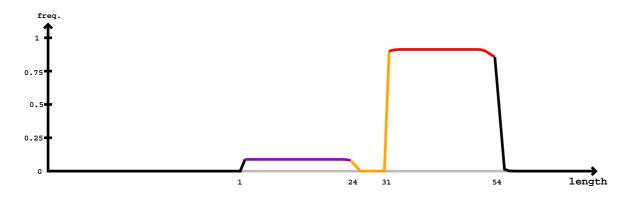| Provisional ID	: NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_99586
Score total	: 954.4
Score for star read(s)	: 3.9
Score for read counts	: 945.9
Score for mfe	: 1.7
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 1867
Mature read count	: 1703
Loop read count	: 0
Star read count	: 164


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

sequence	reads	mm	sample
..................................uggacaaaggcuccaaggaaa..	2	0	seq
..................................uggacaaaggcuccaaggaaau...	11	0	seq
..................................uUgacaaaggcuccaaggaaau...	1	1	seq
..................................uggacaaaggcuccaaggaaUug..	2	1	seq
..................................uggacaaaggcuccaaggaaauA..	11	1	seq
..................................uggGcaaaggcuccaaggaaaug..	1	1	seq
..................................uggacaaaggcuccaaggaaaCg..	2	1	seq
..................................uggacaaaggcuccGaggaaaug..	1	1	seq
..................................uggacaaaggcuccaaggGaaug..	2	1	seq
..................................uggacaaaggcuccaaggaaaug..	54	0	seq
..................................uggaUaaaggcuccaaggaaaug..	1	1	seq
..................................uAgacaaaggcuccaaggaaaug..	1	1	seq
..................................uggacaaaggcuccaaggaaauU..	1	1	seq
..................................uggacaaaggcuccaaggaaaugC...	10	1	seq
..................................uggacaGaggcuccaaggaaaugu..	1	1	seq
..................................uggacGaaggcuccaaggaaaugu..	1	1	seq
..................................uggacaaaggcuccaaggaaaugu..	54	0	seq
..................................uAgacaaaggcuccaaggaaaugu..	1	1	seq
..................................Gggacaaaggcuccaaggaaaugu..	1	1	seq
..................................uggacaaGggcuccaaggaaaugu..	1	1	seq
..................................Aggacaaaggcuccaaggaaaugu..	4	1	seq
..................................uggacaaaggcuccaaggGaaugu..	1	1	seq
....uuugcuaguugcuuuuguc...........................	1	0	seq
....uuugcuaguugcuuuuguU...........................	1	1	seq
....uuugcuaguugcuuuugucA..........................	1	1	seq
....uuugcuaguugcuuuugucc..........................	12	0	seq
....uuugcuaguugcuuuugucU..........................	3	1	seq
....uuugcuaguugcuuuuguccU.........................	14	1	seq
....uuugcuaguugcuuuuguccA.........................	11	1	seq
....uuugcuaguugcuuuuguccc.........................	18	0	seq
....uuugcuaguugcuuuugucccg........................	6	0	seq
....uuugcuaguugcuuuugucccA........................	5	1	seq
....uuugcuaguugcuuuugucccC........................	8	1	seq

Star **Mature**

aaagguauugauaugccaggacuuuuucugguuaagaaauggacaaaggcuccaaggaaauguuuaucauuugcuaguugcuuuugucccguucuuaaacuagacauucaua

Sequence	Count	MM	
uuugcuaguugcuuuugucccU	23	1	seq
uuugcAaguugcuuuugucccgu	1	1	seq
uuugcuaguugcuuuugucccUu	5	1	seq
uuugcuaguugcuuuugucccgu	912	0	seq
uuugcuaguugcuuuuguAccgu	1	1	seq
uuCgcuaguugcuuuugucccgu	5	1	seq
uuugcuaAuugcuuuugucccgu	2	1	seq
uuugcuaguugcuuuugucAcgu	1	1	seq
uuugcuaguugcuuuugucccCu	26	1	seq
uAugcuaguugcuuuugucccgu	1	1	seq
uuugcuaguugcCuuugucccgu	4	1	seq
uuugcuaguuAcuuuugucccgu	1	1	seq
uuugcuaguugcuuuuguccAgu	6	1	seq
uuugcuaguugcuuCugucccgu	2	1	seq
uuugcuaguugcuuuugucccgG	29	1	seq
uuugcuaguugcuuuCgucccgu	8	1	seq
uuugUuaguugcuuuugucccgu	1	1	seq
uuugcuCguugcuuuugucccgu	2	1	seq
uuugcuaguugcuuuuguccUgu	8	1	seq
uuugcuaguugcuuuugucccgA	56	1	seq
uuugcuagCugcuuuugucccgu	7	1	seq
uuugcuaguugcuuuugucccAu	5	1	seq
uuugcuaguugcuuuuguccGgu	1	1	seq
uuugcuaguugcuuuugucccgC	234	1	seq
uuugcuaguugcuuuugCcccgu	3	1	seq
Auugcuaguugcuuuugucccgu	1	1	seq
uuugcuaguugcuuuuAgucccgu	1	1	seq
uuugcuagAugcuuuugucccgu	1	1	seq
uuugcCaguugcuuuugucccgu	2	1	seq
uuugcuaguugcuCuugucccgu	3	1	seq
uuugcuaguCgcuuuugucccgu	2	1	seq
Cuugcuaguugcuuuugucccgu	3	1	seq
uuugcuaguAgcuuuugucccgu	1	1	seq
uuugcuaguugcuuuuguUccgu	3	1	seq
uCugcuaguugcuuuugucccguu	1	1	seq
uuugcAaguugcuuuugucccguu	1	1	seq
uuugcuUguugcuuuugucccguu	2	1	seq
uuugcuaguugcuuuugucccguC	19	1	seq
uuugcuaguugcuuuugucccguu	102	0	seq
uuugcuaguugcuuuugucccgGu	3	1	seq
uuugcuaguugcuuuugucccguA	12	1	seq
uuugcuaguugcuuuuguccUuu	3	1	seq
uuugcuGguugcuuuugucccguu	1	1	seq
uuugcuaguugcAuuugucccguu	1	1	seq
uuugcuaguugcuuuCgucccguu	1	1	seq
uAugcuaguugcuuuugucccguu	1	1	seq
uuugcuaguugcuuuuguccUguu	2	1	seq
uuugcuaguugcuuuugucccgCu	62	1	seq
uuugcuaguugcuuuugucccguuU	15	1	seq
uuugcuaguugcuuuugucccguuA	4	1	seq
uuugcuaguugcuuuugucccgCuc	2	1	seq
uuugcuaguugcuuuugucccguuc	1	0	seq
uuugcuaguugcuuuugucccguuG	2	1	seq
uuugcuaguugcuuuugucccguuUu	2	1	seq
uuugcuaguugcuuuugucccguuAuu	1	1	seq
uugcuaguugcuuuuguccc	1	0	seq
uugcuaguugcuuuugucccgu	10	0	seq
uugcuaguugcuuuugucccguu	4	0	seq
uugcuaguugcuuuugucccguC	1	1	seq
uugcuaguugcuuuugucccguA	2	1	seq
ugcuaguugcuuuuguccc	1	0	seq
ugcuaguugcuuuuguccU	1	1	seq
ugcuaguugcuuuugucccgu	4	0	seq