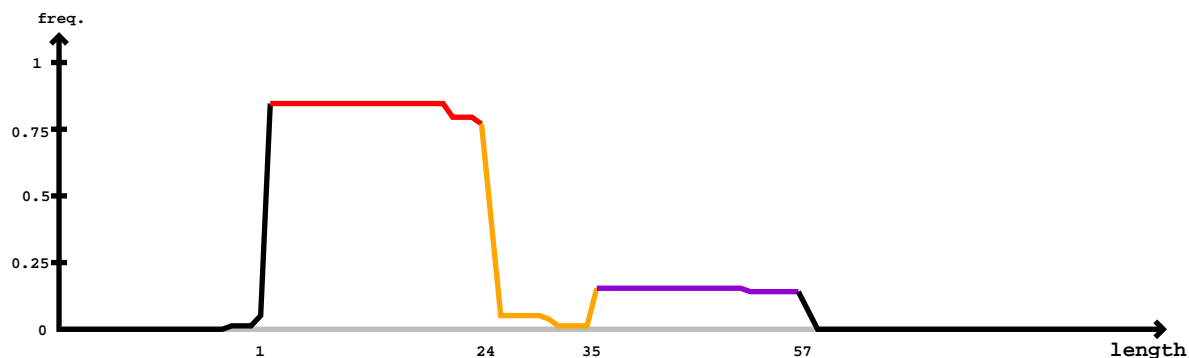


Provisional ID	:	NMRB01000597.1_Notospermus_geniculatus_strain_Ushimado_scaffold597_whole_genome_shotgun_sequence_16652
Score total	:	31.5
Score for star read(s)	:	3.9
Score for read counts	:	30.7
Score for mfe	:	-2.4
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	72
Mature read count	:	61
Loop read count	:	0
Star read count	:	11



Mature

5'	obs	exp	reads	mm	sample
uucuguuuauuuuacuuc <u>ugagauccgagucgagacgagaaa</u> uaaagcauuuu <u>uacguauacgucgagucucuuu</u> caggccuggaagaaggagcaguggggcaugcauuua	-3'				
uucuguuuauuuuacuuc <u>ugagauccgagucgagacgagaaa</u> uaaagcauuuu <u>uacguauacgucgagucucuuu</u> caggccuggaagaaggagcaguggggcaugcauuua					
..(((.((((((((((.(.((((((.(.....(((.((((.(.....)))))).)))).)))).)))).(((((.....)))..)))))..)))).					
.....ugagauccgagucgagacgagaa.....	1	0			Fm1
.....uacguauacgucgagucucuuu.....	4	0			Fm1
.....uacguauacgucgagucucuuu.....	6	0			Fm2
.....uucaugagauccAagucgagacg.....	1	1			Fm3
.....augagauccgagucgagacg.....	3	0			Fm3
.....ugagauccgagucgagacgagaa.....	2	0			Fm3
.....ugagauccgagucgagacgagaa.....	47	0			Fm3
.....ugagauccgagucgagaaAgagaa.....	1	1			Fm3
.....ugagauccgagucgagacgagaaa <u>uaaag</u>	1	0			Fm3
.....ugagauccgagucgagacgagaaa <u>uaaagc</u>	1	0			Fm3
.....ugagauccgagucgagacgagaa.....	2	0			Mm2
.....ugagauccgagucgGgacgagaa.....	4	1			Mm2
.....ugagauccgagucgagacgagaaa <u>uaaagc</u>	2	0			Mm2
.....auuuuacguauacgucgaguu.....	1	0			Mm2
.....ugagaucUgagucgagacgagaa.....	1	1			Mm3
.....uacguauacgucgagucucuuu.....	1	0			Mm3