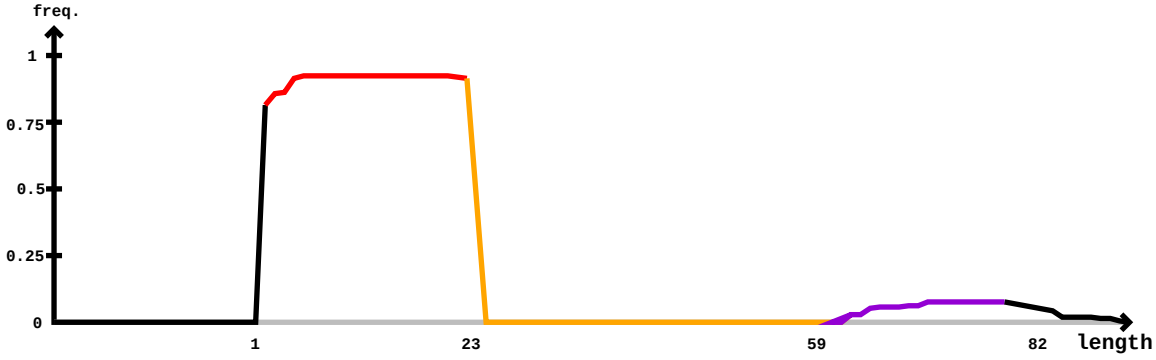
Provisional ID	:	NC_058073.1_Acropora_millepora_isolate_JS-1_chromosome_8_Amil_v2.1_whole_genome_shotgun_sequence_328494
Score total	:	0.8
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.7
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	206
Mature read count	:	194
Loop read count	:	0
Star read count	:	12



Mature **Star**

```
5'- gucuguucagucauuacaggaaaaucucagaccucggccuugagguauugaccgagcgauaacaaggucaauacaucaaggccucggucugagauuucccuguaaugacc  -3'   obs
    gucuguucagucauuacaggaaaaucucagaccucggccuugagguauugaccgagcgauaacaaggucaauacaucaaggccucggucugagauuucccuguaaugacc        exp
    .........((((((((((.(((((((((((..((((((((.(((((((((.........)))))))))..))))))))..))))))))))))).))))))))))).
```

	reads	mm	sample
....aaaaucucagaccucggccu....	1	0	seq
....aaaaucucagaccucggccuug....	130	0	seq
....aaaaucucagaccucggccuuU....	6	1	seq
....Caaaucucagaccucggccuug....	1	1	seq
....aaaaucucagaccucggccuuA....	21	1	seq
....aaaaucucagaccucggccuuC....	1	1	seq
....aaaaucCcagaccucggccuug....	2	1	seq
....aaaaucucagaccucggccuCg....	1	1	seq
....aGaaucucagaccucggccuug....	1	1	seq
....aaaaCcucagaccucggccuug....	1	1	seq
....aaaaucucagaccucggccuuga....	6	0	seq
....aaaucucagaccucggccuuC....	1	1	seq
....aaaucucagaccucggccuug....	6	0	seq
....aaaucucagaccucggccuuA....	1	1	seq
....aaaucucagaccucggccuugG....	1	1	seq
....aaucucagaccucggccuug....	1	0	seq
....aucucagaccucggccuu....	1	0	seq
....aucucagaccucggccuuA....	3	1	seq
....aucucagaccucggccuug....	5	0	seq
....aucucagaccucggccuuC....	1	1	seq
....aucucagaccucggccuuga....	1	0	seq
....ucucagaccucggccuuga....	2	0	seq
....ccucggucugagauuuccc.....	2	0	seq
....ccucggucugagauuucccu.....	4	0	seq
....ucggucugagauuucccuU.....	1	1	seq
....ucggucugagauuucccugC.....	1	1	seq
....ucggucugagauuucccugu.....	3	0	seq
....cggucugagauuucccugA.....	1	1	seq
....ucugagauuucccuguaaug...	1	0	seq
....ugagauuucccuguaaugaA.	2	1	seq
....ugagauuucccuguaaugacU	1	1	seq