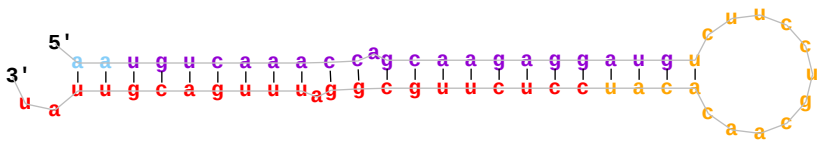
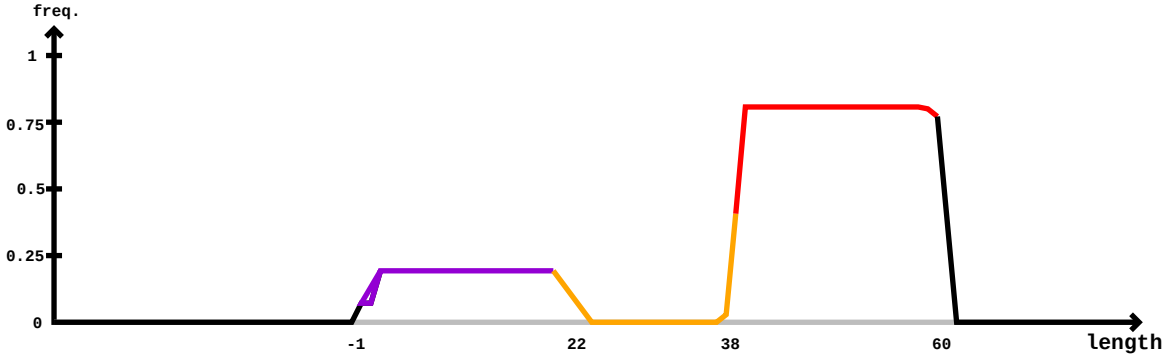
Provisional ID	:	NMRB01000249.1_Notospermus_geniculatus_strain_Ushimado_scaffold249,_whole_genome_shotgun_sequence_8341
Score total	:	0.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	140
Mature read count	:	113
Loop read count	:	0
Star read count	:	27





Star Mature

```
5'- acccugcaucgcgagggcuaggcuacucagaaugucaaaccagcaagaggaugucuuccugcaacacauccucuugcggauuugacguuaugaguggugcuccuauccgau  -3'   obs
    acccugcaucgcgagggcuaggcuacucagaaugucaaaccagcaagaggaugucuuccugcaacacauccucuugcggauuugacguuaugaguggugcuccuauccgau        exp
    .......((((..(((((...(((((((((.((((((((((((.(((((((((((.))))))))))))))).))))))))).))))))))))).)))...))))   reads    mm    sample
    ............................Gaugucaaaccagcaagaggau.......................................................   1        1     Mm2
    ............................aaugucaaaccagcaagaggau.......................................................   2        0     Mm2
    ...............................................................uccucuugcUgauuugacguu.....................   1        1     Mm2
    ...............................................................ccucuugcUgauuugacguua.....................   2        1     Mm2
    ...............................................................ccucuugcUgauuugacguuau....................   2        1     Mm2

    ............................aaCgucaaaccagcaagagga.......................................................   3        1     Fm3
    ...............................................................uccucuugcggauuugacguua.....................   2        0     Fm3
    ...............................................................uccucuugcggauuugacguuau....................   1        0     Fm3
    ...............................................................ccucuugcggauuugacguuGu....................   3        1     Fm3
    ...............................................................ccucuugcggauuugacguuau....................   46       0     Fm3
    ...............................................................cucuugcggauuugacguuau.....................   30       0     Fm3
    ...............................................................cucuugcggauuugacguuGu.....................   3        1     Fm3
    ...............................................................cucuugcggauuugacguuauU....................   23       1     Fm3

    ............................aaugucaaaccagAaagaggau......................................................   4        1     Fm2
    ...........................ugucaaaccagAaagaggaug.......................................................   9        1     Fm2
    ...........................ugucaaaccagAaagaggaugu......................................................   8        1     Fm2
```