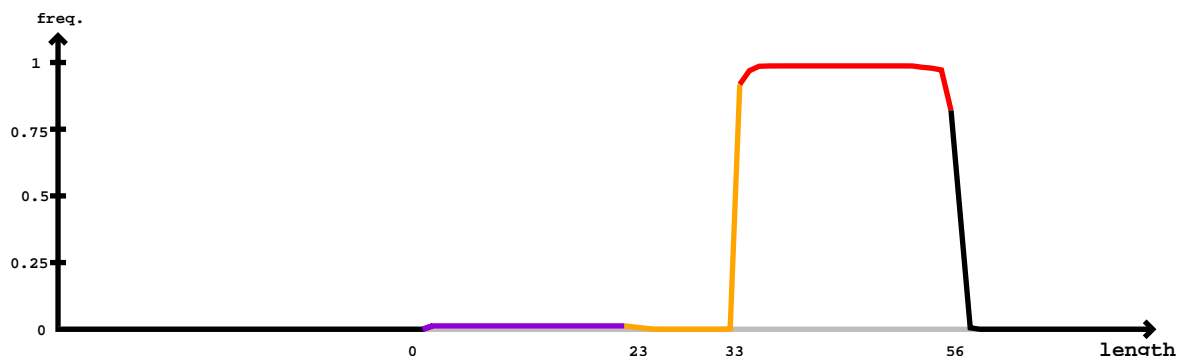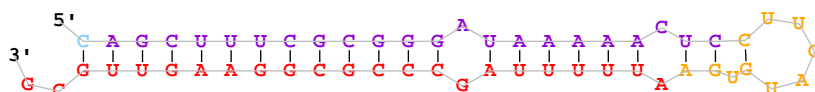| Provisional ID	: NC_058069.1_Acropora_millepora_isolate_JS-1_chromosome_4_Amil_v2.1_whole_genome_shotgun_sequence_193634
Score total	: 7816.8
Score for star read(s)	: 3.9
Score for read counts	: 7808.1
Score for mfe	: 1.8
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 15327
Mature read count	: 15132
Loop read count	: 0
Star read count	: 195


```
                                                                                   Star            Mature
5'- gucagucaguuugggauugauucaagacuauguacucagcuuucgcgggauaaaaacuccuucaugugaauuuuuagcccgcggaaguugcguaaguaugucacauuccacc  -3'    obs
    gucagucaguuugggauugauucaagacuauguacucagcuuucgcgggauaaaaacuccuucaugugaauuuuuagcccgcggaaguugcguaaguaugucacauuccacc          exp
    ((((((((........)))))))....((((((.(((.((((((((((((((.((((((.(((.....).)).)))))).)))))))))))))).)))).))).))).........   reads   mm    sample
```

sequence	reads	mm	sample
..................................agcuuucgcgggauaaaa...	1	0	seq
.................................Ggcuuucgcgggauaaaaacu..	1	1	seq
..................................agcuuucgcgggauaaaaacA..	4	1	seq
..................................agcuuucgcgggauaaaaaUu..	1	1	seq
..................................agcuuucgcgggauaaaaacG..	1	1	seq
..................................agcuuucgcAggauaaaaacu..	1	1	seq
..................................agcuuucgUgggauaaaaacu..	1	1	seq
..................................agcuuucgcgggauaaaaaacC...	8	1	seq
..................................agcuuucgcgggauaaaaaacu...	48	0	seq
..................................agcuuuGgcgggauaaaaacuc...	1	1	seq
..................................agcuuucgcgggauaaaaGcuc...	1	1	seq
..................................agcuuucgcgggauaaaaaacCc..	1	1	seq
.................................Ggcuuucgcgggauaaaaaacuc..	3	1	seq
..................................agcuuucgcgggauaaaaaacuc..	63	0	seq
.................................Ugcuuucgcgggauaaaaaacuc..	2	1	seq
..................................agcuuucgcgggauaaaGacuc..	1	1	seq
..................................agcuuucgcgggauaaaaaacuU..	2	1	seq
..................................agcuuucgcgggauaaaaaacuA..	2	1	seq
..................................agcuuucgcgggauaaaaaacucc...	9	0	seq
..................................agcuuucgcgggauaaaaaacucG...	1	1	seq
..................................agcuuucgcgggauaaaaaacucA...	27	1	seq
..................................agcuuucgcAggauaaaaaacucc..	1	1	seq
..................................agcuuucgcgggauaaaaaacucU...	12	1	seq
..................................agcuuucgcggUauaaaaaacucc..	1	1	seq
..................................agcuuucgcgggauaaaaaacucUu...	1	1	seq
..................................agcuuucgcgggauaaaaaacucAuu..	1	1	seq
....uucaugugaauuuuuagcccgcggaaguugc......................	1	0	seq
....aauuuuuagcccgcggaaguugU.......................	1	1	seq
....aUuuuuuagcccgcggaaguugc.......................	1	1	seq
....aauuuuuagcccgcggaaguugc.......................	6	0	seq
