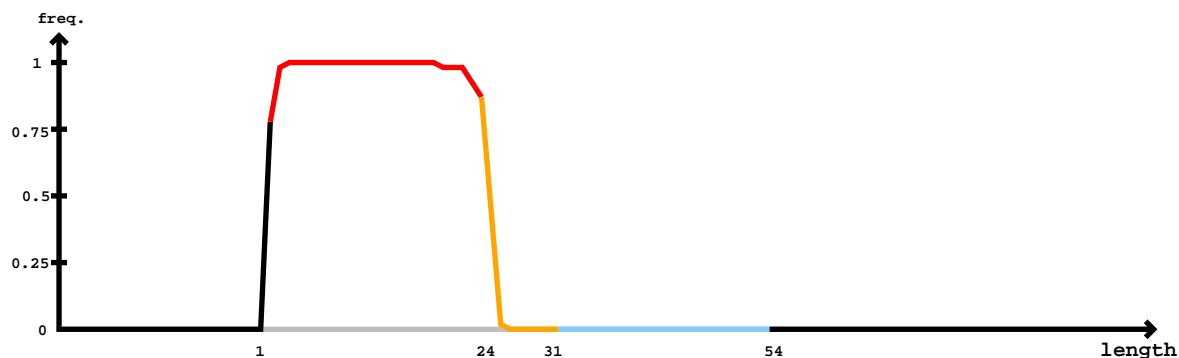


Provisional ID	:	NW_025322622.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000061_whole_genome_shotgun_sequencing
Score total	:	0.6
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.5
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	54
Mature read count	:	54
Loop read count	:	0
Star read count	:	0



Mature		Star					
5'-	aacaacaggccauuucuuuugaauugauuggggaucaggagcuucugagaaagcuccuguaucccaaucaaaaucaaaagagcagccuugcugcagugagagaaggaaggcaaaucg	-3'	exp				
(((((((.(((((((((((((((((((((((...))))))))))))))))))))))))..(((((.(((.....))))..)))))).....	reads	mm				sample
aaugauuggggaucaggaA.....	1	1				seq
aaugauuggggaucaggagU.....	2	1				seq
aaugauuggggaucaggagc.....	1	0				seq
aaugauuggggaucaggagcu.....	2	0				seq
aaugauuggggaucaggagcuC.....	11	1				seq
aaugauuggggaucCggagcuu.....	1	1				seq
aaugauuggggaucaggagcCu.....	3	1				seq
aaugauuggggaucaggagcuu.....	9	0				seq
aaugauuggggaucaggagcuuc.....	2	0				seq
Gauugauuggggaucaggagcuuc.....	1	1				seq
aGuugauuggggaucaggagcuuc.....	1	1				seq
aaugauuggggaucaggagcuuA.....	2	1				seq
aaugauuggggaucaggagcuuU.....	5	1				seq
aaugauuggggaucaggagcuuUu.....	1	1				seq
auugauuggggaucaggagcu.....	1	0				seq
auugauuggggaucaggagcuu.....	2	0				seq
auugauuggggaucaggagcuuA.....	1	1				seq
auugauuggggaucaggagcuuU.....	7	1				seq
uugauuggggaucaggagcuu.....	1	0				seq