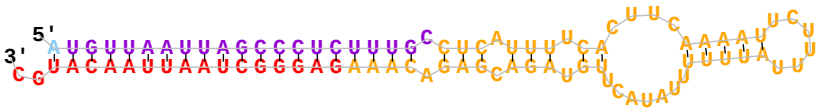
Provisional ID	:	NC_058073.1_Acropora_millepora_isolate_JS-1_chromosome_8_Amil_v2.1_whole_genome_shotgun_sequence_320085
Score total	:	8.7
Score for star read(s)	:	3.9
Score for read counts	:	3.7
Score for mfe	:	1.7
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	19
Mature read count	:	15
Loop read count	:	1
Star read count	:	3



Star **Mature**

```
5'- uuuuguuuacauguuaauuagcccucuuugccucauuuucacuucaaaauucuuuuauuuuuuauacuuguagacgagacaaagagggcuaauuaacaugcaaaccaa  -3'   obs
    uuuuguuuacauguuaauuagcccucuuugccucauuuucacuucaaaauucuuuuauuuuuuauacuuguagacgagacaaagagggcuaauuaacaugcaaaccaa        exp
    ....((((.(((((((((((((((((((.(((.(((.((....(((((......)))))).......)).))).))).)))))))))))))))))))).))))...   reads   mm   sample
    ...........uguuaauuagcccucuuugc.......................................................................   2       0    seq
    ............uuaauuagcccucuuugcA......................................................................   1       1    seq
    ...................gcccucuuugccucauuuu.................................................................   1       0    seq
    ...............................................................uagacgagacaaagaggg....................   1       0    seq
    ......................................................................aagagggcuaauuaacaugc.......   1       0    seq
    ......................................................................aagagggcuaauAaacaugc.......   1       1    seq
    ......................................................................aCgagggcuaauuaacaugc.......   1       1    seq
    ......................................................................aagagggcuaauuaacaugcaaa....   1       0    seq
    ......................................................................agagggcuaauuaacaug........   1       0    seq
    ......................................................................agagggcuaauuaacaugU.......   1       1    seq
    ......................................................................agagggcuaauuaacaugc.......   2       0    seq
    ......................................................................agagggcuaauuaacaugca......   1       0    seq
    ......................................................................agagggcuaauuaacaugcaa.....   1       0    seq
    ......................................................................agagggcuaauuaacaugcaGa....   1       1    seq
    ......................................................................gagggcuaauuaacaugc.......   2       0    seq
    ......................................................................gagggcuaauuaacaugca......   1       0    seq
    ......................................................................gagggcuaauuaacaugcaaacG..   1       1    seq
```