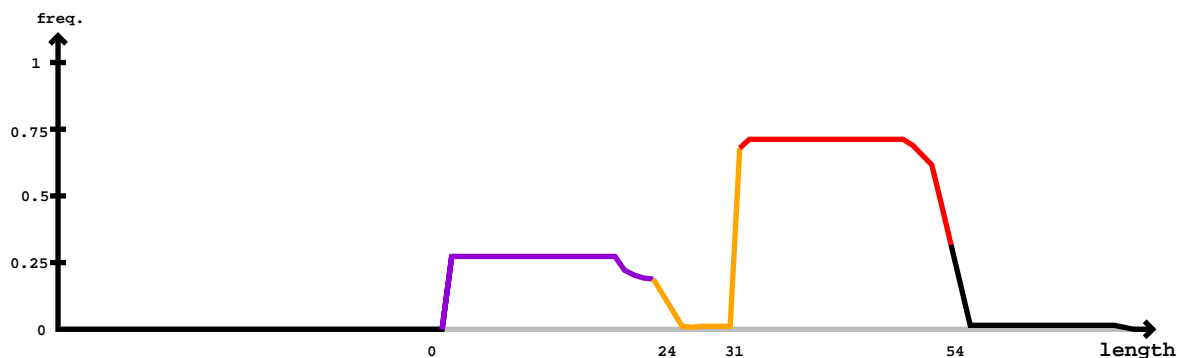|		
Provisional ID	:	NC_058068.1_Acropora_millepora_isolate_JS-1_chromosome_3_Amil_v2.1_whole_genome_shotgun_sequence_152300
Score total	:	132.9
Score for star read(s)	:	3.9
Score for read counts	:	128.6
Score for mfe	:	1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	264
Mature read count	:	190
Loop read count	:	0
Star read count	:	74

Star   **Mature**

```
5'- ucuuagcacaagaggcccucuuauuauaaugagacacaaaugaaaaacacuugaauuaauauuaauaauuguucaaguucuuuucaugauguguuuuauucugcugaacau  -3'   obs
    ucuuagcacaagaggcccucuuauuauaaugagacacacaaugaaaaacacuugaauuaauauuaauaauuguucaaguucuuuucaugauguguuuuauucugcugaacau        exp
    ..(((((((.(((((...)))))....((((((((((..((((((((..(((((((((..............))))))))..)))))))..)))))))))).))))))....  reads   mm    sample
```

sequence	reads	mm	sample
....aaugaaaaacacuugaau.	14	0	seq
....aaugaaaaacacuugaaCu.	1	1	seq
....aUugaaaaacacuugaauu.	1	1	seq
....aaugaaaaacacuugaauu.	3	0	seq
....aaugaaaaacacuugaauua.	3	0	seq
....aaugaaaaacacuugaauuaa.	1	0	seq
....aaugaaaaacacuugaauuaaG.	1	1	seq
....aaugGaaaacacuugaauuaau.	1	1	seq
....aaugaaaaacacuugaauuaau.	18	0	seq
....aaugaaaaacacuugaGuuaau.	1	1	seq
....aaugaaaacGcuugaauuaaua.	1	1	seq
....aaugaaGaacacuugaauuaaua.	1	1	seq
....aaugaaaaacacCugaauuaaua.	1	1	seq
....aaugaaaaacacuugGauuaaua.	1	1	seq
....Gaugaaaaacacuugaauuaaua.	1	1	seq
....aaugaaaaacacuugaauuaaua.	20	0	seq
....aaugaaaaacacuugaauuaaCa.	1	1	seq
....aaugaaaaacacuugaauuaauau.	2	0	seq
....aaugaaaaacacuugaauuaauaA.	1	1	seq
....aaugaaaaacacuugaauuaauaGu.	1	1	seq
....aGuaauaauuguucaaguucuuuucaugau.	2	1	seq
....auaaauuguucaaguucuuuucaugau.	1	0	seq
....uuguucaaguucuuuuca.	6	0	seq
....uuguucaaguucuuuucau.	10	0	seq
....uuguucaaguucuuuucaug.	9	0	seq
....uuguucaaguucuuuucaugU.	2	1	seq
....uuguucaaguucuuuucaugG.	1	1	seq
....uuguucaaguucuuuucauga.	32	0	seq
....uuguucaaguucCuuucauga.	1	1	seq
....uuguucaaguucuuUCcauga.	1	1	seq
....uuguucaaguucuuuucaugaC.	3	1	seq
....uuguucaaguucuuuuGugau.	1	1	seq
....uuguucaaguCcuuuucaugau.	1	1	seq

ucuuagcacaagaggcccucuuauuauaaugagacacacaaugaaaaacacuugaauuaauauuaauaauuguucaaguucuuuucaugauguguuuuauucugcugaacau

....uuguucaaguucuuuucaugaG....................	1	1	seq
....uuguucaaguucuuuucaugau....................	31	0	seq
....uuguucaaguucuuuuUaugau....................	1	1	seq
....uuguucaaguucuuuucaugauU...................	53	1	seq
....uuguucaaguucuuuucaugauA...................	7	1	seq
....uuguucaaguucuuuucaugaug...................	5	0	seq
....uuguucaaguucuuuucaugauC...................	15	1	seq
....uuguucaaguucuuuucaugauUu..................	1	1	seq
....uguucaaguucuuuucaug......................	2	0	seq
....uguucaaguucuuuucauga.....................	3	0	seq
....uguucaaguucuuuucaugaC....................	1	1	seq
....uguucaaguucuuuucaugau....................	2	0	seq
....uguucaaguucuuuucaugaug...................	1	0	seq
....auuguuuuUuucugcugaa...	1	1	seq
....guguuuuauucugcugaa...	1	0	seq
....guguuuCauucugcugaac..	1	1	seq
....guguuuuauucugcugaac..	1	0	seq