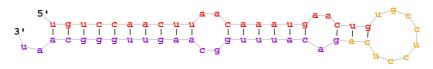
Provisional ID  $: NMRB01000004.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold4,\_whole\_genome\_shotgun\_sequence\_160$ 

Score total 81.1 Score for star read(s) 3.9 Score for read counts 76.1 Score for mfe 1.7 Score for randfold 0 -0.6 Score for cons. seed Total read count 161 107 Mature read count Loop read count 0

54

Mature

Star read count



exp

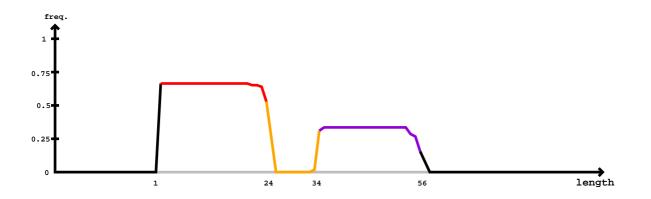
Mm2

Mm 2

Mm2

1

0



Star 5'- ggcggcuuggcuggagcagu<mark>uguccaacuuaacaaaugaacugugccuccuc</mark>agacauuuggcaaguugggcaauauguuucagacaucaaauggucuauguguuccuuugc ggcggcuuggcuggagcagu<mark>uguccaacuuaacaaaugaacugugccuccuca</mark>gacauuuggcaaguugggcaauauguuccagacaucaaauggucuauguguuccuuugc

.....uquccaacuuaacaaaugaacu..... .....uguccaacuuaacaaaugaacug.....

	-		
(((.((((((((((((((((())).))))))	reads	mm	sample
uguccaacuuaacaaaugaacu	7	0	Fm3
uguccaacuuaacaaaugaacug	26	0	Fm3
agacauuuggcaaguugggcGaua	2	1	Fm3
gacauuuggcaaguugggc	6	0	Fm3
gacauuuggcaaguugggca	3	0	Fm3
gacauuuggcGaguugggcaa	1	1	Fm3
gacauuuggcaaguugggcaa	16	0	Fm3
gacauuuggcaaguugggcaau	4	0	Fm3
gacauuuggcaaguugggcaaG	1	1	Fm3
acauuuggcaaguugggcaa	1	0	Fm3
uguccaacuuaacaaauga	1	0	Mm3
uguccaacuuaacaaaugaac	1	0	Mm3
uguccaacuuaacaaaugaacu	1	0	Mm3
uguccaacuuaacaaaugaacugu	1	0	Mm3
uguccaacuuGacaaaugaacugu	1	1	Mm3
uguccaacuuaacaaauCaacugu	1	1	Mm3
agacauuuggcaaguugggc	1	0	Mm3
uguccaacuuaacaaaugaacug	4	0	Fm1
uguccaacuuaacaaaugaacugu	2	0	Fm2
acauuuggcaaguugggcGaua	3	1	Fm2
uguccaacuuaacaaaugaacug	18	0	Mm1
gacauuuggcaaguuggCcaau	1	1	Mm1
gacauuuggcaaguugggcaau	14	0	Mm1

Mature Star
ggcggcuuggcuggagcaguuguccaacuuaacaaaugaacugugccuccucagacauuuggcaaguugggcaauauguuucagacaucaaauggucuauguguuccuuugc