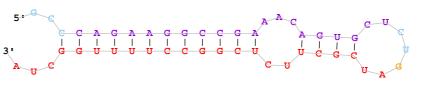|		
Provisional ID	:	NW_025322615.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000015_whole_genome_shotgun_sequen
Score total	:	0
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1.8
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	242
Mature read count	:	234
Loop read count	:	0
Star read count	:	8

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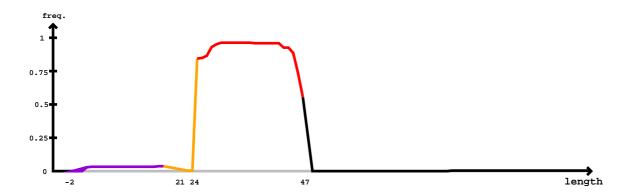
```
5'- aggcccagaaggccgaaacagugcucugaucgcuucucggccuuuuggcuaaguugcgguuguugacaguuuacguugauuuuucagauuguucugccacgaugccgcc  -3'  obs
    aggcccagaaggccgaaacagugcucugaucgcuucucggccuuuuggcuaaguugcgguuguugacaguuuacguugauuuuucagauuguucugccacgaugccgcc       exp
    .((((.(((((((((((...((((.......)))).))))))))))))))).....(((((.((..(((((....)))).))..))....(((((....)))))....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

sequence	reads	mm	sample
.....cagaGggccgaaacagugcu..	3	1	seq
.....cagaaggccgaaacagugcu..	4	0	seq
......agaaggccgaaacagugcuG.......................................	1	1	seq
..................gugcucugaucgcuucucgg...........................	1	0	seq
.....................aucgcuucucggccuuuu..........................	8	0	seq
.....................aucgcuucucggccuuuugg........................	7	0	seq
.....................aucgcuucucggccuuuugA........................	1	1	seq
.....................aucgcuucucggccuuuugC........................	1	1	seq
....................aCcgcuucucggccuuuuggc........................	1	1	seq
.....................aucgcuucucggccuuuuggc.......................	9	0	seq
.....................aucgcuucucggccuuuuggU.......................	6	1	seq
.....................aucgcuucucggccuuuuggA.......................	2	1	seq
.....................aucgcuucucggccuuuugCc.......................	5	1	seq
.....................aucgcuucucggccuuuuggcA......................	10	1	seq
.....................aucgcuucucggccuuuuggUu......................	2	1	seq
.....................aucgcuucucggccuuuuggcu......................	28	0	seq
.....................aucgcuucCcggccuuuuggcua.....................	1	1	seq
.....................aucgcuucucggUcuuuuggcuG.....................	2	1	seq
.....................aucgcuucucggUcuuuuggcua.....................	1	1	seq
.....................aucgcuucucggccuuuuggcuU.....................	3	1	seq
.....................aucgcuucucggccuuuuggcuC.....................	1	1	seq
.....................Uucgcuucucggccuuuuggcua.....................	1	1	seq
.....................Gucgcuucucggccuuuuggcua.....................	2	1	seq
.....................aucgcuucucggccuuuCggcua.....................	3	1	seq
.....................aucAcuucucggccuuuuggcua.....................	1	1	seq
.....................aucgcuucucggccuuuuggcua.....................	105	0	seq
.....................aucgcuucucggccuuuCggcuaa....................	1	1	seq
.....................aucgcuucucggccuuuuggcuaa....................	4	0	seq
......................ucgcuucucggccuuuugg.......................	1	0	seq
.......................cgcuucucggccuuuuggc.....................	1	0	seq
.......................cgcuucucggccCuuuggcu....................	1	1	seq
.......................cUcuucucggccuuuuggcua..................	1	1	seq
.......................cgcuucucggccuuuuggcua..................	1	0	seq

aggcccagaaggccgaaacagugcucugaucgcuucucggccuuuuggcuaaguugcgguuguugacaguuuacuguugauuuuucagauuguucugccacgaugccgcc

...............................gcuucucggccuuuuggU...	2	1	seq
...............................gcuucucggccuuuugAc...	1	1	seq
...............................gcuucucggccuuuuggA...	2	1	seq
...............................gcuucucggccuuuuggc...	8	0	seq
...............................gcuucucggccuuuuggcua...	1	0	seq
...............................Ccuucucggccuuuuggcua...	2	1	seq
................................cuucucggccuuuuggAu...	3	1	seq
................................cuucucggccuuuuggcua...	2	0	seq
.................................uucucggccuuuuggcuaU...	3	1	seq
....uuuucagauuguucugccacgaugccgcc	1	0	seq