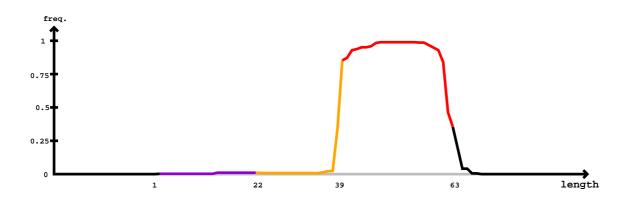|		
Provisional ID	:	NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_28567
Score total	:	181
Score for star read(s)	:	3.9
Score for read counts	:	175.5
Score for mfe	:	2.2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	356
Mature read count	:	355
Loop read count	:	0
Star read count	:	1


```
                           Star                  Mature
5'- uuggacuucaugucauacaauucagggaguaaucgggcucauaauuucaaaucaaaauuacucgcccgauuacucccugaauugugcuccgcucgguccaauuacuauu  -3'   obs
    uuggacuucaugucauacaauucagggaguaaucgggcucauaauuucaaaucaaaauuacucgcccgauuacucccugaauugugcuccgcucgguccaauuacuauu        exp
    (((((((........(((((((((((((((((((((...(((((.......)))))...))))))))))))))))))))))........)))))))........          reads   mm     sample
....................uucagggaguaaucgggcuca.....................................................................      1       0      seq
...........................ucgggcucGuaauuucaaauca.............................................................      3       1      seq
..........................................aaauuacucgcccgauuacu...............................................      1       0      seq
..........................................aaUuacucgcccgauuacucc..............................................      1       1      seq
..........................................aaauuacucgcccgauuacucc..............................................      1       0      seq
..........................................aaauuacucgcccgauuacucA..............................................      1       1      seq
..........................................aaauuacucgcccgauuacuccU.............................................      1       1      seq
...........................................aauuacucgcccgauuacucA..............................................      2       1      seq
...........................................aauuacucgcccgauuacuccc.............................................      1       0      seq
............................................auuacucgcccgauuacucA..............................................      1       1      seq
.............................................uuacucgcccgauuaccA...............................................      1       1      seq
.............................................uuacucgcccgauuacuccc.............................................      1       0      seq
.............................................uuacucgcccgauuacucccu............................................      2       0      seq
.............................................uuacucgcccgauuacucccC............................................      3       1      seq
.............................................uuacucgcccgauuacucccug...........................................      13      0      seq
.............................................uuacucgcccgauuacucccuga..........................................      22      0      seq
.............................................uuacucgcccgauuacucccugC..........................................      9       1      seq
.............................................uuacucgcccgauuacucccugG..........................................      1       1      seq
.............................................uuacucgcccgauuacucccugU..........................................      52      1      seq
.............................................uuacucgcccgauuacucccugaU.........................................      9       1      seq
.............................................uuacucgcccgauuacucccugaUu........................................      3       1      seq
.............................................uuacucgcccgauuacucccugaauu.......................................      1       0      seq
..............................................uacucgcccgauuacucc..............................................      1       0      seq
..............................................uacucgcccgauuacuccc.............................................      2       0      seq
..............................................uacucgcccgauuacuccU.............................................      1       1      seq
..............................................uacucgcccgauuacucccu............................................      1       0      seq
..............................................uacucgcccgauuacucccA............................................      1       1      seq
..............................................uacucgcccgauuacucccug...........................................      11      0      seq
..............................................uacucgcccgauuacucccuC...........................................      3       1      seq
..............................................uacucgcccgauCacucccug...........................................      1       1      seq
..............................................uacucgcccgauuacucccCg...........................................      2       1      seq
..............................................uacucgcccgauuacucccUuga.........................................      2       1      seq
..............................................uacucgcccgauuacucccugG..........................................      1       1      seq
```

uuggacuucaugucauacaauucagggaguaaucgggcucauaauuucaaaucaaaauuacucgcccgauuacucccugaauugugcuccgcucgguccaauuacuauu

Sequence	Count	Mismatch	
....uacucgcccgauuacucccuga.............................	38	0	seq
....uacucgcccgauuacucccugU.............................	8	1	seq
....uacucgcccgauuacucccugaa............................	14	0	seq
....uacucgcccgauuacucccugaU............................	11	1	seq
....uacucgcccgauuacucccugaC............................	3	1	seq
....uacucgcccgauuacucccugaCu...........................	1	1	seq
....uacucgcccgauuacucccugaaA..........................	2	1	seq
....uacucgcccgauuacucccugaau..........................	60	0	seq
....uacucgcccgauuacuccUugaau..........................	1	1	seq
....uacucgcccgauuacucccugaUu..........................	4	1	seq
....uacucgcccgauCacucccugaau..........................	1	1	seq
....uacucgcccgauuacucccugaauC.........................	1	1	seq
....uacucgcccgauuacucccugaauA.........................	1	1	seq
....uacucgcccgauuacucccugaauu.........................	16	0	seq
....acucgcccgauuacuccUug..............................	2	1	seq
....acucgcccgauuacuccUuga.............................	1	1	seq
....acucgcccgauuacuccUugaa............................	1	1	seq
....acucgcccgauuacuccUugaau...........................	3	1	seq
....cucgcccgauuacucccuga.............................	1	0	seq
....cuUgcccgauuacucccuga.............................	3	1	seq
....cuUgcccgauuacucccugaa............................	1	1	seq
....cucgcccgauuacucccugaC............................	1	1	seq
....cucgcccgauuacucccugaau..........................	7	0	seq
....cucgcccgauuacuccUugaau..........................	2	1	seq
....cuUgcccgauuacucccugaau..........................	3	1	seq
....cuUgcccgauuacucccugaauu.........................	1	1	seq
....cucgcccgauuacucccugaauu.........................	2	0	seq
....ucgcccgauuacucccug..............................	1	0	seq
....ucgcccgauuacuccUugaau..........................	1	1	seq
....ucgcccgauuacucccugaauugG.......................	1	1	seq
....Ugcccgauuacucccugaau..........................	2	1	seq
....Ugcccgauuacucccugaauu.........................	3	1	seq
....cccgauuacucccugaauugu.......................	3	0	seq
....ccgauuacucccugaauugu......................	9	0	seq
....cgauuacucUcugaauugugc....................	2	1	seq