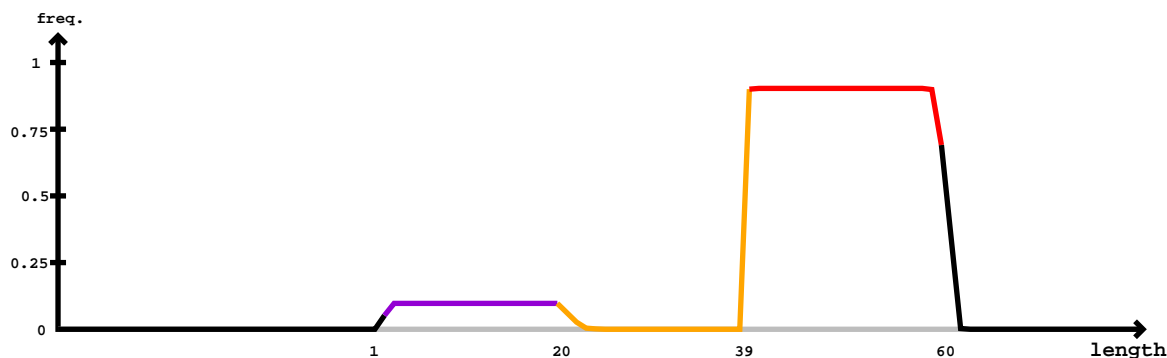


Provisional ID	:	NMRB01008102.1_Notospermus_geniculatus_strain_Ushimado_scaffold8102,_whole_genome_shotgun_sequence_49212
Score total	:	361.8
Score for star read(s)	:	3.9
Score for read counts	:	356.5
Score for mfe	:	2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	711
Mature read count	:	642
Loop read count	:	0
Star read count	:	69



Star

[illegible]

## Star

## Mature

cuucaaguuggaggauaucugcgugcgcggaucuggucacaaacuguguc	ccaaauccaucuaaaauugugacacaguuuguuacgagua	cucgcguccgcgauugcggau			
.....ugacacaguuuguuacgagua.....	31	0	Fm2		
.....ugacacaguuuguuacgaguac.....	47	0	Fm2		
.....ugacacaguuuguuacgaguaA.....	9	1	Fm2		
.....ugacacaguuuguuacgaguaU.....	24	1	Fm2		
.....acuggucacaaacuguguc.....	2	0	Mm3		
.....cuggucacaaacuguguc.....	1	0	Mm3		
.....ugacacaguuuguuacgagu.....	7	0	Mm3		
.....ugacGcaguuuguuacgagu.....	1	1	Mm3		
.....ugacacaguuuguuacgagua.....	7	0	Mm3		
.....ugacacaguuuguuacgaguac.....	9	0	Mm3		
.....ugacacaguuuguuacgaguaU.....	4	1	Mm3		
.....gacacaguuuguuacgaguacu.....	1	0	Mm3		
.....acuggucacaaacugugucc.....	2	0	Fm3		
.....acuggucacaaacugugAcc.....	1	1	Fm3		
.....acuggucacaaacuguguccc.....	11	0	Fm3		
.....cuggucacaaacuguguc.....	6	0	Fm3		
.....cuggucacaaacuguguccaa.....	1	0	Fm3		
.....ugacacaguuuguuacgag.....	3	0	Fm3		
.....ugacacaguuGguuacgagu.....	1	1	Fm3		
.....ugacacaguuuguuacUagu.....	1	1	Fm3		
.....ugacacaguuuguuacgagu.....	100	0	Fm3		
.....ugacacaguuuguuacGgu.....	1	1	Fm3		
.....ugacacaguuuguuacUgagu.....	1	1	Fm3		
.....ugacacaguuuguuacGggu.....	1	1	Fm3		
.....ugacacaguuuguuacgUgua.....	1	1	Fm3		
.....ugacacaguuuguuacgGgua.....	1	1	Fm3		
.....ugacacaguuuguuacgagua.....	101	0	Fm3		
.....ugacacaguuAgguuacgagua.....	1	1	Fm3		
.....ugacacaguuuguuacgaguU.....	7	1	Fm3		
.....ugacacaguuuguuacgaguaU.....	19	1	Fm3		
.....ugacacaguuuguuacgaguac.....	68	0	Fm3		
.....ugacacaguuuguuacgGguac.....	1	1	Fm3		
.....ugacacaguuuguuacgUguac.....	1	1	Fm3		
.....gacacaguuuguuacgaguacu.....	1	0	Fm3		