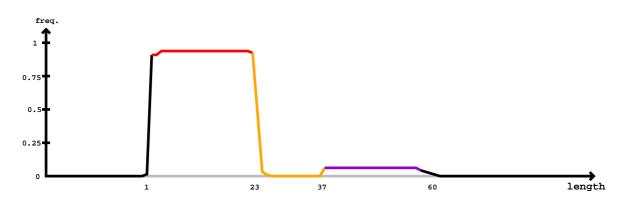
$: NMRB01000231.1_Notospermus_geniculatus_strain_Ushimado_scaffold231,_whole_genome_shotgun_sequence_7797$ Provisional ID

3.4 Score total

Score for star read(s) -1.3 Score for read counts 0 Score for mfe 1.7 Score for randfold 0 Score for cons. seed 3

Total read count 209 196 Mature read count Loop read count 0 Star read count 13



Mature	Star
5'- ucguucccccugagaggugaucagcuguaccgaugccuucuuguccucaa	uguuacaagaagucauuggacggcucaugaauucuuggguaaagucgccaguggaauguuua

.(((((((((((((((((((((((((((((((((((((reads 11 2 1 33	mm 0 1 1	sample Fm2 Fm1 Fm1
.ucagcuguacAgaugccuucuu	2 1 33 1	1	Fm1
	1 33 1	1	
	1 33 1	1	
ucagcuquaccgaugccCucuu	33 1		Fm1
	1	0	
ucagcuguaccgaugccuucuu			Fm1
.ucagcuguaccgaugcAuucuu		1	Fm1
agcuguaccgaugccuucuugucagcuguaccgaugccuucuuguc	2	0	Fm1
.aagaagucauuggacAgcuca	1	1	Fm1
.ucagcuguaccgaugccuucuu	54	0	Mm1
.ucagcuguacAgaugccuucuu.	1	1	Mm1
aagaagucauuggacAgcuc	4	1	Mm1
aagaagucauuggacAgcuca.	6	1	Mm1
ucagcuguaccgaugccuucu	1	0	Mm2
ucagcuguaccgaugccuucuu	43	0	Mm2
ucagcuguaccgaugccGucuu	1	1	Mm2
ucagcuguaccgaugccuucuuC	2	1	Mm2
ucagcuguaccgaugccuucuug	8	0	Mm2
ucagcuguaccgaugccuucuugu	2	0	Mm2
agcuguacegaugccuucuugu	1	0	Mm2
aagaagucauuggacggcuca	1	0	Mm2
aucagcuguaccgaugccuucu	1	0	Fm3
aucagcuguaccgaugchuucuaucagcuguaccgaugchuucu	1	1	Fm3
	1	1	Fm3
ucageuguacagaugeeuucuuucageuguacegaugeeuucuu	15	0	Fm3
ucageuguacegaugeeuucuuucageuguacegaugeeuucuu	1	1	Fm3
ucagcuguacGgaugccuucuu	1	1	Fm3
ucagcuguacegaugccuucuug	5	0	Fm3
ucagcuguaccgaugccuucuugG	1	1	Fm3
agcuquaccgaugccuucuug.	3	0	Fm3
ageuguacegaugeeuucuug	-	1	Fm3

Mature Star

ucguuccccugagaggugaucagcuguaccgaugccuucuuguccucaauguuacaagaagucauuggacggcucaugaauucuuggguaaagucgccaguggaauguuua

aucageuguacegaugceuueuuUu	1	1	Mm3
ucagcuguaccgaugccuucuu	2	0	Mm3
nesacriaris ecastracemicina	1	0	Mm 3