....aauuuuuagcccgcggaaguugcU......................	5	1	seq
....auuuuuagcccgcggaag...........................	5	0	seq
....Uuuuuuagcccgcggaagu...........................	1	1	seq

```
gucagucaguuugggauugauucaagacuauguacucagcuuucgcgggauaaaaacuccuucaugugaauuuuuagcccgcggaaguugcguaaguaugucacauuccacc
```

Sequence	Count	A	B
....auuuuuagcccgcggaagC....................	8	1	seq
....auuuCuagcccgcggaagu....................	1	1	seq
....auuuuuagcccgcggaagA....................	3	1	seq
....auuuuuagcccgcggaagG....................	2	1	seq
....auuuuuagcccgcggaagu....................	54	0	seq
....auuuuuagcccgcggaaguu...................	41	0	seq
....auuuuuagcccgcggaagGu...................	1	1	seq
....auuuuuagcccgcggaaguC...................	6	1	seq
....auuuuuagcccgcggaaguA...................	3	1	seq
....auuuuuagcccgcAgaaguug..................	1	1	seq
....auuuuuagcccgcggaaguuA..................	8	1	seq
....auuuuuagcccgcggGaguug..................	1	1	seq
....auuuuuagcccgcggaaguug..................	73	0	seq
....auuuuuagcccgUggaaguug..................	1	1	seq
....auCuuuagcccgcggaaguug..................	2	1	seq
....Guuuuuagcccgcggaaguug..................	2	1	seq
....auuuuuagcccgcggaaguuC..................	9	1	seq
....auuuuuagcccgcggaaguGg..................	1	1	seq
....auuuuuagUccgcggaaguugc.................	1	1	seq
....Cuuuuuagcccgcggaaguugc.................	1	1	seq
....auuuuuagcccAcggaaguugc.................	1	1	seq
....auuuuuagcccgcggaaguuCc.................	2	1	seq
....auuuuuagcUcgcggaaguugc.................	2	1	seq
....auuuuuagcccgcggaaguugG.................	16	1	seq
....auuuuuagcccgcggaaguugU.................	761	1	seq
....auuuuuagcccgcggaaUuugc.................	3	1	seq
....auuuuuaAcccgcggaaguugc.................	1	1	seq
....auGuuuagcccgcggaaguugc.................	1	1	seq
....auuuCuagcccgcggaaguugc.................	1	1	seq
....auuuuuagcccgUggaaguugc.................	2	1	seq
....auuuuuGgcccgcggaaguugc.................	1	1	seq
....auuuAuagcccgcggaaguugc.................	1	1	seq
....auuuuuagcccgcggGaguugc.................	1	1	seq
....auuuuuagcccgcggaaguugA.................	93	1	seq
....auuuuuagcccgcggaaguCgc.................	5	1	seq
....auuuuuagcccgcggaaAuugc.................	2	1	seq
....auuuuuagcccgcggaaguugc.................	1284	0	seq
....auuuuuagcccgcggaagGugc.................	2	1	seq
....Uuuuuuagcccgcggaaguugc.................	7	1	seq
....auuuuCagcccgcggaaguugc.................	5	1	seq
....auuuuuagcccgcggaaguuAc.................	7	1	seq
....auuuuuagccUgcggaaguugc.................	1	1	seq
....auuuuuagcccgcggaagCugc.................	6	1	seq
....Guuuuuagcccgcggaaguugc.................	11	1	seq
....auuAuuagcccgcggaaguugc.................	4	1	seq
....aCuuuuagcccgcggaaguugc.................	5	1	seq
....auuuuuagcccgcggaGguugc.................	1	1	seq
....auCuuuagcccgcggaaguugc.................	5	1	seq
....auuCuuagcccgcggaaguugc.................	2	1	seq
....auuuuuagcccgcggUaguugc.................	1	1	seq
....aCuuuuagcccgcggaaguugcg................	1	1	seq
