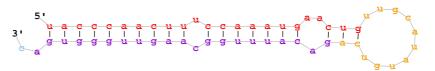
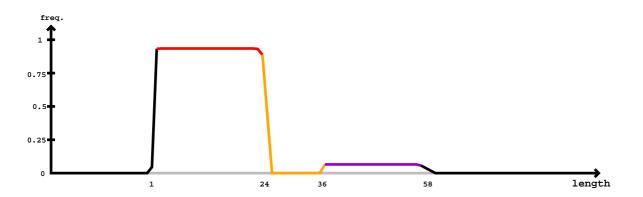
Provisional ID : NMRB01000004.1_Notospermus_geniculatus_strain_Ushimado_scaffold4,_whole_genome_shotgun_sequence_162

482.6 Score total Score for star read(s) 3.9 477.3 Score for read counts Score for mfe 2 Score for randfold 0 -0.6 Score for cons. seed Total read count 948 886 Mature read count Loop read count 0

62

Star read count





5'- ucacauuugccuagagcaguu<mark>acccaacuuuccaaaugaacug</mark>uugcauaugucagacauuuggcaaguugggugacuguuccaggauugaaaugucugacaguauaggggc

Mature	Star
Mature	ol.al

ucacauuugccuagagcaguuacccaacuuuccaaaugaacuguugcauaugucagacauuuggcaaguugggugacuguuccaggauugaaaugucugacaguauaggggc	exp		
(((((.(((((((((((((((((((((((((((((((reads	mm	sample
uuacccaacuuuccaaaCgaacug	1	1	Fm3
uuacccaacuuuccaaaAgaacug	1	1	Fm3
uuacccaacuuuccaaaugaacug	9	0	Fm3
uacccaacuuuccaaaugaac	1	0	Fm3
uacccaacuuuccaaaugaacu	8	0	Fm3
uaAccaacuuuccaaaugaacug	1	1	Fm3
uacccaacCuuccaaaugaacug	1	1	Fm3
uacccaacuuuccaaaGgaacug	1	1	Fm3
uacccaacuuuccaaauCaacug	1	1	Fm3
uUcccaacuuuccaaaugaacug	1	1	Fm3
uacccaacuuuccaaaugaacug	231	0	Fm3
uacccGacuuuccaaaugaacug	1	1	Fm3
uacccaacuuuccaaaugaacugu	4	0	Fm3
.acccaacuuuccaaaugaacug	1	0	Fm3
agacauuuggcaaguuggguga	2	0	Fm3
gacauuuggcaaguuggguga	3	0	Fm3
gacauuuggcaaguugggugU	1	1	Fm3
gacauuuggcaaguugggugac	10	0	Fm3
gacauuuggcaaguugggugaA	1	1	Fm3
gacauuuggcaaguugggugacu	5	0	Fm3
gacauuuggcaaguugggugacA	2	1	Fm3
uuacccaacuuuccaaaugaacu	1	0	Mm3
uuacccaacuuuccaaaugaacug	8	0	Mm3
uacccaacuuuccaaaugaacu	4	0	Mm3
uacccaacuuuccaaaugaacug	29	0	Mm3
Cacccaacuuuccaaaugaacug	1	1	Mm3
uacccaacuuAccaaaugaacug	1	1	Mm3
uacccaacuuuccaaaugaacugu	1	0	Mm3
gacauuuggcaaguugggug	1	0	Mm3
gacauuuggcaaguugggugac	1	0	Mm3
gacauuuggcaGguugggugac	1	1	Mm3
gacauuuggcaaguugggugacG	1	1	Mm3

Mature Sta

Mature S	Star Star			
ucacauuugccuagagcaguuacccaacuuuccaaaugaacuguugcauaugucaga	acauuuggcaaguugggugacuguuccaggauugaaaugucugacaguauaggggc			
ga	acauuuggcaaguugggugacu	3	0	Mm3
uacccaacuuuccaaaugaacug		22	0	Fm1
ga	acauuuggcaaguuggguga	4	0	Fm1
uuacccaacuuuccaaaugaacug		15	0	Fm2
uacccaacuuuccaaaugaacug		29	0	Fm2
CaacuuuccaaauCaacug		1	1	Fm2
uacccaacuuuccaaauAaacug		1	1	Fm2
uacccaacuuuccaaaugaacuU		1	1	Fm2
uGcccaacuuuccaaaugaacug		1	1	Fm2
ggaacug		1	1	Fm2
ga	acauuuggcaaguugggug	7	0	Fm2
uacccaacuuuccaaaugaacu		6	0	Mm1
UacccaacuuucUaaaugaacug		1	1	Mm1
uacccaacuuuccaaaCgaacug		1	1	Mm1
uacAcaacuuuccaaaugaacug		1	1	Mm1
uacccaacuuuccaaaugaacug		88	0	Mm1
ga	acauuuggcaaguugAguga	1	1	Mm1
ga	acauuuggcaaguuggguga	10	0	Mm1
uuacccaacuuuccaaaugaacu		2	0	Mm2
uuacccaacuuuccaaUugaacug		1	1	Mm2
uuacccaacuuuccaaauCaacug		1	1	Mm2
uuacccaacuuuccaaaugaacug		4	0	Mm2
uacccaacuuuccaaauCaac		1	1	Mm2
uacccaacuuuccaaaugaac		1	0	Mm2
uacccaacuuuccaaaugaacu		14	0	Mm2
uacccaacuuuccaaauCaacu		7	1	Mm2
UacccaacuuuccaaaugUacug		1	1	Mm2
Grant and the control of the co		1	1	Mm2
uacccaacuuuccaaauAaacug		2	1	Mm2
acccaacuuuccaaauCaacug		3	1	Mm2
uacccaacuuuccaaauUaacug		2	1	Mm2
Gaugaacug		1	1	Mm2
UacccaacuuuUcaaaugaacug		1	1	Mm2
uGcccaacuuuccaaaugaacug		3	1	Mm2
uacccaacuuuccaaaAgaacug		1	1	Mm2
acccaacuuuccaaaugaacug		352	0	Mm2
uUcccaacuuuccaaaugaacug		1	1	Mm2
gacccaacuuuccaaaCgaacug		4	1	Mm2
uacccaacGuuccaaaugaacug		1	1	Mm2
uaccAaacuuuccaaaugaacug		2	1	Mm2
uacccGacuuuccaaaugaacug		1	1	Mm2
UacccaacuuucUaaaugaacug		1	1	Mm2
gacccaacuuuccaaaugaacugu		2	0	Mm2
acccaacuuuccaaaugaacug		1	0	Mm2
aga	acauuuggcaaguugggugU	1	1	Mm2
ga	acauuuggcaaguuggguga	3	0	Mm2
ga	acauuuggcaaguugggugac	1	0	Mm2
ga	acauuuggcaaguugggugacA	1	1	Mm2

Mm2