Provisional ID	:	Porites_evermani_scaffold_138_7424
Score total	:	578.7
Score for star read(s)	:	3.9
Score for read counts	:	572.2
Score for mfe	:	1
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	1134
Mature read count	:	1074
Loop read count	:	0
Star read count	:	60

Mature   **Star**

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

sequence	reads	mm	sample
..................ucacugaagccgucuauA..	6	1	seq
..................ucacugaagccgucuauU..	16	1	seq
..................ucacugaagccgCcuauc..	1	1	seq
..................ucacugaagccgucuauc..	79	0	seq
..................ucacugGagccgucuauc..	1	1	seq
..................Ccacugaagccgucuaucu...	1	1	seq
..................ucacugaagccgCcuaucu...	1	1	seq
..................ucacugaagccgucuaucA...	5	1	seq
..................ucacugaagccgucuaGcu...	1	1	seq
..................ucacugaagccgucuaucu...	158	0	seq
..................ucacugaagccgucuaucC...	13	1	seq
..................ucGcugaagccgucuaucu...	2	1	seq
..................ucacugaagccgucuaucuu..	29	0	seq
..................ucacugaagccgucuaucuA..	2	1	seq
..................ucacugaagccgucuaucuC..	2	1	seq
..................ucacugaagccAucuaucuu..	1	1	seq
..................ucacugaagccgucuaucuug...	45	0	seq
..................ucacugaagccgucuaucuuC...	1	1	seq
..................ucacugaagccgucuaucuuU...	13	1	seq
..................ucacugaagccgucuaucuuA...	4	1	seq
..................ucacugaGgccgucuaucuuga..	1	1	seq
..................ucacugaagccgucuaucuugU..	167	1	seq
..................ucacugaagccgucCaucuuga..	1	1	seq
..................ucacugaagccgucuaucuugC..	19	1	seq
..................ucacugaagccgucuaucuuga..	84	0	seq
..................ucacugaagccgucuaucuugG..	1	1	seq
..................uUacugaagccgucuaucuuga..	2	1	seq
..................ucacugaGgccgucuaucuugau...	4	1	seq
..................ucacugaaUccgucuaucuugau...	1	1	seq
..................ucacugaagccgucuaucuCgau...	1	1	seq
..................ucacugaaAccgucuaucuugau...	1	1	seq
..................ucacugaagccgucuaCcuugau...	1	1	seq
..................ucacugaagccgCcuaucuugau...	2	1	seq

```
gaaggagcaacuuauccuuuucacugaagccgucuaucuugauuuuaaccaaggucaaauggcuuuuguguuuaaaggauuuguugaaugugagcgcuaacaauaaaa
```

Sequence	Count	Flag	
....................ucacCgaagccgucuaucuugau..	1	1	seq
....................ucacugaagccgucuaucuugaC..	20	1	seq
....................ucacugaagccgucuaucuugau..	305	0	seq
....................ucacugaagccgucuaucuugGu..	3	1	seq
...................Ccacugaagccgucuaucuugau..	2	1	seq
...................Gcacugaagccgucuaucuugau..	1	1	seq
...................uUacugaagccgucuaucuugau..	3	1	seq
....................ucacugaagccgucuaucuugUu..	1	1	seq
....................ucacugaagccgucuaucCugau..	1	1	seq
....................ucacugaagccgucuaucuugauA...	2	1	seq
....................ucacugaagccgucuaucuugaAu...	17	1	seq
....................ucacugaagccgucuaucuugauu...	4	0	seq
....................ucacugaagccgucuaucuugauAu..	31	1	seq
.....................cacugaagccgucuaucC...	3	1	seq
.....................cacugaagccgucuaucu...	2	0	seq
.....................cacugaagccgucuaucuugaC...	1	1	seq
.....................cacugaagccgucuaucuugau...	7	0	seq
......................acugaagccgucuaucuuga..	4	0	seq
....aaggucaGauggcuuuuguguu..	26	1	seq
....aaggucaGauggcuuuuguguuu...	8	1	seq
....aaggucaaauggcuuuuguguuua.......................................	3	0	seq
....aaggucaGauggcuuuuguguuua.......................................	15	1	seq
....aaggucaGauggcuuuuguguuuaa......................................	4	1	seq
....ucaGauggcuuuuguguuua.......................................	4	1	seq
....aggauuuguugaauguga..................	3	0	seq
....aggauuuguugaaugugag.................	1	0	seq
....aggauuuguugaaugugagU................	17	1	seq
....aggauuuguugaaugugagc................	2	0	seq
....aggGuuuguugaaugugagcg...............	1	1	seq
....aggauuuguugaaugugagUg...............	11	1	seq
....aggauuuguugaaugugagcgcA.............	1	1	seq