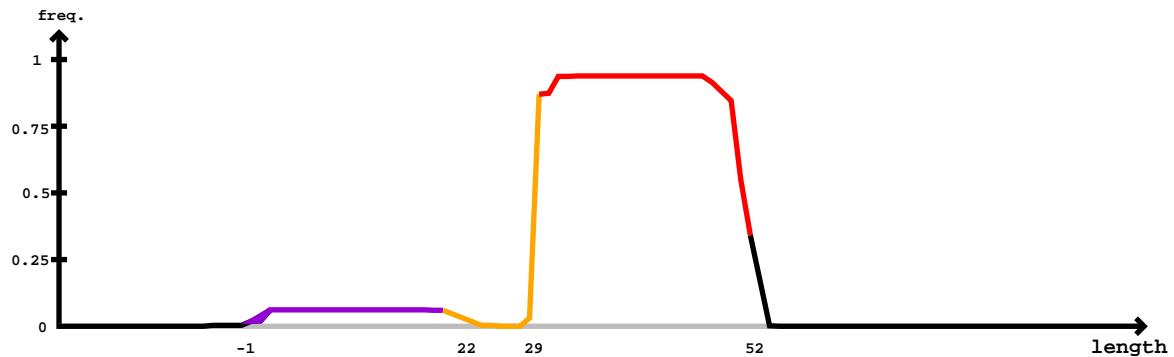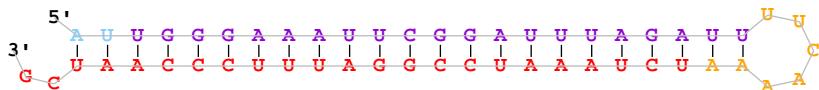Provisional ID   : Porites_evermani_scaffold_434_14215
Score total   : 2.8
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe   : 2.5
Score for randfold   : 1.6
Score for cons. seed   :
Total read count   : 582
Mature read count   : 548
Loop read count   : 0
Star read count   : 34

RNA secondary structure:

```
        5' AUUGGGAAAUUCGGAUUUAGAUU          UUC
           ||||||||||||||||||||||||        C
     3' GC UAACCCUUUUAGGCCUAAAUCUAA        A A
                                            A
```

freq. (y-axis: 1, 0.75, 0.5, 0.25, 0)
length (x-axis: -1, 22, 29, 52)

Star   **Mature**

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

sequence	reads	mm	sample
uucgauugggaaauCcggauuuagauuuc	2	1	seq
auugggaaauucggauuua	1	0	seq
auugggaaauucggauuuagauu	6	0	seq
auugggaaauucggauuuagauC	2	1	seq
uugggaaauucggauuuagauu	1	0	seq
ugggaaauucggauuuagauu	23	0	seq
ugggaaauucggauuuagauC	1	1	seq
Cucuaaauccggauuuccca	1	1	seq
aucuaaauccggauuucccaaC	1	1	seq
aucuaaauccggauuucccaau	10	0	seq
aucuaaauccggauuucccaUu	1	1	seq
aucuaaauccggauuucccaauc	2	0	seq
aucuaaauccggauuucccaauU	3	1	seq
ucuaaauccggauuuccc	13	0	seq
ucuaaauccggauuuccA	1	1	seq
ucuaaauccggauuuccca	10	0	seq
ucuaaauccggauuucccC	1	1	seq
ucuaaauccggauuucccU	7	1	seq
ucuaaauccggauuucccaU	7	1	seq
ucuaaauccggauuucccaa	12	0	seq
ucuaaauccggauuucccaC	1	1	seq
ucuaaauccggauuucccaaC	12	1	seq
ucuaaauccggauuucccaUu	8	1	seq
ucuaaauccggauuucccaaA	3	1	seq
ucuaaauccggauuucccGau	1	1	seq
ucuaaauccggauCucccaau	1	1	seq
ucuaaauccggauuGcccaau	2	1	seq
ucuaaauccggauuucccaau	112	0	seq
ucuaaauccggauuucccaauc	67	0	seq
ucuaaauccggauuucccaauA	1	1	seq
ucuaaauccggauuucccaauU	41	1	seq
ucuaaCuccggauuucccaauc	1	1	seq
Ccuaaauccggauuucccaaucg	3	1	seq

aaauuuguggugcguucgauugggaaauucggauuuagauuuucaaaaucuaaauccggauuucccaaucgaacgcgaaauccgaaaacggauuucaccuccgagaaaucc

....ucuaaauccggauuucccaaucU...	21	1	seq
....ucuaaauccggauuucccaaucC...	4	1	seq
....ucuaaauccggauuucccaaucg...	120	0	seq
....uAuaaauccggauuucccaaucg...	1	1	seq
....ucuaaauccggauuuUccaaucg...	1	1	seq
....ucuaaauccggauuucccGaucg...	2	1	seq
....ucuaaauccggauuucccaaucA...	34	1	seq
....ucuaaauccggauuucccaaucgU..	1	1	seq
....ucuaaauccggauuucccaaucgC..	1	1	seq
....ucuaaauccggauuucccaaucgaU.......................................	1	1	seq
....cuaaauccggauuucccA..	1	0	seq
....cuaaauccggauuucccaaucA...	1	1	seq
....uaaauccggauCucccaau..	1	1	seq
....uaaauccggauuucccaau..	15	0	seq
....uaaauccggauuucccaaC..	4	1	seq
....uaaauccggauuucccaGu..	2	1	seq
....uaaauccggauuucccaauc...	4	0	seq
....uaaGuccggauuucccaauc...	1	1	seq
....uaaauccggaCuucccaauc...	1	1	seq
....uaaauccggauuucccaauU...	1	1	seq
....uaaauccggauuucccaaucA..	1	1	seq
....uaaauccggauuucccaaucg..	6	0	seq
....uaaauccggauuucccaaucU..	1	1	seq
....aauccggauuucccaaucg..	1	0	seq