....auuuuuagcccgcggaaguugcA................	1933	1	seq
....auuuuuagcccgcggaaguugcU................	7236	1	seq
....auuuuuagcccgcggaaguugcC................	1516	1	seq
....auuuuuagcccgcggaaUuugcg................	1	1	seq
....auuuuuagcccgcggGaguugcg................	2	1	seq
....auuuAagcccgcggaaguugcg................	1	1	seq
....auuuCuagcccgcggaaguugcg................	1	1	seq
....auuuuuagcccgcggaaguCgcg................	1	1	seq
....auCuuuagcccgcggaaguugcg................	1	1	seq
....Cuuuuuagcccgcggaaguugcg................	1	1	seq
....auuuuuagcccgcggaaguugcg................	342	0	seq
....auuuuuaUcccgcggaaguugcg................	1	1	seq
....auuuuuagcccgcggaGguugcg................	5	1	seq
....auuuuuagcccgcggaaguugcgu...............	5	0	seq
....auuuuuagcccgcggaaguugcCu...............	18	1	seq
....auuuuuagcccgcggaaguugcgC...............	2	1	seq
....auuuuuagcccgcggaaguugcAu...............	176	1	seq
gucagucaguuugggauugauucaagacuauguacucagcuuucgcgggauaaaaacuccuucaugugaauuuuuagcccgcggaaguugcUu...............	272	1	seq
....auuuuuagcccgcggaaguugcUua..............	45	1	seq

gucagucaguuugggauugauucaagacuauguacucagcuuucgcgggauaaaaacuccuucaugugaauuuuuagcccgcggaaguugcguaaguaugucacauuccacc

Sequence	Count	Flag	Type
....auuuuuagcccgcggaaguugcAua..................	39	1	seq
....uuuuuUgcccgcggaagu........................	1	1	seq
....uuuuuagcccgcggaagu........................	3	0	seq
....uuuuuagcccgcggaaguu.......................	2	0	seq
....uuuuuagcccgcggaaguug......................	1	0	seq
....Auuuuagcccgcggaaguugc.....................	2	1	seq
....uuuuuagcccgcggaaguugA.....................	3	1	seq
....Guuuuagcccgcggaaguugc.....................	1	1	seq
....uuuuuagcccgcggaaguugU.....................	21	1	seq
....uuuuuagcccgcggaaguugc.....................	33	0	seq
....uuuuuagcccgcggaaguugcU....................	460	1	seq
....uuuuuagcccgcggaaguugcA....................	107	1	seq
....uuuuuagcccgcggaaguugcC....................	114	1	seq
....uuuuuagcccgcggaaguugcg....................	22	0	seq
....uuuuuagcccgcggaaguugcCu...................	1	1	seq
....uuuuuagcccgcggaaguugcAu...................	6	1	seq
....uuuuuagcccgcggaaguugcUu...................	9	1	seq
....uuuuagcccgcggaaguug.....................	3	0	seq
....uuuuagcccgcggaaguugU....................	5	1	seq
....uuuuagcccgcggaaguugA....................	2	1	seq
....uuuuagcccgcggaaguugc....................	12	0	seq
....uuuuagcccgcggaaguugcg...................	4	0	seq
....uuuuagcccgcggaaguugcC...................	52	1	seq
....uuuuagcccgcggaaguugcU...................	134	1	seq
....uuuuagcccgcggaaguugcA...................	32	1	seq
....uuuuagcccgcggaaguugcUu..................	7	1	seq
....uuuuagcccgcggaaguugcAua.................	1	1	seq
....uuuagcccgcggaaguugcU...................	12	1	seq
....uuuagcccgcggaaguugcC...................	3	1	seq
....uuuagcccgcggaaguugcA...................	4	1	seq
....uuuagcccgcggaaguugcUu..................	2	1	seq
....uuagcccgcggaaguugU...................	1	1	seq
....uagcccgcggaaguugcU..................	1	1	seq