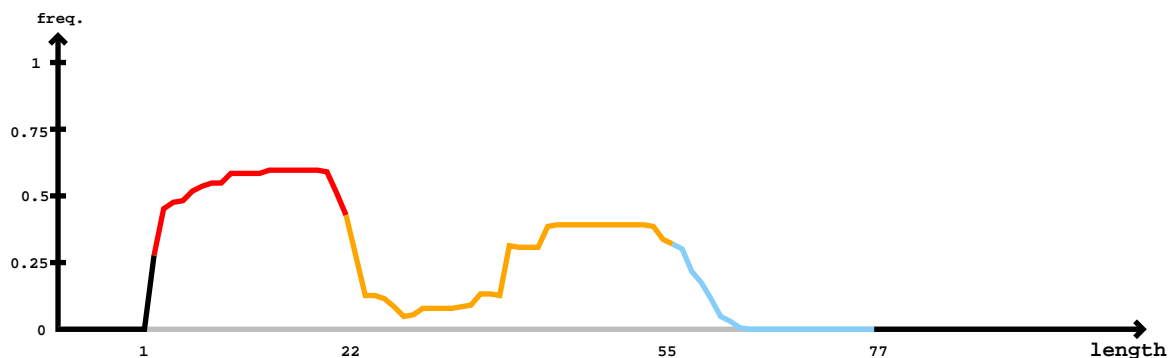


Provisional ID : NW\_025322832.1\_Acropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_Amil\_v2.1\_xfSc0000179\_whole\_genome\_shotgun\_sequ  
 Score total : 2.2  
 Score for star read(s) : -1.3  
 Score for read counts : 0  
 Score for mfe : 0.5  
 Score for randfold : 0  
 Score for cons. seed : 3  
 Total read count : 155  
 Mature read count : 96  
 Loop read count : 59  
 Star read count : 0



## Mature

[illegible]

## Mature

## Star

uaucuaucuaauaauaacuaacugcagacaguaacuaauuuucggccuucugggccucaucagugcagugcuguugucuggggaugaagguaaggcacuaagcccaccucgaguguc

.....ucggccuucugggccucaucagugc.....	1	0	seq
.....ucggccuucugggccucaucagugcag.....	1	0	seq
.....ucggcUcucugggccucaucagugcagu.....	1	1	seq
.....ucggccuucugggccCcaucagugcagugc.....	1	1	seq
.....ccuucugggccucaucagugcaC.....	1	1	seq
.....cuucugggccucaucagugca.....	1	0	seq
.....cuucugggccucauGagugca.....	1	1	seq
.....uucugggccucaucaguA.....	1	1	seq
.....uucugggccucaucagugA.....	1	1	seq
.....uucugggccucaucagugc.....	2	0	seq
.....uucugggccucaucagugca.....	1	0	seq
.....uucugggccucaucagugcaUu.....	2	1	seq
.....ugggccucaucagugcag.....	1	0	seq
.....ugggcUucaucagugcagu.....	6	1	seq
.....ugggccucaucagugcagC.....	2	1	seq
.....ugggccucaucagugcaUu.....	2	1	seq
.....ugggccucaucagugcagu.....	1	0	seq
.....ugggccucaucagugcagug.....	4	0	seq
.....ugggccucaucagugcagCg.....	1	1	seq
.....ugggccucaucagugcaguU.....	2	1	seq
.....ugggccucaucagugcagugc.....	4	0	seq
.....ugggccucaucagugcagugU.....	4	1	seq
.....ugggccucaucagugcagugA.....	1	1	seq
.....ugggccucaucagugcagugcu.....	2	0	seq
.....ugggccucaucagugcagugcA.....	1	1	seq
.....ccucaucagugcagugcu.....	8	0	seq
.....ccucaucagugcagugcug.....	3	0	seq
.....ccucaucagugcagugcugu.....	2	0	seq
.....ccucaucagugcagugcugC.....	1	1	seq
.....ccucaucagugcGgugcuguu.....	1	1	seq
.....cucaucagugcagugcuUu.....	1	1	seq