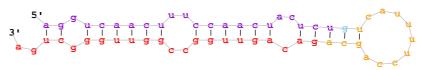
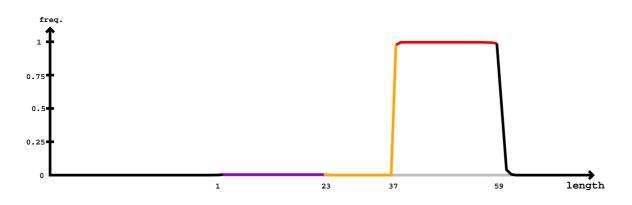
$: NMRB01000076.1_Notospermus_geniculatus_strain_Ushimado_scaffold 76,_whole_genome_shotgun_sequence_2805$ Provisional ID

769.4 Score total Score for star read(s) 3.9 764.9 Score for read counts 1.3 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 1512 Mature read count 1506 Loop read count 0 Star read count 6





Star	Mature	
5'- guugugagggucaaagugaaugguuugugacauaaggucaacu	uuccaacuacucugucauuuuccagcagacaguuggccgguugggcugauguuuacagaucagugcau	1

guugugagggucaaagugaaugguuugugacauaaggucaacuuuccaacuaugucauuuuccagcagacaguuggecgguuggggcugauguuuacagaucagugcau	exp		
((((((((((((((((((((((((((((reads	mm	sample
gacaguugggcugacaguugggcu	2	0	Fm3
gacaguuggccgguuggAcug	1	1	Fm3
gacaguuggccgguuggCcuga	3	1	Fm3
Uacaguuggccgguugggcuga	1	1	Fm3
gacaguugagccgguugAgcuga	4	1	Fm3
gacaguuggccgguuggAcuga	3	1	Fm3
gacaguuggcGgguugggcuga	2	1	Fm3
gacaguuggccgguuggUcuga	1	1	Fm3
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gacaguuggccgguugggcugŪu	1	1	Fm3
gacaguuggccgguugggcugau	5	0	Fm3
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acaguuggccgguugggcuga	10	0	Fm3
acaguuggccgguugggGuga	1	1	Fm3
acaguuggccgguugggcugau	1	0	Fm3
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gacaguuggcogguugggcugG	1	1	Mm3
gacaguuggcogguugggcugU	3	1	Mm3
gacaguuggcogguugggcugaA	2	1	Mm3
gacaguuggcogguugggcugau	6	0	Mm3

.....gacaguuggccgguugggcugauU.....

Star Mature

guugugagggucaaagugaaugguuugugacauaaggucaacuuuccaacuacucugucauuuuccagcagacaguuggccgguugggcugauguuuacagaucagugcau

gacaguuggccgguugggcuga	203	0	Fm1
gacGguuggccgguugggcuga	1	1	Fm1
gacaguuggUcgguugggcuga	1	1	Fm1
gacaguuggccgguuggCcuga	1	1	Fm1
gacaguuggccgguugggcugU	2	1	Fm1
gacaguuggccgguugggcugG.	4	1	Fm1
gacaguuggccgguugCgcuga.	1	1	Fm1
gacaAuuggccgguugggcuga	1	1	Fm1
gaAaguuggccgguugggcuga	1	1	Fm1
gacaguuggecgguugggcuAa	1	1	Fm1
		1	
gacaguuggccgguuggAcuga	3		Fm1
gacaguugggccgguugggcugaA	5	1	Fm1
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	1	1	Fm2
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	1	1	Mm1
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aaggucaacuuuccaacuacucu	1	0	Mm2
aggucaacuuuccaacuacucu	4	0	Mm2
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