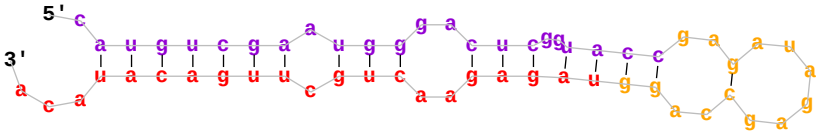
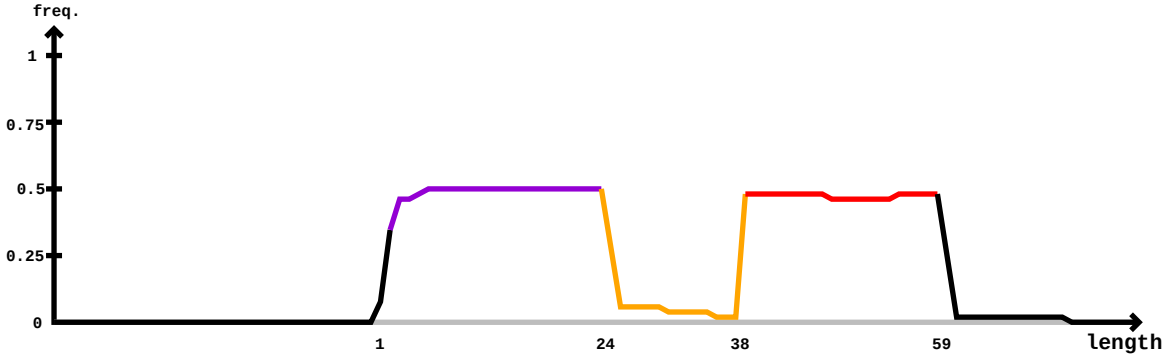
Provisional ID	:	NMRB01005750.1_Notospermus_geniculatus_strain_Ushimado_scaffold5750,_whole_genome_shotgun_sequence_47920
Score total	:	20.9
Score for star read(s)	:	3.9
Score for read counts	:	18.5
Score for mfe	:	-0.8
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	48
Mature read count	:	24
Loop read count	:	0
Star read count	:	24





Star Mature

```
5'- gagauagagccacguaaagaacugcuugacauacaugucgaaugggacucgguaccgagauagagccaggguagagaacugcuugacauacaugucaaaugggacuagguac  -3'   obs
    gagauagagccacguaaagaacugcuugacauacaugucgaaugggacucgguaccgagauagagccaggguagagaacugcuugacauacaugucaaaugggacuagguac        exp
    ....(((..(((.(((......)))(((((((..(((((((.(((..(((..((((..(.....)..)))))))..)))).)))))))..)))))))..)))).))).....   reads   mm   sample
    ......................................................................uagagaacugcuugacauaca....................   4       0    Mm3

    ................................caugucgaaugggacucAguacc...............................................   1       1    Mm1
    ................................caugucgaaugggacucgguacc...............................................   3       0    Mm1
    ......................................................................uagagaacugcuugacauaca....................   10      0    Mm1

    ..............................acaugucgaaugggacucgguaccgagaua..........................................   1       0    Mm2
    ...................................gucgaaugggacucgguaccgagauagagcc.....................................   1       0    Mm2
    ..................................................................uagagccagguagagaacu...............   1       0    Mm2
    ......................................................................uagagaacugcuugacauacC....................   1       1    Mm2
    ......................................................................uagagaacugcuugacauaca....................   1       0    Mm2
    ..................................................................................aAacaugucaaaugggac.......   1       1    Mm2

    ..............................acaugucgaaugggacucgguacc...............................................   3       0    Fm3
    ................................augucgaaugggacucgguacc...............................................   1       0    Fm3
    ....................................ucgaaugggacucgguaccgaga...........................................   1       0    Fm3
    ......................................................................uagagaacugcuugacauaca....................   2       0    Fm3

    ..............................caugucAaaugggacucgguacc................................................   10      1    Fm2

    ................................augucgaaugggacucgguacc...............................................   5       0    Fm1
    ......................................................................uagagaacugcuugacauacU....................   6       1    Fm1
```