$Provisional\ ID \\ : NMRB01000794.1_Notospermus_geniculatus_strain_Ushimado_scaffold794,_whole_genome_shotgun_sequence_20940$

0.4 Score total Score for star read(s) -1.3 Score for read counts 0 2.3 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 114 Mature read count 105 Loop read count 3

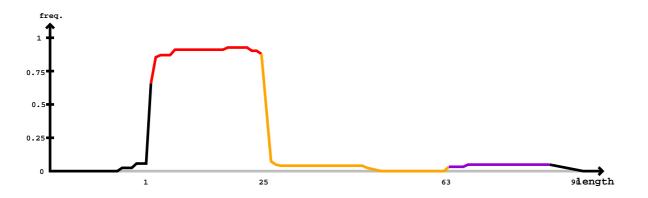
6

 ${\tt 5'-auggcaacgcugccauuagcaaggcugggucuggccuggcaccaccuggcucugggcugg}$

Star read count



iggcaugccaaccuuggcccaguccaggcaugacu



Mature Star

auggcaacgcugccauuagcaaggcugggucuggccuggcaccaccuggcucugggcuggcaagcccgggcauuggcuuggcaugccaugccauggccaguccaggcaugacu	exp	•	
(((((())))))(((((((((((((((((((reads	mm.	sample
caaggcugggCcuggcac	1	1	Mm3
caaggcugggCcuggcacc	1	1	Mm3
cugggCcuggcacgacca	1	1	Mm3
caccuggcucugggcuggcaagcc	1	0	Mm3
caaggcugggCcuggccuggc	3	1	Fm3
caaggcugggCcuggcac	4	1	Fm3
caaggcugggCcuggcacc	3	1	Fm3
aaggcugggCcuggcacc	1	1	Fm3
cugggCcuggcacga	1	1	Fm3
.cugggCcuggccuggcaccac	3	1	Fm3
caccuggcucugggcuggcaagc	1	0	Fm3
caugccaaccuuggcccagueU	2	1	Fm3
caugccaaccuuggcccaguccU	1	1	Fm3
ccaaccuuggcccaguccaggc	2	0	Fm3
cauuagcaaggcugggCcuggcacc	3	1	Mm1
uagcaaggcugggCcuggcacc	4	1	Mm1
caaggcugggCcuggcacc	30	1	Mm1
caaggcugggucuggcacc	1	0	Mm1
aaggcugggucuggcaccU	16	1	Mm1
.aggcugggCcuggcaccaggcugggggCcuggcacc	2	1	Mm1
caaggcugggCcuggcacc	3	1	Mm2
aaggcugggCcuggcacc	1	1	Mm2
.cuggcaccaccuggcucuggcuggcaag	2	0	Mm2
.ccuggcucugggcuggcaagccc	1	0	Mm2
caugccaaccAuggcccagucc	1	1	Mm2
caaggcugggCcuggccuggcacc	2	1	Fm1
caaggeugggCeuggeeuggeace	26	1	Fm2
aaggcugggCcuggcaccaaggcuggcacc	6	1	Fm2

Mature Star

augg caac g cug c cauu ag caa g g cug g g u cug g c cug g cac cac cug g cucu g g c cug g caa g c c c g g c a u u g g cau g c ca g c c a g c cag g cau g c ca g c cag g cau g c cag g cau g c cag g c