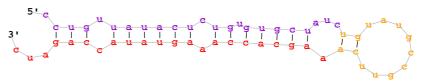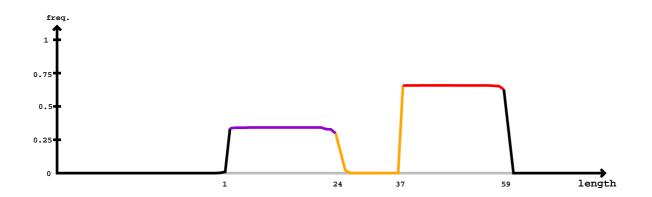| Provisional ID	: NMRB01001254.1_Notospermus_geniculatus_strain_Ushimado_scaffold1254,_whole_genome_shotgun_sequence_26708
Score total	: 1221.8
Score for star read(s)	: 3.9
Score for read counts	: 1222.2
Score for mfe	: -7.2
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 2409
Mature read count	: 1585
Loop read count	: 0
Star read count	: 824


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

sequence	reads	mm	sample
...............................cuccuguuauacucugugugc...	1	0	Mm2
...............................cuccuguuauacucugugugcu..	4	0	Mm2
................................uccuguuauacucugugugcu..	2	0	Mm2
................................uccuguuauacucugugugcuau..	3	0	Mm2
................................uccuguuauacucugugugcuauc...	1	0	Mm2
................................uccuguuauacucugugugcuauU...	2	1	Mm2
.................................ccuguuauacucugugugcu...	5	0	Mm2
.................................ccuguuauacucugugugcua..	3	0	Mm2
.................................ccuguuauacucugugUcuau...	1	1	Mm2
.................................ccuguuauacucugugugcuau...	49	0	Mm2
.................................ccuguuauacucugugAgcuauc..	1	1	Mm2
.................................ccuguuauacucugugugcuauA..	1	1	Mm2
.................................ccuguuauacucugugGgcuauc..	1	1	Mm2
.................................ccuguuauacAcucugugcuauc..	1	1	Mm2
.................................ccuguuauacucugugUAcuauc..	4	1	Mm2
.................................ccuguuauacucugugugcuauU..	4	1	Mm2
.................................ccugAuauacucugugugcuauc..	1	1	Mm2
.................................ccuguuauacucugugugcuauc..	237	0	Mm2
.................................ccuguuauacucugugugcuaucA...	2	1	Mm2
.................................ccuguuauacucugugugcuaucu...	172	0	Mm2
.................................ccuguuauacucuAugugcuaucu...	1	1	Mm2
.................................ccuguuauacucugugGgcuaucu...	1	1	Mm2
.................................ccuguuauacucugugugAcuaucu..	1	1	Mm2
.................................ccuguuauacucugugugcuaucG...	2	1	Mm2
.................................ccuguuauacucugugUcuaucu...	1	1	Mm2
.................................ccuguuauacucugugCgcuaucu...	2	1	Mm2
.................................ccuguuauacucugugugcuaucug..	23	0	Mm2
.................................ccuguuauacucugugGgcuaucug..	1	1	Mm2
.................................ccuguuauacucugugugcuaucuguaugcc..	1	0	Mm2
..................................cuguuauacucugugugcuauc..	3	0	Mm2
..................................cuguuauacucugugugcuaucu...	1	0	Mm2
..................................cuguuauacucugugugcuaucug..	2	0	Mm2
....uaugccguucaaagcaccaaag..	1	0	Mm2

Reference: aagaucuugcuuuuacuguccuugccacgugucu**ccuguuauacucugugugcuauc**uguaugccguuca**aagcaccaaaguauaccagauc**acaaguucacugcacuugcu

