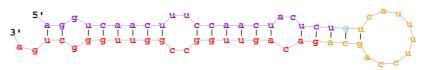
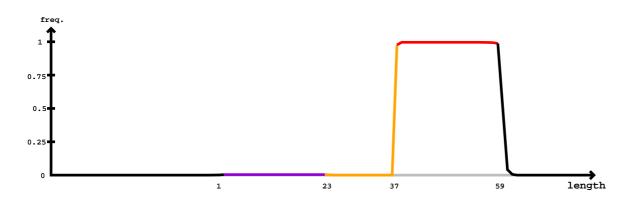
$: NMRB01000004.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold4,\_whole\_genome\_shotgun\_sequence\_173$ Provisional ID

769.6 Score total Score for star read(s) 3.9 764.9 Score for read counts Score for mfe 1.4 Score for randfold 0 Score for cons. seed -0.6 Total read count 1512 1506 Mature read count Loop read count 0 Star read count 6





	Star	Mature
5'- uuugucagggucugagugaaugguuugugacau	aaggucaacuuuccaacuacucugucauuuuccago	cagacaguuggccgguugggcugauguuuacagaucagug

uuugucagggucugagugaaugguuugugacauaaggucaacuuuccaacuacucug <mark>ucauuuuccagcagacaguuggccgguugggcugau</mark> guuuacagaucagugcau	exp		
(((())))(((((((((((((((((((((	reads	mm	sample
Uacaguuggccgguugggcuga	1	1	Fm2
gacaguuggccgguugggcugG	10	1	Fm2
gacaguuggcGgguugggcuga	1	1	Fm2
gacaguuggccgguuggUcuga	2	1	Fm2
gacaguuggccgguuggAcuga	1	1	Fm2
gacaguuggccgguugAgcuga	1	1	Fm2
gacaguuggccgguugggcuga	242	0	Fm2
gacaguuggccgguugggcugau	11	0	Fm2
gacaguuggccgguugggcugaA	25	1	Fm2
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gacaguuggcogguugggcugauk	6	1	Fm2
acaguuggcogguugggcuga	5	0	Fm2
acaguuggccgguugggcugau	3	0	Fm2
gacaguuggcogguugggcuga	203	0	Fm1
gacaguuggccgguuggAcuga	3	1	Fm1
gacaguuggcogguugggcugU	2	1	Fm1
gacGguuggccgguugggcuga	1	1	Fm1
gacaguuggUcgguugggcuga	1	1	Fm1
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gacaAuuggccgguugggcuga	1	1	Fm1
gacaguuggccgguugggcugG	4	1	Fm1
gacaguuggccgguugggcuAa	1	1	Fm1
gacaguuggccgguugCgcuga	1	1	Fm1
gacaguuggccgguuggCcuga	1	1	Fm1
gacaguuggccgguugggcugau.	23	0	Fm1
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gacaguuggccgguugggcugauA	4	1	Fm1
acaguuggccgguugggcuga	2	0	Fm1
acaguuggccgguuggcugaA.	3	1	Fm1
2 22 23 33 33			

Star Mature

Star Ma	ature			
uuugucagggucugagugaaugguuugugacauaaggucaacuuuccaacuacucugucauuuuccagcagac	aguuggccgguugggcugauguuuacagaucagugcau			
gaca	aguuggccgguuggAcug	1	1	Fm3
gaca	aguuggcAgguugggcuga	1	1	Fm3
gaca	aguuggGcgguugggcuga	1	1	Fm3
gaca	aguuggccgguuUggcuga	2	1	Fm3
gaca	aguuggccgguuggUcuga	1	1	Fm3
gaca	aguuggccggAugggcuga	1	1	Fm3
gaca	aguuggccgguugggAuga	1	1	Fm3
gaca	aguuggccgguugggcugU	1	1	Fm3
	aguuggccgguugggcuga	1	1	Fm3
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	aguuggcGgguugggcuga	2	1	Fm3
	aguuggccgguugAgcuga	4	1	Fm3
	aguuggccgguugggcuga	1	1	Fm3
	aguuggccgguugCgcuga	3		Fm3
	aguuggcegguuggCeuga	3		Fm3
	agAuggccgguugggcuga			Fm3
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	aguuggccgguugggcugaA			Fm3
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	aguuggccgguugggcugauA			Fm3
	aguuggccgguugggcuga			Fm3
	aguuggccgguugggGuga			Fm3
	aguuggccgguugggcugau			Fm3
	<u>agaaggeeggaaggeagaa</u>	-	•	LING
gaca	aguuggccgguuggcug	5	0	Mm1
	aguuggccgguugggGug			Mm1
	aguuggccgguugUgcuga			Mm1
				Mm1
	aguuggccgguuggUcuga			Mm1
	aguuggcegguuggAcuga			Mm1
	aguuggcegguugCgcuga			Mm1
	aguuggccgguugggcuga			Mm1
	aguuggccgguugggcugU			Mm1
	aguuggccgguugggcugaG			Mm1
	aguuggccgguugggcugau			Mm1
	aguuggccgguugggcugaA			Mm1
	aguuggccgguugggcugauU			Mm1
gaca	ayuuggeegguugggeugangu	,	-	PHILL
uaaggucaacuuuccaacuacucu		1	0	Mm2
aaggucaacuuuccaacuacucuaaggucaacuuuccaacuacucu	•••••			Mm2
.aggucaacuuuccaacuacucu	•••••			Mm2
gaca				Mm2 Mm2
gaca				
gaca				Mm2
gaca				Mm2 Mm2
gaca				
gaca				Mm2
aca				Mm2
aca	aguuggccgguugggcugau	1	0	Mm2
		24	0	16- 7
gaca				Mm3
gaca				Mm3
gaca				Mm3
gaca				Mm3 Mm3
				Mm3
gac	~_~	-	_	حسب