccuca				6	0	seq
.....	ucuuugcauuuuuauaggaccucag				11	0	seq
.....	ucuuugcauuuuuauaggaccCag				1	1	seq
.....	ucuuugcauuuuuauaggaccucaA				3	1	seq
.....	ucuuugcauuuuuauaggaccucaC				1	1	seq
.....	ucuuugcauuuuuauaggaccucagA				2	1	seq
.....	ucuuugcauuuuuauaggaccucagcuuug				1	0	seq
.....	cuuugcauuuuuauaggaccuU				1	1	seq
.....	cuuugcauuuuuauaggaccuA				2	1	seq
.....	cuuugcauuuuuauaggaccuc				2	0	seq
.....	cuuugcauuuuuauaggaccuca				3	0	seq
.....	cuuugcauuuuuauaggaccucaC				1	1	seq
.....	cuuugcauuuuuauaggaccucaU				2	1	seq
.....	cuuugcauuuuuauaggaccucaA				2	1	seq
.....	cuuugcauuuuuauaggaccucag				4	0	seq
.....	cuuugcauuuuuauaggaccucagU				3	1	seq
.....	cuuugcauuuuuauaggaccucagc				1	0	seq
.....	cuuugcauuuuuauaggaccucagcu				2	0	seq
.....	cuuugcauuuuuauaggaccucagcuu				1	0	seq
.....	uuugcauuuuuauaggaccucag				1	0	seq
.....	uuugcauuuuuauaggGccucagc				1	1	seq
.....	uugcauuuuuauaggaccucagcC				1	1	seq
.....	uugcauuuuuauaggaccucagcu				1	0	seq
.....	gcuuugcucucguccauaaaaacAc				1	1	seq
.....	aaagagaacuuggccaUu				1	1	seq