Sequence	Count	Mismatch	Label
....Caagcaccaaaguauaccagauc....................	1	1	Mm2
....aagcaccaaaguauaccaga.....................	4	0	Mm2
....aagcaccaaaguauaccagau....................	21	0	Mm2
....aagcaccaaaguauaccagauc...................	796	0	Mm2
....aagcaccaaaguaCaccagauc...................	1	1	Mm2
....aagcaUcaaaguauaccagauc...................	1	1	Mm2
....aagcaccaaaguauaccagauU..................	4	1	Mm2
....aagUaccaaaguauaccagauc...................	1	1	Mm2
....aagGaccaaaguauaccagauc...................	1	1	Mm2
....aGgcaccaaaguauaccagauc...................	1	1	Mm2
....aagcaccaaaAuauaccagauc...................	1	1	Mm2
....aagcaccaaaCuauaccagauc...................	1	1	Mm2
....Uagcaccaaaguauaccagauc...................	1	1	Mm2
....aagcaAcaaaguauaccagauc...................	1	1	Mm2
....aagcaccaaaguauaGcagauc...................	1	1	Mm2
....aaUcaccaaaguauaccagauc...................	1	1	Mm2
....aagcaccaaaguauaccagUuc...................	1	1	Mm2
....aagcaccaaaguauaccGgauc...................	2	1	Mm2
....aagcaccaaaguaGaccagauc...................	1	1	Mm2
....aagcacUaaaguauaccagauc...................	1	1	Mm2
....aagcaccaaaguauaccUgauc...................	1	1	Mm2
....aagcaccaaaguauaccCgauc...................	4	1	Mm2
....aagcaccaaaguauacAagauc...................	2	1	Mm2
....aagcaccaaaguauaccagGuc...................	1	1	Mm2
....aagcaccaaaUuauaccagauc...................	2	1	Mm2
....aagcaccaaaguauaccagauca..................	1	0	Mm2
....aagcaccaaaguauaccagaucU..................	9	1	Mm2
..............cuccuguuauacucugugugcu...	1	0	Mm1
....................ccuguuauacucugugugcuauc...	47	0	Mm1
....................ccuguuauacucugAgugcuauc...	1	1	Mm1
....................ccuguuauacGcugugugcuauc...	1	1	Mm1
....................ccuguuauacucugugUcuauc..	1	1	Mm1
....................ccuguuauacucugugugcuaucu..	13	0	Mm1
....................ccuguuauacucugugUccuaucu..	1	1	Mm1
....................ccugAuauacucugugugcuaucu..	1	1	Mm1
........................uuauacucugugugcuaucug...	4	0	Mm1
....aagcaccaaaguauaccagau...................	11	0	Mm1
....aagcaccaaaguauacAagauc..................	1	1	Mm1
....aagcaccaaaguauaccagauc..................	38	0	Mm1
...................uccuguuauacucugugugcG...	1	1	Fm2
...................uccuguuauacucugugugcu...	6	0	Fm2
....................ccuguuauacucugugUcuauc..	1	1	Fm2
....................ccuguuauacucugugugcuauc...	44	0	Fm2
....................ccuguuauacucugugugcuaucug...	10	0	Fm2
.....................Nuguuauacucugugugcuauc...	1	1	Fm2
.....................cuguuauacucugugugcuauc...	5	0	Fm2
....aagcaccaaaguauacUagauc..................	1	1	Fm2
....aagcaccaaaguauacAagauc..................	2	1	Fm2
....aagcaccaaaguauaccagauc..................	142	0	Fm2
....aagcaccaaaAuauaccagauc..................	1	1	Fm2
....aagcaccaaaguauaccUgauc..................	1	1	Fm2
....Uagcaccaaaguauaccagauc..................	1	1	Fm2
....................ccuguuauacucugugugcuau..	5	0	Fm1
....................ccuguuauacucugugugcuauc...	8	0	Fm1
....................ccuguuauacucugugugcuaucu..	3	0	Fm1
....................ccuguuauacucugugugcuaucug...	6	0	Fm1
....aagcaccaaaguauaccagau...................	8	0	Fm1
....aagcaccaaaguauaccGgauc..................	1	1	Fm1
....aagcaccaaaguauacAagauc..................	1	1	Fm1
....aagcaccaaaguauaccUgauc..................	1	1	Fm1
....aagcaccaaaguauaccagauc..................	120	0	Fm1
....aagcaccaaaguauaccagaucU.................	4	1	Fm1
....................ccuguuauacucugugugcu...	1	0	Mm3
....................ccuguuauacucugugugcua..	1	0	Mm3
....................ccuguuauacucugugugcuau.uguaugccguucaaagcaccaaaguauaccagaucacaaguucacugcacuugcu........	9	0	Mm3
....................ccuguuauacucugugugcuauc..	41	0	Mm3

aagaucuugcuuuuacuguccuugccacgugucuccguguauacucugugugcuaucuguaugccguucaaagcaccaaaguauaccagaucacaaguucacugcacuugcu

Sequence	Count	Mismatch	Sample
......ccuguuauacucugugugcuauU.......	2	1	Mm3
......ccuguuauacucugugugUuauc.......	1	1	Mm3
......ccuguuauacucugugugUcuauc......	1	1	Mm3
......ccuguuauacucugugugGgcuauc.....	1	1	Mm3
......ccuguuauacucugugugcuaucG......	1	1	Mm3
......ccuguuauacucugugugcuaucu......	33	0	Mm3
......ccuguuauacucugugugcuaucug.....	4	0	Mm3
......cuguuauacucugugugcuaucu.......	1	0	Mm3
......aagcaccaaaguauaccagau........	5	0	Mm3
......aagcaccaaaguauacAagauc.......	1	1	Mm3
......aagcaccaaaguauaccagauc.......	200	0	Mm3
......aagcaccaaaguauaccUgauc.......	1	1	Mm3
......Nagcaccaaaguauaccagauc.......	1	1	Mm3
......aagcaccaCaguauaccagauc.......	1	1	Mm3
......aagcaccaaaguauaccagaucU......	5	1	Mm3
......aagcaccaaaguauaccagauca......	1	0	Mm3
......agcaccaaaguauaccagauc........	1	0	Mm3
......accaaaguauaccagauc..........	1	0	Mm3
......uccuguuauacucugugugcu........	2	0	Fm3
......ccuguuauacucugugugcu.........	6	0	Fm3
......ccuguuauacucugugugcua........	2	0	Fm3
......ccuguuauacucugugugcuau.......	6	0	Fm3
......ccuguuauacucugugugcuauc......	15	0	Fm3
......ccuguuauacucugugugcuauU......	1	1	Fm3
......ccuguuauacucugugugcuaucu.....	3	0	Fm3
......aagcaccaaaguauaccag.........	8	0	Fm3
......aagcaccaaaguauaccagau.......	16	0	Fm3
......aagcaccaaaguauaccagauc......	143	0	Fm3
......aagcaccaaaAuauaccagauc......	1	1	Fm3
......aagcaccaaaguauaccagauA......	1	1	Fm3
......aagcaccaaaguauaccagaucU.....	3	1	Fm3
......aagcaccaaaguauaccagauca.....	1	0	Fm3

aagaucuugcuuuuacuguccuugccacgugucuccguguauacucugugugcuaucuguaugccguucaaagcaccaaaguauaccagaucacaaguucacugcacuugcu