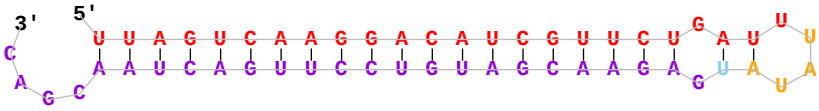
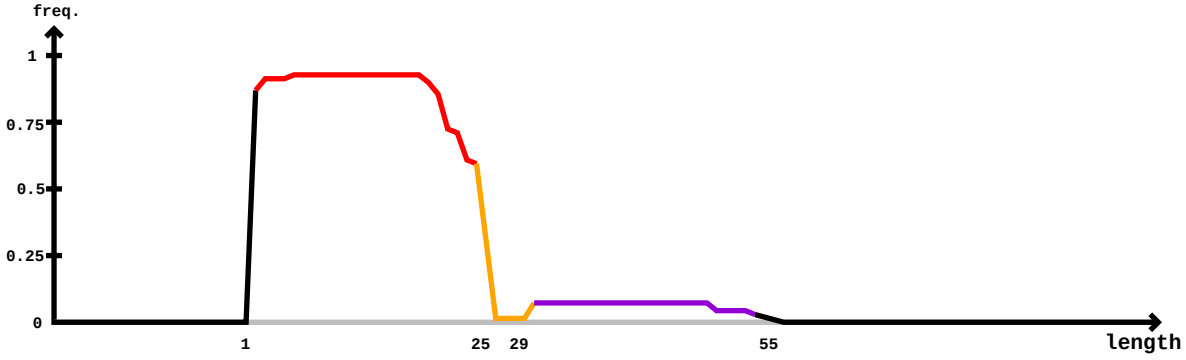
Provisional ID	:	NC_058074.1_Acropora_millepora_isolate_JS-1_chromosome_9_Amil_v2.1_whole_genome_shotgun_sequence_364993
Score total	:	34.5
Score for star read(s)	:	3.9
Score for read counts	:	28.7
Score for mfe	:	2.5
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	68
Mature read count	:	64
Loop read count	:	0
Star read count	:	4






```
                    Mature                Star
5'- ccaaaaacugcgacaguuguuagucaaggacaucguucugauuuauaugagaacgauguccuugacuaacgacuuucgcaguacucuucuaaauuugcuugcaaguuucuuuc  -3'    obs
    ccaaaaacugcgacaguuguuagucaaggacaucguucugauuuauaugagaacgauguccuugacuaacgacuuucgcaguacucuucuaaauuugcuugcaaguuucuuuc         exp
    ......(((((((.(((((((((((((((((((((((((.((....)).))))))))))))))))))))))))))).)))))))........(((((((....)))))))..... reads      mm         sample
    ..................uuagucaaggacaucguu.......................................................................... 2          0          seq
    ..................uuagucaaggacaucguuc......................................................................... 3          0          seq
    ..................uuagucaaggacaucguCcu........................................................................ 1          1          seq
    ..................uuagucaaggacaucguucu........................................................................ 5          0          seq
    ..................uuagucaaggacaucguucA........................................................................ 2          1          seq
    ..................uuagucaaggacaucguucG........................................................................ 1          1          seq
    ..................uuagucaaggacaucguucug....................................................................... 1          0          seq
    ..................uuGgucaaggacaucguucuga...................................................................... 1          1          seq
    ..................uuagucaaggacaucguucuga...................................................................... 5          0          seq
    ..................uuagucaaggacaucguucugau..................................................................... 1          0          seq
    ..................uuGgucaaggacaucguucugau..................................................................... 1          1          seq
    ..................uuagucaaggacaucguucugauA.................................................................... 2          1          seq
    ..................uuagucaaggacaucguucugauC.................................................................... 2          1          seq
    ..................uuagucaaggacaucguucugaCu.................................................................... 1          1          seq
    ..................uuagucaaggacaucguucugauu.................................................................... 32         0          seq
    ...................uagucaaggacaucguucugauC..................................................................... 1          1          seq
    ...................uagucaaggacaucguucugauu..................................................................... 1          0          seq
    ...................uagucaGggacaucguucugauu..................................................................... 1          1          seq
    ......................ucaaggacaucguucuga....................................................................... 1          0          seq
    ........................................uuauaugagaacgauguccuugacCaacg.......................................... 1          1          seq
    ...............................................gagaacgauguccuugacu.......................................... 1          0          seq
    ...............................................gagaacgauguccuugacA.......................................... 1          1          seq
    ...............................................gagaacgauguccuugacCaacgac.................................... 2          1          seq
```