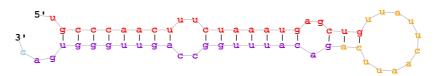
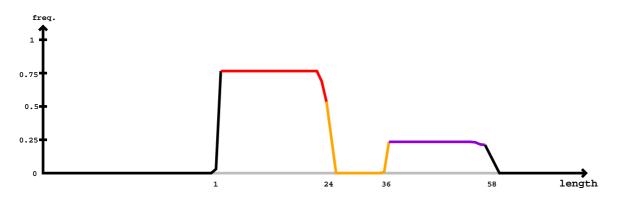
$Provisional\ ID \\ : NMRB01000076.1_Notospermus_geniculatus_strain_Ushimado_scaffold76, \\ _whole_genome_shotgun_sequence_2797 \\ : NMRB01000076.1_Notospermus_geniculatus_strain_sequence_2797 \\ : NMRB01000076.1_Notospermus_geniculatus_$

131.6 Score total Score for star read(s) 3.9 Score for read counts 126.6 1.8 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 260 199 Mature read count Loop read count 0 Star read count 61





Mature	Star

.....gacauuuggccaguugggugac.....

5'- aucggaguuuccaaagaacauugguccuggauuguugcccaacuuucuaaaugagcuguuauucaauucagacaguuuggccaguugggugacuguuucagguugagcucca

aucggaguuuccaaagaacauugguccuggauuguu <mark>gcccaacuuucuaaaugagcuguuauucaauuca</mark>	exp		
((((((((((((((((((((((((((((((((((((reads	mm	sample
uugcccaacuuucuaaaugagc	1	0	Fm3
gacauuuggccaguugggu	3	0	Fm3
gacauuuggccaguugggu	1	0	Fm1
gacauuuggccaguuggguga	6	0	Fm1
gacauuuggccaguugggugG	1	1	Fm1
gacauuuggccaguugggugacu	2	0	Fm1
uugcccaacuuucuaaaCgagc	1	1	Mm2
uuAcccaacuuucuaaaugagcu	1	1	Mm2
uugcccaacuuucuaaaugagcu	4	0	Mm2
	1	1	Mm2
uAcccaacuuucuaaaugagc	18	1	Mm2
uAcccaacuuucuaaaugagcu	35	1	Mm2
uAcccaacuuucuaaaugagcug.	136	1	Mm2
uAcccaacuuucuaaaugagcugu	2	1	Mm2
.agacauuuggccUguuggguga	1	1	Mm2
.agacauuuggccaguuggguga	1	0	Mm2
gacauuuggccaguuggg	1	0	Mm2
gacauuuggccaguugggug	1	0	Mm2
gacauuuggccaguugggugG	1	1	Mm2
gacauuuggccaguuggguga	10	0	Mm2
gacauuuggccaguugCguga	1	1	Mm2
gacauuuggccaguugggugaU	2	1	Mm2
gacauuuggccaguuggCugac	1	1	Mm2
gacauuuggccaguugggugacgacauuuggccaguugggugac	6	0	Mm2
gacauuuggccaguugggugacu	1	0	Mm2
Uacauuuggccaguuggguga	1	1	Mm1
gacauuuggccaguuggguga	12	0	Mm1
gacauuuggccaguugggugaU	4	1	Mm1