

Provisional ID : NW\_025322617.1\_Acropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_Amil\_v2.1\_Sc0000030\_whole\_genome\_shotgun\_sequencing

Score total : 44.8

Score for star read(s) : 3.9

Score for read counts : 36.3

Score for mfe : 1.6

Score for randfold : 0

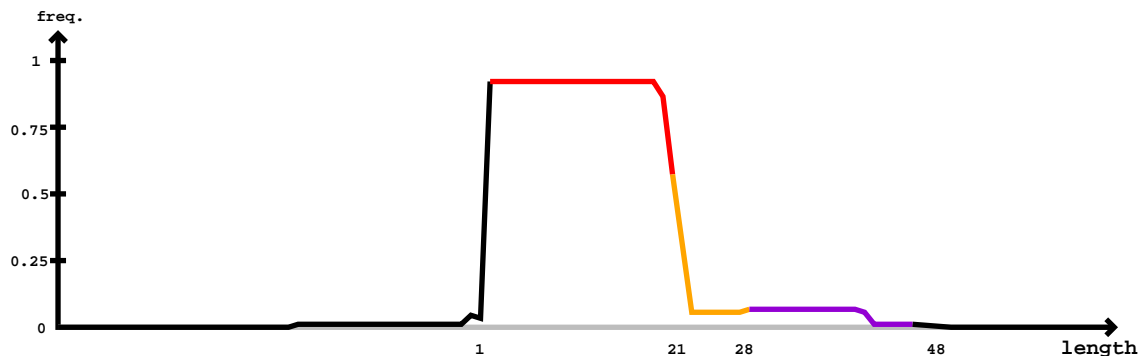
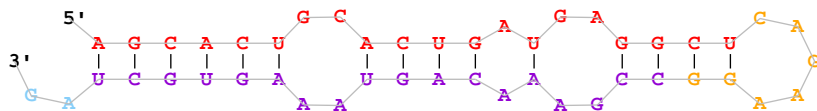
Score for cons. seed : 3

Total read count : 83

Mature read count : 82

Loop read count : 0

Star read count : 1



### Mature Star

5'	obs	exp	reads	mm	sample
aagcucgagggcgccuuuaaagguucgccuucacucagcaucagcacugcacugaugaggcucagaaggccgaaacaguaaagugcuagugacguuuucuuugcuuu					
aagcucgagggcgccuuuaaagguucgccuucacucagcaucagcacugcacugaugaggcucagaaggccgaaacaguaaagugcuagugacguuuucuuugcuuu					
((((( ((((((((((.....)))))).)))))).....((((((( ((((((((((.....)))))).)))))).....((((((( ((((((((((.....)))))).)))))).....)))))).))))).					
.....uucgccuucacucagcau.....			1	0	seq
.....ucagcacugcacugaugagg.....			2	0	seq
.....ucagcacugcacugaugaggc.....			1	0	seq
.....agcacugcacugaugagg.....			2	0	seq
.....agcacugcacugaugagC.....			1	1	seq
.....agcacugcacugaugaggc.....			23	0	seq
.....agcacugcacugaugaggA.....			2	1	seq
.....agcacugcGcugaugaggcu.....			1	1	seq
.....agcacugcacugGugaggcu.....			5	1	seq
.....agcacugcacugaugGggcu.....			2	1	seq
.....agcacGgcacugaugaggcu.....			1	1	seq
.....agcaAugcacugaugaggcu.....			1	1	seq
.....agcacugcacugaCgaggcu.....			1	1	seq
.....agcacugcacugUugaggcu.....			3	1	seq
.....agcacugcacugaugaggcu.....			31	0	seq
.....agcaGugcacugaugaggcu.....			1	1	seq
.....Ggcacugcacugaugaggcu.....			1	1	seq
.....agcacugcacugaugaggcG.....			4	1	seq
.....cagaaggccgaaacaguaCa.....			4	1	seq
.....agaaggAcgaaacaguaa.....			1	1	seq
.....ccgaaacaguaaagugcu.....			1	0	seq