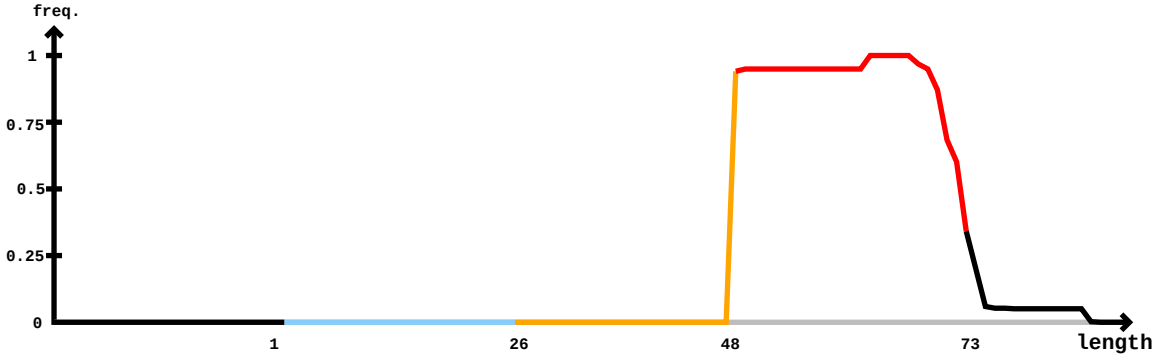
Provisional ID	:	NW_025322653.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000146_whole_genome_shotgun_sequen
Score total	:	0.1
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	-1.5
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	450
Mature read count	:	450
Loop read count	:	0
Star read count	:	0



Star Mature

```
5'- aaaggaaagggaacaucuugaucaggaguuuugucaagcugaccuuguaaugaugcccucaagaacugaagguuuugguuuggcugcacuugaaccuuugucugcagguc  -3'   exp
    (((((...(....).......(((((.((...((((((((((((((..(((.....)))...)).)))))).)))))))).)).)))))).)))))..........
    ..........................................................................agguuuugguuuggcugcU.....................
    ..........................................................................agguuuugguGuggcugca.....................
    ..........................................................................agguuuugguGuggcugcac....................
    ..........................................................................agguuuugguuuggcugcaUu...................
    ..........................................................................agguuuugguuuggcugcacA...................
    ..........................................................................agguuuugguGuggcugcacu...................
    ..........................................................................agguuuugguuuggUugcacu...................
    ..........................................................................agguuuugguuuAgcugcacuu..................
    ..........................................................................agguuuugguGuggcugcacuu..................
    ..........................................................................agguuuugguGuggcugcacuug.................
    ..........................................................................agguuuuggCuuggcugcacuug.................
    ..........................................................................agguuuugguGuggcugcacuuga................
    ..........................................................................agguuuugguuuggcugcacuGga................
    ..........................................................................agguuuuggAuuggcugcacuuga................
    ..........................................................................agguuuuggCuuggcugcacuuga................
    ..........................................................................agguuuugguGuggcugcacuugaa...............
    ..........................................................................agguuuuggCuuggcugcacuugaa...............
    ..........................................................................agguuuugguCuggcugcacuugaac..............
    ..........................................................................agguuuugguCuggcugcacuugaacc.............
    ..........................................................................agguuuugguCuggcugcacuugaaccuu...........
    ..........................................................................gguuuugguGuggcugcacuug.................
    ..........................................................................gguuuugguGuggcugcacuugaa...............
    ...............................................................................cugcacuugaaGcuuugucugca....
    ...............................................................................cugcacuugaaGcuuugucugcag...
```

reads	mm	sample
1	1	seq
14	1	seq
9	1	seq
1	1	seq
2	1	seq
33	1	seq
1	1	seq
1	1	seq
88	1	seq
36	1	seq
2	1	seq
115	1	seq
2	1	seq
2	1	seq
4	1	seq
129	1	seq
1	1	seq
1	1	seq
3	1	seq
1	1	seq
1	1	seq
3	1	seq
23	1	seq
1	1	seq