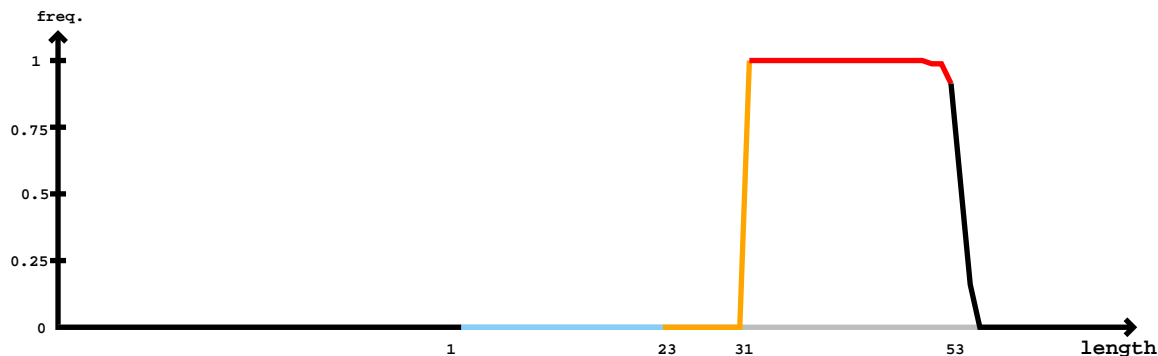
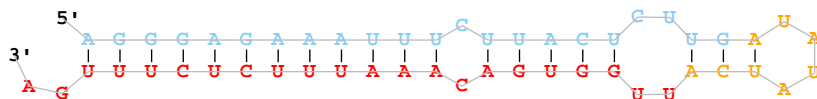


Provisional ID : NW_025322644.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000127_whole_genome_shotgun_sequenc
Score total : 2.5
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 0.8
Score for randfold : 0
Score for cons. seed : 3
Total read count : 81
Mature read count : 81
Loop read count : 0
Star read count : 0



Star Mature

| 5' - | guggaaauaaaaaagcacugaaugucgggguaaaaagagggagaaauucuuacucucugauauaaucauuggugacaaaauucucuugaauuuugaccugagauuu | -3' | exp | | |
|-------|--|-------|-----|--------|--|
| | (((((.....)))..)).....(((((((.....((((((((((((((((((((.....)))))))))))))))))))).....))))..... | reads | mm | sample | |
| |uuggugacaaaauucCuu..... | 1 | 1 | seq | |
| |uuggugacaaaauucucuug..... | 5 | 0 | seq | |
| |uuggugacaaaauucucuuuU..... | 1 | 1 | seq | |
| |uuggugacGaaauucucuuga..... | 1 | 1 | seq | |
| |uuggugacaaaauucucuuga..... | 25 | 0 | seq | |
| |uuggugacaaaauucucuugU..... | 2 | 1 | seq | |
| |uuggugGcaaaauucucuuga..... | 2 | 1 | seq | |
| |uuggugacaaaauucucuuuUa..... | 1 | 1 | seq | |
| |uuggugacaaaCuucucuugau..... | 1 | 1 | seq | |
| |uuggugacaaaauucucuugaC..... | 8 | 1 | seq | |
| |uuggugacaaaauucucuuCgau..... | 1 | 1 | seq | |
| |uCGgugacaaaauucucuugau..... | 1 | 1 | seq | |
| |uuggugacaaaauucucuugau..... | 18 | 0 | seq | |
| |uuggugacaaaauucucuuuUau..... | 1 | 1 | seq | |
| |uuggugacaaaauucucuuuUauu..... | 1 | 1 | seq | |
| |uuggugacaaaauucucuugauC..... | 2 | 1 | seq | |
| |uuggugacaaaauucucuugaau..... | 10 | 0 | seq | |