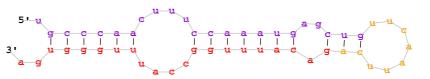
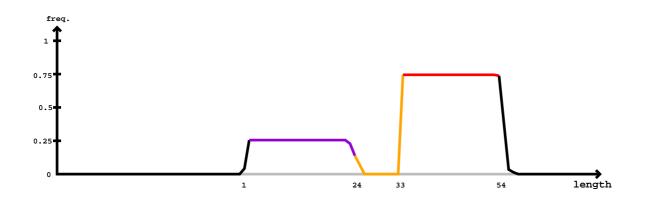
Provisional ID: NMRB01000076.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold76,\_whole\_genome\_shotgun\_sequence\_2799

72.4 Score total Score for star read(s) 3.9 67.9 Score for read counts 1.2 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 145 108 Mature read count Loop read count 0 Star read count 37





5'- uccauuagaguugccaggaauauugcucccagauugguugcccaacuuuccaaaugagcuguucaauucagacauuuggccauuugggugacuguuucgggguuuaaccug

Star

uccauuagaguugccaggaauauugcucccagauugguugcccaacuuuccaaaugagcug <mark>uucaauucagacauuugggcauuugggggac</mark> uguuuaggggguuuaaccug	exp	)	
((((((((.((((((((((((((((((((	reads	mm	sample
.gacauuuggccauuugggug	1	0	Mm3
.gacauuuggccauuuggtuga	1	1	Mm3
.gacauuuggccauuugggugG	1	1	Mm3
.gacauuuggccauuuggguga	14	0	Mm3
	1	1	Mm3
.gacauuuggccauuugggugac	1	0	Mm3
.gacauuuggccauuugggugaU	3	1	Mm3
.gacauuuggccUuuugggugac	1	1	Mm3
.gacauuuggccauuugggugaA.	1	1	Mm3
.gacauuuggccauuugggugacu	1	0	Mm3
ugcccaacuuuccaaaugagcu	3	0	Fm3
.ugcccaacuuuccaaaugagcug	4	0	Fm3
	1	0	Fm3
.gacauuuggccauuuggUuga.	1	1	Fm3
.gacauuuggccauuuggguga.	13	0	Fm3
.gacauuuggccauuugUgugac	1	1	Fm3
.gacauuuggccauuugggugac	1	0	Fm3
.gacauuuggccauuugggugaUu	1	1	Fm3
.gacauuuggccauuugggugacu	1	0	Fm3
.gacauuuggccauuugggugacuU	1	1	Fm3
	4	0	Mm2
.ugcccaacuuuccaaaugagcu	3	0	Mm2
.ugcccaacuuGccaaaugagcu	1	1	Mm2
.ugcccaacuuuccaaaugaAcug	3	1	Mm2
.ugcccaacuuuccaaaugagcug	5	0	Mm2
.gacauuuggccauuugggugG.	1	1	Mm2
	1	1	Mm2
	4	0	Mm2
	3	0	Mm2
gacauuuggccauuuggggugacuU	1	1	Mm2

Mature

Star Mature

uccauuagaguugccaggaauauugcucccagauugguugcccaacuuuccaaaugagcuguucaauucagacauuuggccauuugggugacuguuucgggguuuaaccug			
ugcccaacuuuccaaaugageug	6	0	Mm1
gacauuuggccauuuggguga	7	0	Mm1
ugcccaacuuuccaaaugaAcug	1	1	Fm2
gGcauuuggccauuuggguga	1	1	Fm2
gacauuuggcUauuuggguga	1	1	Fm2
gacauuuggccauuuggCuga	1	1	Fm2
gacauuuggccauuugggugG	4	1	Fm2
gacauuuggccauuuggguga	23	0	Fm2
gacauuuggccauuugggugac	10	0	Fm2
gacauuuggccauuuggguga	5	0	Fm1
gacauuuggccauuugggugG	2	1	Fm1