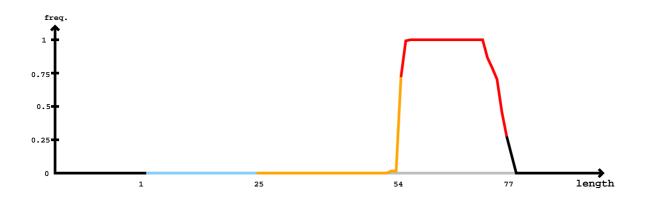
$Provisional\ ID \\ \hspace*{0.5cm} : NC_058067.1_A cropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_84731$

Score total : 0.3
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : -1.3
Score for randfold : 0
Score for cons. seed : 3

AACAU UCAUU CAAUU AAAUU AAAUUU AAAU

Score for cons. seed : 3
Total read count : 246
Mature read count : 246
Loop read count : 0
Star read count : 0



Star Mature			
5'- uaucugagaggagauuasaaugcccacuaccuauucauucuucacaauuuauaacaauuauu	-3' exp		
((((((((((((((((((((((((((((((((((reads	mm	sample
	1	0	seq
	1	1	seq
ugaaugagacuuagugggcuaca	2	0	seq
aaugagacuuagugggcuaaugagacuuagugggcu	25	0	seq
aaugagacuuagugggcCaaugagacuuagugggcC	2	1	seq
Uaugagacuuagugggcu	1	1	seq
aaugagacuuagugggch	4	1	seq
aaugagacuuagugggcuUaaugagacuuagugggcuU	4	1	seq
aaugagacuuagugggcua	7	0	seq
Gaugagacuuagugggcua	1	1	seq
aaugagacuuagugggcCa	2	1	seq
aaugagacuuagugggTua	1	1	seq
aaugagacuuagugggcuaU	8	1	seq
aaugagacuuagugggcuac	6	0	seq
aaugagacuuaguggcuacGaaugagacuuaguggcuacG	1	1	seq
aaugGgacuuagugggcuacaaaugGgacuuagugggcuaca	1	1	seq
aaugagacuuagugggcuaca	18	0	seq
aaugagacuuagugggcuacUaaugagacuuagugggcuacU	1	1	seq
aaugagacuuagugggcuacaa	25	0	seq
aaugagGcuuagugggcuacaa	1	1	seq
aaCgagacuuagugggcuacaa	1	1	seq
aaugagacuuagugggcuacaaGaaugagacuuagugggcuacaaG	1	1	seq
aaugagacuuagugggcuacaaCaaugagacuuagugggcuacaa	6	1	seq
aaugagacuuagugggcuacaaU	41	1	seq
aaugagacuuagugggcuacaCaaaugagacuuagugggcuacaCa	1	1	seq
aaugagacuuaguggcuacaaaaaugagacuuagugggcuacaaa	2	0	seq
aaugagacuuagugggcuacaaUu	13	1	seq
augagacuuaguggcCa	1	1	seq
augagacuuagugggcua	3	0	seq
augagacuuagugggcuac	6	0	seq
	1	1	seq
augagacuuagugggcuaca	32	0	seq
aCgagacuuagugggcuaca	1	1	seq

augagacuuagugggcuacU.....

Star Mature

uaucugagaggagauuasaaugcccacuaccuauucauucuucacaauuuuauaacaauuauu			
augagacuuagugggcuaUa	1	1	seq
augagacuuagugggcuacaa	15	0	seq
Gugagacuuagugggcuacaa	1	1	seq
augagacuuagugggcuacaaU	2	1	seq
augagacuuagugggcuacaaUu	2	1	seq
yanga guya guya guya guya gu	2	0	202