

Provisional ID : NMRB01000035.1_Notospermus_geniculatus_strain_Ushimado_scaffold35_whole_genome_shotgun_sequence_1516

Score total : 164.8

Score for star read(s) : 3.9

Score for read counts : 159.7

Score for mfe : 1.8

Score for randfold : 0

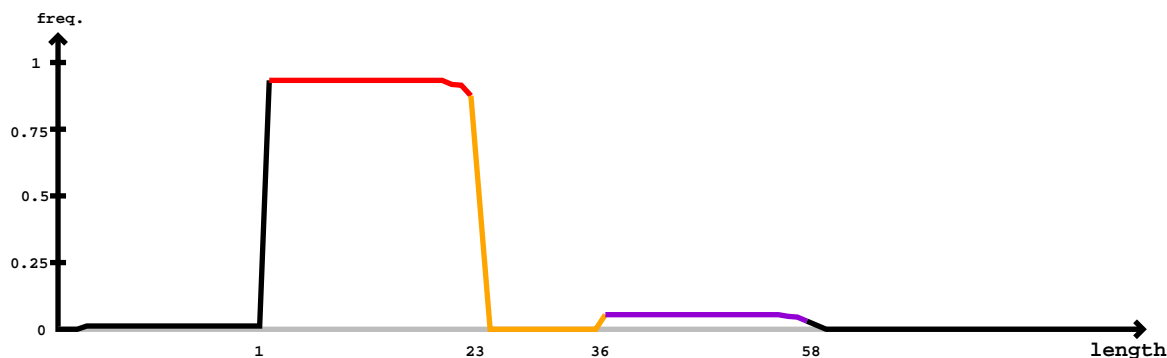
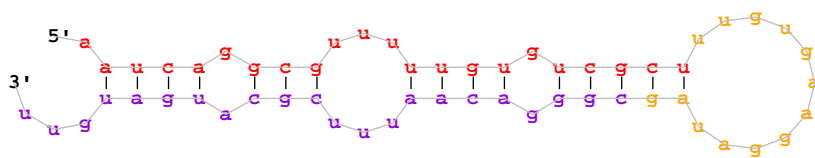
Score for cons. seed : -0.6

Total read count : 325

Mature read count : 307

Loop read count : 0

Star read count : 18



Mature

Star

5' -		-3'	obs	
	gcagauagcgacuggcgga	aaucaggcgguuuuugugucgcuuuugugaaggauagcgggacaauuucgcaugauguucgaccaguugaacacacacacacauacaacguga	reads	mm
	gcagauagcgacuggcgga	aaucaggcgguuuuugugucgcuuuugugaaggauagcgggacaauuucgcaugauguucgaccaguugaacacacacacacauacaacguga	exp	sample
(((((((((((.....)))))))))))).....((((.....)))).			
aaucaggcgguuuuugugucgcu.....	20	0	Mm2
aaucaggcgguuuuugugucgcG.....	1	1	Mm2
aaucaggcgguuuuugugucgcC.....	1	1	Mm2
aaucaggcgguuuuugugucgc.....	5	0	Mm1
aaucaggcgguuuuugugucgc.....	6	0	Mm1
aaucaggcgguuuuugugCgcgu.....	1	1	Mm1
aaucaggcgguuuuugugucgc.....	38	0	Fm2
aaucaggcgguuuuugugucgcG.....	2	1	Fm2
aaucaggcgguuuuugugCgcgu.....	1	1	Fm2
cgggacaauuucgcaugau.....	2	0	Fm2
aaucaggcgguuuuuguguc.....	3	0	Fm1
aaucaggcgguuuuugugucgc.....	6	0	Fm1
aaucaggcgguuuuugugAgcgu.....	1	1	Fm1
Nauaggcgguuuuugugucgcu.....	1	1	Fm1
aaucaggcgguuuuugUucgcu.....	1	1	Fm1
aaucaggcgguuuuugugAgcgu.....	2	1	Fm1
aaucaggcgguuuuugugCgcgu.....	3	1	Fm1
aaucaggcgguuuuugugucgcC.....	1	1	Fm1
aaucaggcgguuuuugugucgcG.....	6	1	Fm1
aaucaggcgguuuuugugucgcu.....	184	0	Fm1
aaucaggcgguuuuugugGcgcgu.....	1	1	Fm1
aaucaggcgguuuuugugucgcu.....	1	1	Fm1
aaucaggcgguuuuugugucgcu.....	4	0	Fm1
cgggacaauuucgcaugaugu.....	6	0	Fm1
aaucaggcgguuuuuguguc.....	1	0	Mm3
aaucaggcgguuuuugugAgcg.....	1	1	Mm3
aaucaggcgguuuuugugucgcu.....	6	0	Mm3
aaucaggcgguuuuugugucgcuA.....	1	1	Mm3
cgggacaauuucgcaugaug.....	1	0	Mm3

Mature

Star

gcagauagcgacuggucgga	aaucaggcg	uuuuugugucgc	uuugugaaggauag	cgggaca	uuucgaugaugu	ucgaccaguugaacacacacacacauacaacguga			
.....	1	0	Mm3
.....			
.cagauagcgacuggucgga.....	4	0	Fm3
.....	.aaucaggcg	uuuuuguguA.....	1	1	Fm3
.....	.aaucaggcg	uuuuugugucg.....	1	0	Fm3
.....	.aaucaggcg	uuuuugugucgc.....	1	0	Fm3
.....	.aaucaggcg	uuuuugugucgcu.....	5	0	Fm3
.....	.aaucaggcg	uuuuugugucgcu.....	1	0	Fm3
.....	cgggaca	uuucgaugaugu.....	5	0	Fm3
.....	cgggaca	uuucgaugaugu.....	3	0	Fm3