

Provisional ID : NW_025322643.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000126_whole_genome_shotgun_sequencing

Score total : 58.6

Score for star read(s) : 3.9

Score for read counts : 55.7

Score for mfe : -0.3

Score for randfold : 0

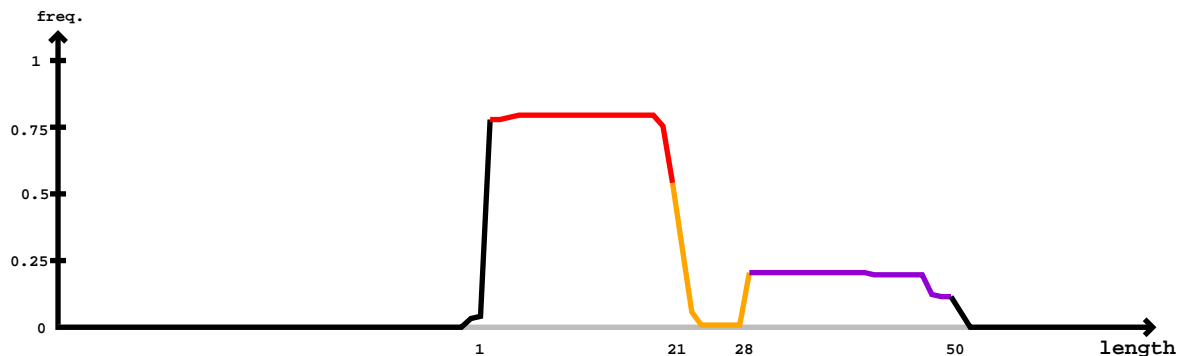
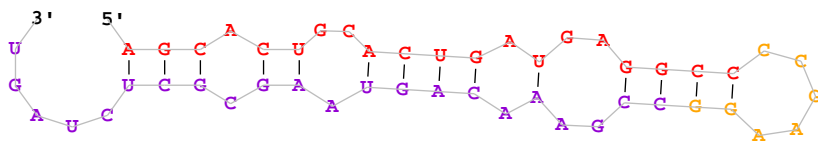
Score for cons. seed : -0.6

Total read count : 121

Mature read count : 97

Loop read count : 0

Star read count : 24



Mature Star

5'	aagcucgagguggcuuuuaaagcuucacccuacucacagcaucagcacugcacugaugaggcccggaaggccgaaacaguaagcgcucuaagaaagucucguuuuguuuuuu	-3'	obs	
	aagcucgagguggcuuuuaaagcuucacccuacucacagcaucagcacugcacugaugaggcccggaaggccgaaacaguaagcgcucuaaguaaaagucucguuuuguuuuuu		exp	
	..(((.((((((((.....)))).....))))))..(((.(.(((.(.(((.....))))..)).))))..)).....	reads	mm	sample
ucagcacugcacugaugagg.....	2	0	seq
ucagcacugcacugaugaggc.....	1	0	seq
ucagcacugcacugaugaggcc.....	1	0	seq
cagcacugcacugaugaggcc.....	1	0	seq
agcacugcacugaugagg.....	2	0	seq
agcacugcacugaugagC.....	1	1	seq
agcacugcacugaugaggc.....	23	0	seq
agcacugcacugaugaggA.....	2	1	seq
agcacugcacugaugaggcG.....	4	1	seq
agcacugcacugaugaggcc.....	25	0	seq
agcacugcacugaugaggccG.....	1	1	seq
agcacugcacugaugaggccc.....	8	0	seq
agcacugcacugaugaggccA.....	18	1	seq
agcacugcacugUugaggcccc.....	1	1	seq
agcacugcacugaugGggcccc.....	2	1	seq
agcacugcacugaugaggccAc.....	3	1	seq
cacugcacugaugaggccA.....	1	1	seq
acugcacugaugaggccc.....	1	0	seq
ccgaaggccgaaacaguaCg.....	1	1	seq
ccgaaacaguaagcgcucu.....	7	0	seq
ccgaaacaguaCgcgcucu.....	1	1	seq
ccgaaacaguaagcgcucG.....	1	1	seq
ccgaaacaguaCgcgcucua.....	1	1	seq
ccgaaacaguaagcgcucuagC.....	1	1	seq
ccgaaacaguaagcgcucuagA.....	2	1	seq
ccgaaacaguaagcgcucuagu.....	10	0	seq
ccgaaacaguaCgcgcucuagu.....	1	1	seq