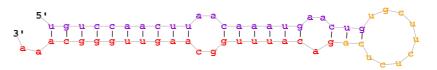
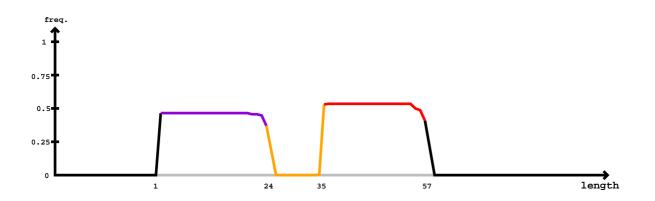
Provisional ID : NMRB01000004.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold4,\_whole\_genome\_shotgun\_sequence\_142

116.4 Score total Score for star read(s) 3.9 111.3 Score for read counts Score for mfe 1.8 Score for randfold 0 Score for cons. seed -0.6 Total read count 230 123 Mature read count Loop read count 0 Star read count 107

Star





Mature

5'- ggugacuuggcuggagcaguuguccaacuuaacaaaugaacugugcuucucucagacauuuggcaaguugggcaaaauguuucagacaucaaauauuggcaaccaauauuggu	-3'	obs	
ggugacuuggcuggagcaguuguccaacuuaacaaaugaacug <mark>ugcuucucagacauuuggcaaguugggcaaa</mark> auguuucagacaucaaauauuggcaaccaauauuggu		exp	
((())))))))))))))))))))))))))))))	read	ds mm	sample
.uguccaacuuaacaaaugaacug	18	0	Mm1
gacauuuggcaaguugggcaaa	6	0	Mm1
uguccaacuuaacaaauga	1	0	Mm2
.uguccaacuuaacaaaugaac	1	0	Mm2
.uguccaacuuaacaaaGgaacu	1	1	Mm2
uguccaacuuaacaaaugaacu	9	0	Mm2
.uguccaacuuaacaaaugaacug	29	0	Mm2
uguccaacuuaacaaaugaacugu	3	0	Mm2
gacauuuggcaaguugggcgacauuuggcaaguugggc	1	0	Mm2
gacauuuggcaaguugggcaaa	3	0	Mm2
gacauuuggcaaguugggcaaaG	1	1	Mm2
gacauuuggcaaguugggcaaaU	1	1	Mm2
gacauuuggcaaguugggcaaaa	2	0	Mm2
uguccaacuuaacaaaugaacug	4	0	Fm1
uguccaacuuaacaaaugaacugu	2	0	Fm2
gacauuuggcaaguugggcaaa	3	0	Fm2
.uguccaacuuaacaaauga	1	0	Mm3
.uguccaacuuaacaaaugaac	1	0	Mm3
.uguccaacuuaacaaaugaacu	1	0	Mm3
uguccaacuuGacaaaugaacugu	1	1	Mm3
.uguccaacuuaacaaauCaacugu	1	1	Mm3
uguccaacuuaacaaaugaacugu	1	0	Mm3
agacauuuggcaaguugggc	1	0	Mm3
gacauuuggcaaguugggcaaa	1	0	Mm3
.uguccaacuuaacaaaugaacu	7	0	Fm3
.uguccaacuuaacaaaugaacug	26	0	Fm3
gacauuuggcaaguugggc	6	0	Fm3

Star Mature

ggugacuuggcuggagcaguuguccaacuuaacaaaugaacug <del>ugcuucucucagacauuuggcaaguugggcaaaa</del> uguuucagacaucaaauauuggcaaccaauauuggu			
gacauuuggcaaguugggcaa	16	0	Fm3
gacauuuggcGaguugggcaa	1	1	Fm3
gacauuuggcaaguugggcaaG	1	1	Fm3
gacauuuggcaaguugggcaaa	50	0	Fm3
gacauuuggcaaguuggCcaaaagacauuuggcaaguuggCcaaaa	1	1	Fm3
gacauuuggcaaguugggcaaaa.	22	0	Fm3
gacauuuggcaaguugggcaaaG	3	1	Fm3
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