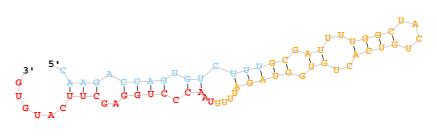
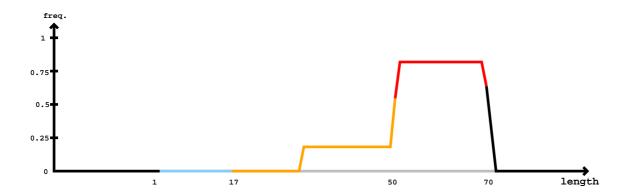
$Provisional\ ID \\ : NW\_025322655.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_shotgun\_sequential mil_v2.1\_Sc0000149\_whole\_genome\_sequential mil_v2.1\_Sc0000149\_whole_genome\_sequential mil_v2.1\_Sc0000149\_whole_genome\_sequential mil_v2.1\_Sc0000149\_whole_genome\_sequential mil_v2.1\_Sc0000149\_w$ 

0.3 Score total Score for star read(s) -1.3 -0.3 Score for read counts -0.9 Score for mfe Score for randfold 0 Score for cons. seed 3 Total read count 11 9 Mature read count 2 Loop read count 0 Star read count





|   | Star  | Mature                                     |         |     |        |
|---|---|--|---------|-----|--------|
| 5 | 5'- uuucacuuuaauuaccacuacaagaccagugucuuugcgauuuuggcuacugucacugugguagauu | uuuaacccuggagcuucaugugguggacaucuccacauaaau | -3' exp |     |        |
|   | ((((((((((((((((((((((((((((((((((                                      | )).)))))).)))))))                          | reads   | mm. | sample |
|   | ugucacugugguagauu   | ıu   | 2       | 0   | seq    |
|   |   | uaacccuggagcuucaugu                        | 1       | 0   | seq    |
|   |   | uaacccuggagcuucaugC                        | 1       | 1   | seq    |
|   |   | uaacccuggagcuucaugug                       | 4       | 0   | seq    |
|   |   | aacccuggagcuucaugug                        | 2       | 0   | seq    |
|   |   |  | _       | _   |        |