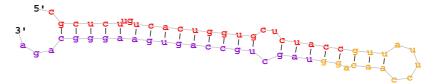
Provisional ID $: NMRB01003202.1_Notospermus_geniculatus_strain_Ushimado_scaffold3202,_whole_genome_shotgun_sequence_42604$

674.5 Score total 3.9 Score for star read(s) Score for read counts 669.5 Score for mfe 1.6 Score for randfold 0 -0.6 Score for cons. seed Total read count 1325 956 Mature read count 0 Loop read count

369

Mature

Star read count



sample

Fm3 Fm3

Fm3

Fm3

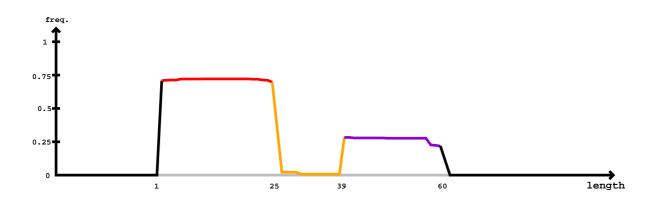
0

1

1

0

0



5'-	uguuaaaaguguugacgauc <mark>cgcucuugucacuggugcucuaccg</mark> uuauuccaacagguagcugccagugaagggcagacuguugacauagauucuucagcaucaagagguggu	-3'	obs	
	uguuaaaaguguugacgauc <mark>cgcucuugucacuggugcucuaccguuauuccaacagg</mark> uagcugccagugaagggcagacuguugacauagauucuucagcaucaagagguggu		exp	
	(((((((((((((((((((((((((((((((((read	s	mm
		3		0
		2		0
		2		0
	cgcucuugucacuggugcucuacc	81		0

Star

.....cacuggugcucuaccguuauuccaacagg.....uagcugccaguAaagggc...... 1 1 36 0uagcugccagugaagggc..... 1 1uagcugccagugaagggca.....uagcugccagugaagggcag..... uagcuqccaquqaaqqCcaqa. 3 1 Fm3uagcCgccagugaagggcaga..... 166 0 1uagcugccagugaagggAaga..... 0 1 0 1 1cgcucuugucacugguAcucuacc..... 2 1 1 1uagcugccagugaagggc...... 6 0

.....uagcugccagugaagggcGg.....uagcugccagugaagggcaga.....

Mature Star			
uguuaaaaguguugacgauccgcucuugucacuggugcucuaccguuauuccaacagguagcugccagugaagggcagacuguugacauagauucuucagcaucaagagguggu			
.cgcucuugucacuggugcucuacc	201	0	Fm2
.egcucuugueCcuggugcucuacc.	1	1	Fm2
.cgcucuugucacuggugAucuacc.	2	1	Fm2
.cgcucuugueUcuggugcucuacc.	1	1	Fm2
.cgcucuugucacuggugcucuaccgCuau.	1	1	Fm2
cgcucuugucacuggugcucuaccguuau	14	0	Fm2
uagcugccagugaagggc	1	0	Fm2
	2	1	Fm1
	1	1	Fm1
	100	0	Fm1
.cgcucuugucacuggugGucuacc	1	1	Fm1
.cgcucuuguUacuggugcucuacc	1	1	Fm1
.cgcucuugucacuggugcucuaccU	6	1	Fm1
.cgcucuugucacuggugcucuaccg.	4	0	Fm1
cucuugucacuggugcucuacc.	2	0	Fm1
accguuauuccaacagguagcugcca	3	0	Fm1
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.uagcugccagugaagggcag	4	0	Fm1
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.cgcucuugucacuggugcG	1	1	Mm2
.cgcucuugucacuggugcu	1	0	Mm2
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. cgcucuugucacuggugcCcuacc.	1	1	Mm2
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cgcucuugucUcuggugcucuacc	1	1	Mm2
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Ugcucuugucacuggugcucuacc	1	1	Mm2
cgcucuugucacugguAcucuacc	1	1	Mm2
cgcucuugucCcuggugcucuacc	1	1 1	Mm2 Mm2
	4	0	Mm2
cgcucuugucacuggugcucuaccg	6	1	Mm2
	1	1	Mm2
	4	0	Mm2
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cuugucacuggugcucuacc	4	0	Mm2
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uagcugccagugaagggcaga	16	0	Mm2
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cgcucuugucacuggugcucu	3	0	Mm1
Cgcucuugucacuggugcucuac	10	0	Mm1
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cgcucuugucacuggugcucuacc	86	0	Mm1
cuugucacuggugcucuaccguuauuccaacaggua	6	0	Mm1
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uagcugccagugaagggcaga	8	0	Mm1