

Provisional ID : NW_025322617.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000030_whole_genome_shotgun_sequenc

Score total : 41.2

Score for star read(s) : 3.9

Score for read counts : 36.3

Score for mfe : 1.6

Score for randfold : 0

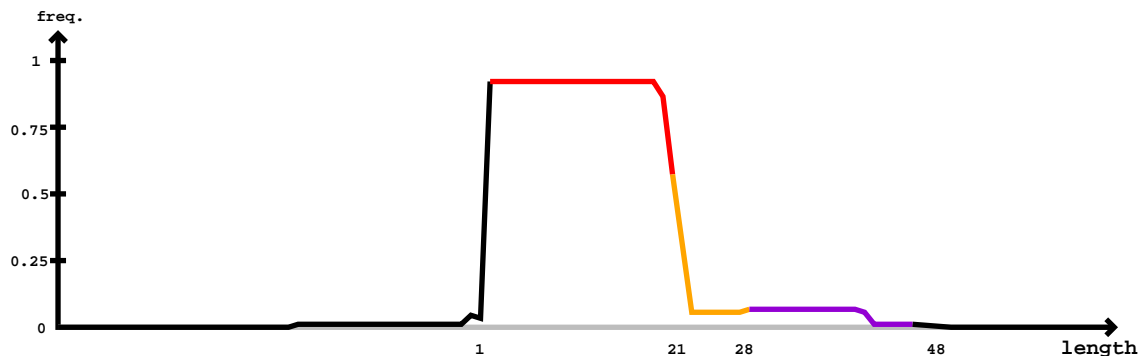
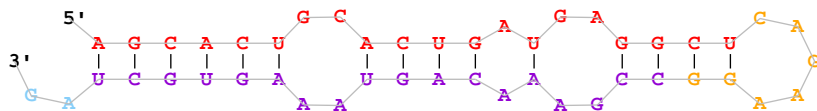
Score for cons. seed : -0.6

Total read count : 83

Mature read count : 82

Loop read count : 0

Star read count : 1



Mature Star

5' -	3' -	obs	exp	reads	mm	sample
aagcucgagggcgccuuuaaagguucgccuucaucucagcaucagcacugcacugacugagggcucagaaggccgaaacaguaaagugcuagugacguuuucuuugcuuu						
aagcucgagggcgccuuuaaagguucgccuucaucucagcaucagcacugcacugacugagggcucagaaggccgaaacaguaaagugcuagugacguuuucuuugcuuu						
(((((.....)))))).....(((.....(((.....)))))).....(((.....)))).....						
.....ucgccuucaucucagcau.....				1	0	seq
.....ucagcacugcacugaggg.....				2	0	seq
.....ucagcacugcacugagggc.....				1	0	seq
.....agcacugcacugagggc.....				1	1	seq
.....agcacugcacugaggg.....				2	0	seq
.....agcacugcacugagggA.....				2	1	seq
.....agcacugcacugagggc.....				23	0	seq
.....agcacugcacugUugagggcu.....				3	1	seq
.....agcacugcacugagggcG.....				4	1	seq
.....agcaAugcacugagggcu.....				1	1	seq
.....agcacugcacugGugagggcu.....				5	1	seq
.....agcaGugcacugagggcu.....				1	1	seq
.....Ggcacugcacugagggcu.....				1	1	seq
.....agcacugcacugagggcu.....				31	0	seq
.....agcacGgcacugagggcu.....				1	1	seq
.....agcacugcacugaGagggcu.....				1	1	seq
.....agcacugcGcugagggcu.....				1	1	seq
.....agcacugcacugagggGggcu.....				2	1	seq
.....cagaaggccgaaacaguaCa.....				4	1	seq
.....agaaggAcgaaacaguaa.....				1	1	seq
.....ccgaaacaguaaagugcu.....				1	0	seq