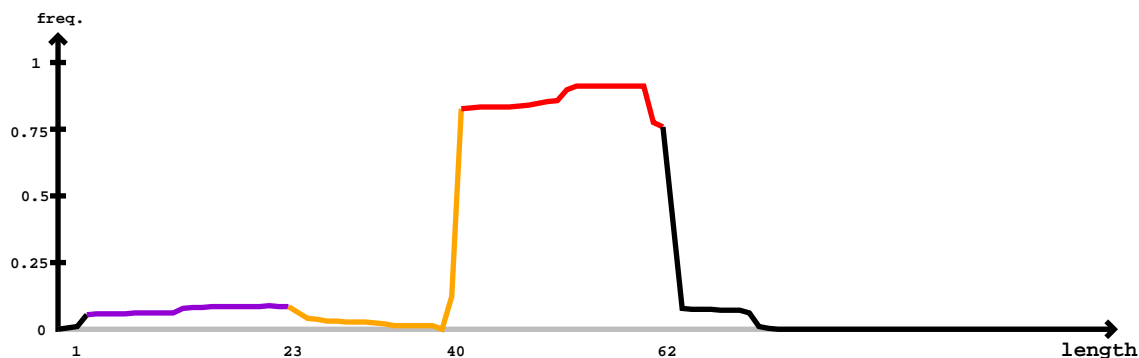


Provisional ID : NW_025322617.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc00000030_whole_genome_shotgun_sequencing
 Score total : 138.8
 Score for star read(s) : 3.9
 Score for read counts : 129.1
 Score for mfe : 2.8
 Score for randfold : 0
 Score for cons. seed : 3
 Total read count : 265
 Mature read count : 247
 Loop read count : 0
 Star read count : 18



Star

[illegible]

Star

Mature

caggguucucucucucugccuccauugucguugagaauggaggcagagaagagagaccucgggaacgagguugguaugaccucugcacucuagucugaugcuagagc			
.....aggcagagaagagagaccUug.....	4	1	seq
.....aggcagagaagagagaccucg.....	1	0	seq
.....aggcagagaagagagaccUugg.....	30	1	seq
.....aggcagagaGgagagaccucgg.....	21	1	seq
.....aggcagagaagagagaccucgg.....	79	0	seq
.....aggcagagaagagagaccucG.....	1	1	seq
.....aggcagagaagagagaccucgC.....	3	1	seq
.....aggcagagaagagagaccucgU.....	6	1	seq
.....aggcagagaagagagaccucgA.....	25	1	seq
.....aggcagagaagagagaccucggC.....	2	1	seq
.....aggcagagaagagagaccucggg.....	1	0	seq
.....aggcagagaagagagaccucggU.....	2	1	seq
.....ggcagagaGgagagaccucgg.....	1	1	seq
.....gcagagaGgagagaccucgg.....	1	1	seq
.....agaagaUagaccucggga.....	1	1	seq
.....gaagagagaccUugggaacg.....	1	1	seq
.....aagagagaGccuggggaacgagguu.....	2	1	seq
.....agagagaccucggggaacgaggu.....	2	0	seq
.....gagagacUcuggggaacgaggu.....	1	1	seq
.....agagaccucggggaacgaggg.....	2	0	seq
.....agagaccucggggaacgaggu.....	10	0	seq
.....gagaccucggggaacgaggg.....	1	0	seq
.....gagaccucggggaacgaggu.....	2	0	seq
.....gagaccucggggaacgagguU.....	1	1	seq