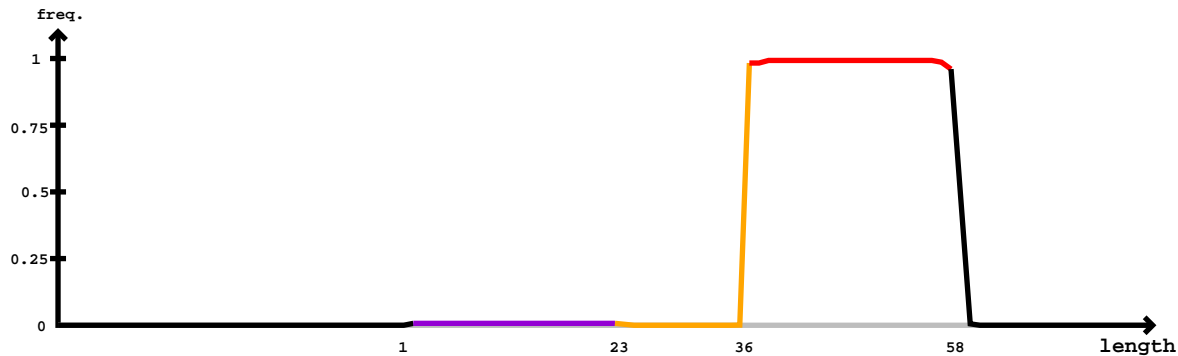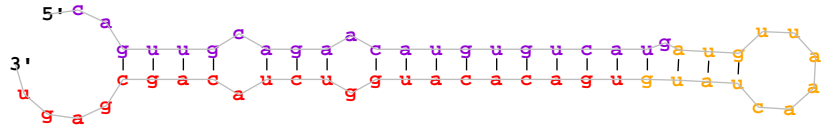| Provisional ID	: NMRB01001155.1_Notospermus_geniculatus_strain_Ushimado_scaffold1155,_whole_genome_shotgun_sequence_25652
Score total	: 846
Score for star read(s)	: 3.9
Score for read counts	: 840.8
Score for mfe	: 1.9
Score for randfold	: 0
Score for cons. seed	: -0.6
Total read count	: 1661
Mature read count	: 1649
Loop read count	: 0
Star read count	: 12

Star   Mature

```
5'- uugaugcaugcgguauagauacugucacgacgcaccaguugcagaacaugugucaugauguuaaacuaugugacacauggcucacagcgagugucauggguugggcagggguuucaac  -3'   obs
    uugaugcaugcgguauagauacugucacgacgcaccaguugcagaacaugugucaugauguuaaacuaugugacacauggcucacagcgagugucauggguugggcagggguuucaac        exp
    ...(((((.....))))((((.(((((.(((((((((..(((((.(((.((((((((((((..(((.......)))))))))))))).))).)))).))))))))))))).))))....     reads   mm    sample
```

Sequence	reads	mm	sample
...............................caguugcagaacaugugucaug..	1	0	Mm3
...............................caguugcagaacaugugucaugU...	1	1	Mm3
....ugacacauggcucacagcgG.....................	1	1	Mm3
....ugacacauggcucacagcga.....................	1	0	Mm3
....ugacacauggcucacagcgag....................	4	0	Mm3
....ugacacauggcucacagUgagu...................	2	1	Mm3
....ugacacauggcucacagcgagu...................	106	0	Mm3
....ugacacauggcucacCgcgagu...................	1	1	Mm3
....ugacacauggcucacaAcgagu...................	1	1	Mm3
....ugacacauggcucacagAgagu...................	1	1	Mm3
....ugacacauggcucacagGgagu...................	1	1	Mm3
....ugacacauggcucacagcgaguA..................	1	1	Mm3
....ugacacauggcucacagcgaguU..................	6	1	Mm3
....acacauggcucacagcgagu...................	1	0	Mm3
....Cugacacauggcucacagcgagu...................	1	1	Fm3
....ugacacauggcucacagcga.....................	1	0	Fm3
....ugacacauggcucacagcgU.....................	2	1	Fm3
....ugacacauggcucacagGga.....................	1	1	Fm3
....ugacacauggcucacaCcga.....................	1	1	Fm3
....ugacacauggcucacagcgag....................	17	0	Fm3
....ugacacauggcucacagAgag....................	2	1	Fm3
....ugacacauggcucacaAcgagu...................	3	1	Fm3
....ugacacauggcucacagcUagu...................	1	1	Fm3
....ugacacauggcucacagAgagu...................	8	1	Fm3
....ugacacauggcucacagcgagA..................	1	1	Fm3
....ugacacauggcucacagGgagu...................	2	1	Fm3
....ugacacauggcucacagcgagu...................	758	0	Fm3
....ugacacauggcucacagUgagu...................	5	1	Fm3
....uCacacauggcucacagcgagu...................	1	1	Fm3
....ugacacauggCcuacagcgagu...................	1	1	Fm3
....ugacacauggcucacaUcgagu...................	1	1	Fm3
....ugacacauggcucacagcgaguU..................	32	1	Fm3

uugaugcaugcgguauagauacugucacgacgcaccaguugcagaacaugugucaugauguuaaacuaugugacacauggucuacagcgaguguguugggcagggguuucaac

....caguugcagaacaugugucaug..	4	0	Mm2
....ugacacauggucuacagcgag.....................	2	0	Mm2
....ugaAacauggucuacagcgagu....................	1	1	Mm2
....ugacacauggucuacaAcgagu....................	1	1	Mm2
....ugacacauggucuacagcgagu....................	292	0	Mm2
....ugacacauggucuacagGgagu....................	1	1	Mm2
....ugacacauUgucuacagcgagu....................	1	1	Mm2
....ugacaUauggucuacagcgagu....................	1	1	Mm2
....ugacacauggucuacagAgagu....................	3	1	Mm2
....ugacacauggucuacagcgaguU...................	10	1	Mm2
....ugacacauggucuacagcgaguA...................	1	1	Mm2
....ugacacauggucuacagcgaguUu..................	6	1	Mm2
....ugacacauggucuacagcgaguAu..................	2	1	Mm2
....acacauggucuacagcgaguUu..................	1	1	Mm2
....ugacacauggucuacagcgag.....................	16	0	Mm1
....ugacacauggucuacaAcgagu....................	1	1	Mm1
....ugacacauggucuacagAgagu....................	1	1	Mm1
....ugacacauggucuacagcgagu....................	50	0	Mm1
....ugacacauggucuacagcgaguU...................	7	1	Mm1
....acacauggucuacagcgagu....................	13	0	Mm1
....acacauggucuacagcgGgu....................	1	1	Mm1
....ugacacauggucuacagcga......................	4	0	Fm1
....ugacacauggucuacagcgag.....................	2	0	Fm1
....ugacacauggucuacagUgagu....................	1	1	Fm1
....ugacacauggucuacagcGgu....................	1	1	Fm1
....ugacacauggucuacagGgagu....................	1	1	Fm1
....ugacacauggucuacagcgagu....................	159	0	Fm1
....ugacacauggucuacagcgagG...................	1	1	Fm1
....ugacacauggucuacagcgaguU...................	2	1	Fm1
...............................caguugcagaacauguCucaug..	1	1	Fm2
..............................caguugcagaacaugugucaug..	5	0	Fm2
....ugacacauggucuacagcgagC...................	1	1	Fm2
....ugacacauggucuacagcgagu....................	99	0	Fm2
....ugacacauggucuacagGgagu....................	1	1	Fm2
....ugacacauggucuacagUgagu....................	1	1	Fm2
....ugacacauUgucuacagcgagu....................	1	1	Fm2