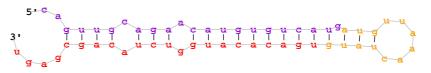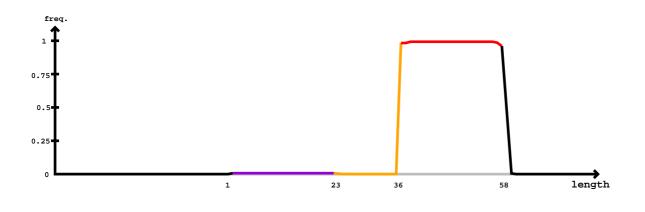|		
Provisional ID	:	NMRB01002238.1_Notospermus_geniculatus_strain_Ushimado_scaffold2238,_whole_genome_shotgun_sequence_36957
Score total	:	846
Score for star read(s)	:	3.9
Score for read counts	:	840.8
Score for mfe	:	1.9
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	1661
Mature read count	:	1649
Loop read count	:	0
Star read count	:	12


```
                                        Star                Mature
5'- uugaugcaugcgguauagauacugucacgacgcaccaguugcagaacaugugucaugauguuaaacuaugugacacauggcuacagcgaguguguugggcaggguuucaa  -3'   obs
    uugaugcaugcgguauagauacugucacgacgcaccaguugcagaacaugugucaugauguuaaacuaugugacacauggcuacagcgaguguguugggcaggguuucaa        exp
    ...(((((.....))))(((((.(((((.((((((((..((((.(((.((((((((((.(((......))))))))))))).))).))))..))))))))))))).))))...   reads   mm    sample
```

sequence	reads	mm	sample
....Cugacacauggcuacagcgagu...................	1	1	Fm3
....ugacacauggcuacagcga.....................	1	0	Fm3
....ugacacauggcuacagcgU.....................	2	1	Fm3
....ugacacauggcuacagGga.....................	1	1	Fm3
....ugacacauggcuacaCcga.....................	1	1	Fm3
....ugacacauggcuacagcgag....................	17	0	Fm3
....ugacacauggcuacagAgag....................	2	1	Fm3
....ugacacauggcuacagUgagu...................	5	1	Fm3
....uCacacauggcuacagcgagu...................	1	1	Fm3
....ugacacauggcuacagGgagu...................	2	1	Fm3
....ugacacauggcuacagAgagu...................	8	1	Fm3
....ugacacauggcuacagcgagA...................	1	1	Fm3
....ugacacauggcuacagcgagu...................	758	0	Fm3
....ugacacauggcuacagcUagu...................	1	1	Fm3
....ugacacauggCcuacagcgagu..................	1	1	Fm3
....ugacacauggcuacaUcgagu...................	1	1	Fm3
....ugacacauggcuacaAcgagu...................	3	1	Fm3
....ugacacauggcuacagcgaguU..................	32	1	Fm3
....................................caguugcagaacaugugucaug......................................	1	0	Mm3
....................................caguugcagaacaugugucaugU.....................................	1	1	Mm3
....ugacacauggcuacagcga.....................	1	0	Mm3
....ugacacauggcuacagcgG.....................	1	1	Mm3
....ugacacauggcuacagcgag....................	4	0	Mm3
....ugacacauggcuacCgcgagu...................	1	1	Mm3
....ugacacauggcuacagAgagu...................	1	1	Mm3
....ugacacauggcuacagUgagu...................	2	1	Mm3
....ugacacauggcuacagcgagu...................	106	0	Mm3
....ugacacauggcuacaAcgagu...................	1	1	Mm3
....ugacacauggcuacagGgagu...................	1	1	Mm3
....ugacacauggcuacagcgaguU..................	6	1	Mm3
....ugacacauggcuacagcgaguA..................	1	1	Mm3
....acacauggcuacagcgagu...................	1	0	Mm3

Star   **Mature**

uugaugcaugcgguauagauacugucacgacgcac caguugcagaacaugugucaug auguuaaacuaug ugacacauggcuacagcgagu guguugggcaggguuucaa

Sequence	Count	MM	Label
....ugacacauggcuacagcga....................	4	0	Fm1
....ugacacauggcuacagcgag...................	2	0	Fm1
....ugacacauggcuacagUgagu..................	1	1	Fm1
....ugacacauggcuacagcgagu..................	159	0	Fm1
....ugacacauggcuacagcgagG.................	1	1	Fm1
....ugacacauggcuacagGgagu.................	1	1	Fm1
....ugacacauggcuacagcGgu..................	1	1	Fm1
....ugacacauggcuacagcgaguU................	2	1	Fm1
.................................caguugcagaacaugugucaug...	5	0	Fm2
.................................caguugcagaacauguCucaug...	1	1	Fm2
....ugacacauUgucuacagcgagu..................	1	1	Fm2
....ugacacauggcuacagGgagu.................	1	1	Fm2
....ugacacauggcuacagcgagu..................	99	0	Fm2
....ugacacauggcuacagcgagC.................	1	1	Fm2
....ugacacauggcuacagUgagu..................	1	1	Fm2
.................................caguugcagaacaugugucaug...	4	0	Mm2
....ugacacauggcuacagcgag...................	2	0	Mm2
....ugacacauUgucuacagcgagu..................	1	1	Mm2
....ugacacauggcuacaAcgagu..................	1	1	Mm2
....ugacacauggcuacagcgagu..................	292	0	Mm2
....ugaAacauggcuacagcgagu..................	1	1	Mm2
....ugacacauggcuacagGgagu..................	1	1	Mm2
....ugacaUauggcuacagcgagu..................	1	1	Mm2
....ugacacauggcuacagAgagu..................	3	1	Mm2
....ugacacauggcuacagcgaguU.................	10	1	Mm2
....ugacacauggcuacagcgaguA.................	1	1	Mm2
....ugacacauggcuacagcgaguAu................	2	1	Mm2
....ugacacauggcuacagcgaguUu................	6	1	Mm2
....acacauggcuacagcgaguUu................	1	1	Mm2
....ugacacauggcuacagcgag...................	16	0	Mm1
....ugacacauggcuacaAcgagu..................	1	1	Mm1
....ugacacauggcuacagcgagu..................	50	0	Mm1
....ugacacauggcuacagAgagu..................	1	1	Mm1
....ugacacauggcuacagcgaguU.................	7	1	Mm1
....acacauggcuacagcgagu..................	13	0	Mm1
....acacauggcuacagcGgu..................	1	1	Mm1