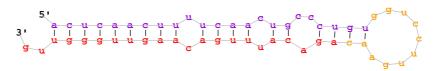
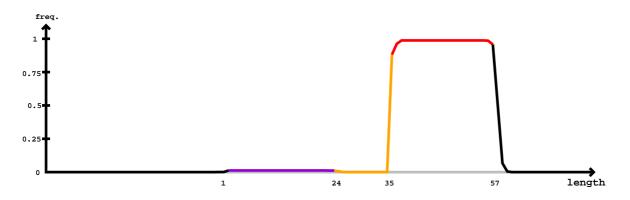
Provisional ID: NMRB01000004.1_Notospermus_geniculatus_strain_Ushimado_scaffold4,_whole_genome_shotgun_sequence_171

603.2 Score total Score for star read(s) 3.9 598.7 Score for read counts 1.2 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 1186 1171 Mature read count Loop read count 0 Star read count 15





Star	Mature

gaagagacagugacuggacuggaguugguaauaucgacucaacuuuucaacugcccugu <mark>gguccuugaacagacauuugacaaguuggguuga</mark> uguucacaaaccaagcauu	exp		
$\dots \dots (((((\dots \dots)))) \dots (((((((((((((((((((($	reads	mm	sample
agacauuugacaaguuggguCg	1	1	Fm1
aAacauuugacaaguuggguug	1	1	Fm1
agacauuugacaaguuggguGg	2	1	Fm1
agacauuugacaaguugCguug	1	1	Fm1
agacauuugacaaguuggguug	62	0	Fm1
agacauuugacaaguuCgguug	1	1	Fm1
agacauuugacaaguuggguuga	11	0	Fm1
agacauuugacaaguuggguugU	70	1	Fm1
agacauuugacaaguuggguugG	1	1	Fm1
agacauuugacaaguuggguugaA	2	1	Fm1
agacauuugacaaguuggguugUuagacauuugacaaguuggguugUu	4	1	Fm1
agacauuugacaaguuggguugau	3	0	Fm1
gacauuugacaaguuggguuga	8	0	Fm1
acauuugacaaguugggCugau	1	1	Fm1
acauuugacaaguuggguugau	4	0	Fm1
acauuugacaaguuggguugauh	5	1	Fm1
agacauuugacaaUuuggguu	1	1	Fm2
agacauuugacaaguugguu	14	0	Fm2
agacauuugacaaguuggguGg	1	1	Fm2
agacauuugacaaguuggguug	180	0	Fm2
agacauuugacaaguugCguug	3	1	Fm2
agacauuugacaGguuggguug	4	1	Fm2
agacauuugacaaguugAguug	2	1	Fm2
agacauuugacaaguuggguugU	94	1	Fm2
agacauuugacaaguuggguuga	50	0	Fm2
agacauuugacaGguuggguuga	4	1	Fm2
agacauuugacaaguuggguugG	2	1	Fm2
agacGuuugacaaguuggguuga	1	1	Fm2
agacauuugacaaguuggguugau	5	0	Fm2
agacauuugacaaguuggguugUu	8	1	Fm2
gacauuugacaaguuggguug	6	0	Fm2
gaganinga gaagungghunga	1	1	Em 2

Star Mature

Star Mature				
gaagagacagugacuggacuggaguugguaauaucgacucaacuuuucaacugcccugugguccuugaac <mark>agacauuugacaaguuggguu</mark> gauguucacaa	accaagcauu			
		41	0	Fm2
gacauuugacaaguuggguuga				
acauuugacaaguuggguugaA		2	1	Fm2
acauuugacaaguuggguugau	• • • • • • • • • • • • • • • • • • • •	9	0	Fm2
agacauuugacaaguugggu		1	0	Mm1
agacauuugacaaguugguug		14	0	Mm1
agacauuugacaaguuggguugU		24	1	Mm1
.agacauuugacaaguuggguugŪu		3	1	Mm1
		1	1	Mm2
cgacucaacuuuucaacugcccG	• • • • • • • • • • • • • • • • • • • •			
acucaacuuuucaacugcecugu		9	0	Mm2
agacauuugacaaguugggu.		1	0	Mm2
agacauuugacaaguuggguuagacauuugacaaguuggguu		6	0	Mm2
agacauuugaUaaguugguugagacauuugaUaaguugguug		2	1	Mm2
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agacauuugCcaaguugguug		1	1	Mm2
agacauuugacaaguuggguuU		1	1	Mm2
.agacauuugacaaguugAguug.		1	1	Mm2
agacauuugacaaguuggguug		62	0	Mm2
agacauuugacaaguCggguug		1	1	Mm2
agacauuugacaaguuggguuga		22	0	Mm2
agacauuugacaaguuggguugU	• • • • • • • • • • • • • • • • • • • •	170	1	Mm2
agacauuugacaaguuggguugG		2	1	Mm2
agacauuugacaaguuggguugau		2	0	Mm2
agacauuugacaaguuggguugUu		6	1	Mm2
agacauuugacaaguuggguugaG		1	1	Mm2
gacauuugacaaguuggguuga		13	0	Mm2
acauuugacaaguuggguuga		1	0	Mm2
acauuugacaaguuggguugU		1	1	Mm2
			0	
acauuugacaaguuggguugau		6		Mm2
acauuugacaaguuggguugauU	• • • • • • • • • • • • • • • • • • • •	1	1	Mm2
acucaacuuuucaacugcccugu	• • • • • • • • • • • • • • • • • • • •	1	0	Fm3
agacauuugacaaguuggguuagacauuugacaaguuggguu	• • • • • • • • • •	10	0	Fm3
agacauuugacaaguugUguu		1	1	Fm3
agacauuugacaaguuggguuU		1	1	Fm3
agacauuugacaaguuggguug		54	0	Fm3
agacauuugacaaguuggguuga		18	0	Fm3
agacauuugacaaguuggAuuga		1	1	Fm3
agacauuugacaaguuggguugU		41	1	Fm3
		2	0	Fm3
agacauuugacaaguuggguugau		2		
gacauuugacaaguuggguug	• • • • • • • • • • • • • • • • • • • •		0	Fm3
gacauuugacaaguuggguuga		14	0	Fm3
gacauuugacaaguuggguugU		2	1	Fm3
gacauuugacaaguuggguugau		3	0	Fm3
gacauuugacaaguuggguugaA		6	1	Fm3
acucaacuuuucaacugeecugu		4	0	Mm3
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agacauuugacaaguuggguuU		1	1	Mm3
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			0	Mm3
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agacauuugacaaguuggguugU		45	1	Mm3
.agacauuugacaaguuggguugau.		2	0	Mm3
agacauuugacaaguuggguugUu		3	1	Mm3
.agacauuugacaaguuggguugaA	• • • • • • • • • • • • • • • • • • • •	1	1	Mm3