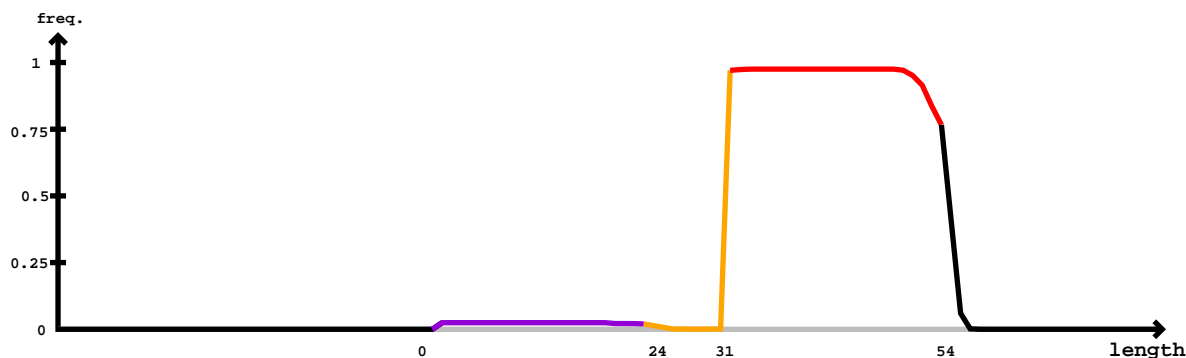|		
Provisional ID	:	NC_058073.1_Acropora_millepora_isolate_JS-1_chromosome_8_Amil_v2.1_whole_genome_shotgun_sequence_341478
Score total	:	25654.9
Score for star read(s)	:	3.9
Score for read counts	:	25646.4
Score for mfe	:	1.6
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	50316
Mature read count	:	49080
Loop read count	:	0
Star read count	:	1236


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                                                         Star                 Mature
5'- ugauaaugcaacugauugucagcaggugugugguugcgaggguauugucuaugaucaaaaauuuuauuauugauuguagacaagccucugacaaccauccauauuauuggacc  -3'   obs
    ugauaaugcaacugauugucagcaggugugugguugcgaggguauugucuaugaucaaaaauuuuauuauugauuguagacaagccucugacaaccauccauauuauuggacc        exp
    ((((((((.......))))))))....(((..(((((((((..(((.(((((((((..(((((.(((...))).))))).)))))))))..)).))))))))..)))..))))...........   reads   mm   sample
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    ...................................aggguauugucuaugau..............................................................   1       0    seq
    ...................................aggguauugucuaCgauc.............................................................   2       1    seq
    ...................................aggguauugucuaugGuc.............................................................   1       1    seq
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    ...................................Gggguauugucuaugauc.............................................................   3       1    seq
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    ...................................aggguauugucuaugaucaU...........................................................   1       1    seq
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    ...................................aggguauCgucuaugaucaaa..........................................................   1       1    seq
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    ...................................aggguauugucuaugaucGaaa.........................................................   4       1    seq
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    ...................................aggguaCugucuaugaucaaaa.........................................................   1       1    seq
    ...................................aggguauugucuaugGucaaaa.........................................................   1       1    seq
    ...................................agggCauugucuaugaucaaaa.........................................................   1       1    seq
    ...................................aggguauugucuaCgaucaaaa.........................................................   1       1    seq
    ...................................Gggguauugucuaugaucaaaa.........................................................   7       1    seq
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    ...................................AggguauugucuaugaucaGaaa........................................................   1       1    seq
    ...................................agggCauugucuaugaucaaaaa........................................................   4       1    seq
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