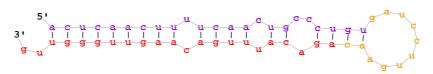
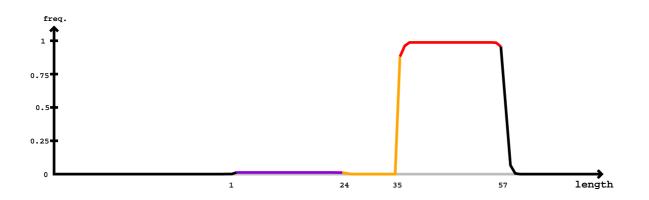
Provisional ID : NMRB01000004.1_Notospermus_geniculatus_strain_Ushimado_scaffold4,_whole_genome_shotgun_sequence_155

Score total 603 Score for star read(s) 3.9 598.7 Score for read counts Score for mfe 1.1 Score for randfold 0 Score for cons. seed -0.6 Total read count 1186 Mature read count 1171 Loop read count 0 Star read count 15



exp



5'- aacugaagacuaaagugaugggaguugguaauaucgacucaacuuuucaacugcccugugauccuugaacagacauuugacaaguuggguugauguucacaaaccaggcauc
aacugaagacuaaagugaugggaguugguaauaucgacucaacuuuucaacugcccugugauccuugaacagacauuugacaaguuggguugauguucacaaaccaggcauc

agacauuugacaaguugAguug......agacauuugacaaguugAguug.....

Star

(((.(((((((((((((((((reads	mm.	sample
acucaacuuuucaacugeecugu	1	0	Fm3
agacauuugacaaguugUguu	1	1	Fm3
agacauuugacaaguugguu	10	0	Fm3
agacauuugacaaguuggguuU	1	1	Fm3
agacauuugacaaguuggguug	54	0	Fm3
agacauuugacaaguuggAuuga	1	1	Fm3
agacauuugacaaguuggguuga	18	0	Fm3
agacauuugacaaguuggguugU	41	1	Fm3
agacauuugacaaguuggguugau	2	0	Fm3
gacauuugacaaguuggguug	2	0	Fm3
gacauuugacaaguuggguugU	2	1	Fm3
gacauuugacaaguuggguuga	14	0	Fm3
gacauuugacaaguuggguugah.	6	1	Fm3
gacauuugacaaguuggguugau	3	0	Fm3
.acucaacuuuucaacugcccugu.	4	0	Mm3

Mature

agacauuugacaaguuggguugau	2	0	Fm3
.gacauuugacaaguuggguug	2	0	Fm3
.gacauuugacaaguuggguugU	2	1	Fm3
gacauuugacaaguuggguuga	14	0	Fm3
gacauuugacaaguuggguugaA	6	1	Fm3
.gacauuugacaaguuggguugau	3	0	Fm3
acucaacuuuucaacugeccugu	4	0	Mm3
agacauuugacaaguuggguu	2	0	Mm3
agacauuugacaaguuggguuC	1	1	Mm3
agacauuugacaaguuggguug	10	0	Mm3
agacauuugacaaguuggguuU	1	1	Mm3
agacauuugacaaguuggguugU	45	1	Mm3
agacauuugacaaguuggguuga	3	0	Mm3
agacauuugacaaguuggguugaA	1	1	Mm3
agacauuugacaaguuggguugUu	3	1	Mm3
agacauuugacaaguuggguugau	2	0	Mm3
agacauuugacaaguuggguu	14	0	Fm2
agacauuugacaaUuuggguu	1	1	Fm2
agacauuugacaaguuggguug	180	0	Fm2
agacauuugacaaguuggguGg	1	1	Fm2
agacauuugacaaguugCguug	3	1	Fm2
agacauuugacaGguuggguugagacauuugacaGguuggguug	4	1	Fm2

Star Mature

Star Mature				
aacugaagacuaaagugaugggaguugguaauaucgacucaacuuuucaacugcccugugauccuugaacagacauuugacaaguuggguuga	auguucacaaaccaggcauc			
agacauuugacaaguuggguugl		94	1	Fm2
		4	1	Fm2
agacauuugacaGguuggguuga				
agacGuuugacaaguuggguuga	1	1	1	Fm2
agacauuugacaaguuggguugaagacauuugacaaguuggguuga	1	50	0	Fm2
agacauuugacaaguuggguug0	3	2	1	Fm2
agacauuugacaaguuggguuga	au	5	0	Fm2
agacauuugacaaguuggguugl	Ju	8	1	Fm2
gacauuugacaaguuggguug.		6	0	Fm2
gacauuugacaaguuggguuga		41	0	Fm2
		1	1	Fm2
gacauuugacaaguuggAuuga				
acauuugacaaguuggguuga		9	0	Fm2
acauuugacaaguuggguuga	AA	2	1	Fm2
agacauuugacaaguuggguCg		1	1	Fm1
agacauuugacaaguuCgguug.		1	1	Fm1
agacauuugacaaguugCguug.		1	1	Fm1
aAacauuugacaaguuggguug		1	1	Fm1
			0	
agacauuugacaaguuggguug.		62		Fm1
agacauuugacaaguuggguGg.		2	1	Fm1
agacauuugacaaguuggguugaagacauuugacaaguuggguuga	1	11	0	Fm1
agacauuugacaaguuggguugtagacauuugacaaguuggguugt	J	70	1	Fm1
agacauuugacaaguuggguug0agacauuugacaaguuggguug0	3	1	1	Fm1
agacauuugacaaguuggguugs	aA	2	1	Fm1
agacauuugacaaguuggguuga		3	0	Fm1
agacauuugacaaguuggguugl		4	1	Fm1
gacauuugacaaguuggguuga		8	0	Fm1
acauuugacaaguugggCuga	au	1	1	Fm1
acauuugacaaguuggguugaacauuugacaaguuggguuga	au	4	0	Fm1
acauuugacaaguuggguuga	auA	5	1	Fm1
agacauuugacaaguugggu		1	0	Mm1
agacauuugacaaguuggguug.		14	0	Mm1
agacauuugacaaguuggguugl		24	1	Mm1
agacauuugacaaguuggguugī		3	1	Mm1
agacaaagacaagagagag	,	3	-	Panta
		_	_	
		1	1	Mm2
acucaacuuuucaacugcccugu		9	0	Mm2
agacauuugacaaguugggu		1	0	Mm2
agacauuugacaaguuggguu		6	0	Mm2
agacauuugacaaguuggguuU		1	1	Mm2
agacauuugacaaguuUgguug.		1	1	Mm2
agacauuugacaaguugAguug.		1	1	Mm2
agacauuugCcaaguuggguug.		1	1	Mm2
				Mm2
agacauuugacaaguuggguug.		62	0	
agacauuugaUaaguuggguug		2	1	Mm2
agacauuugacaaguCggguug.		1	1	Mm2
agacauuugacaaguuggguug(3	2	1	Mm2
agacauuugacaaguuggguugl	J	170	1	Mm2
agacauuugacaaguuggguug	a	22	0	Mm2
agacauuugacaaguuggguugl	Ju	6	1	Mm2
agacauuugacaaguuggguuga		2	0	Mm2
agacauuugacaaguuggguuga		1	1	Mm2
gacauuugacaaguuggguuga		13	0	Mm2
acauuugacaaguuggguuga		1	0	Mm2
acauuugacaaguuggguugt	J	1	1	Mm2
acauuugacaaguuggguug	au	6	0	Mm2
		_	_	

acauuugacaaguuggguugauU 1 1 Mm2