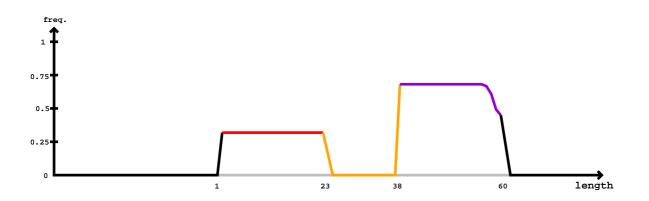
$: NMRB01000004.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold4, \underline{\ \ } whole\_genome\_shotgun\_sequence\_165$ Provisional ID

33.7 Score total

Score for star read(s) 3.9 Score for read counts 29.2 1.2 Score for mfe

Score for randfold 0 Score for cons. seed -0.6 Total read count 69 22 Mature read count Loop read count 0 Star read count 47



5'- agauggcagugcuguggauugcuccuggauugcugcucaacuuuccaaaugaacuguaauauucaauucagacauuuggccaguuggguaacuguuucaggguugaacuucu

Mature

agauggcagugcuguggauugcuccuggauugc <mark>ugcucaacuuuccaaaugaacu</mark> guaauauucaauucagacauuuggccaguuggguaac <b>u</b> guuucaggguugaacuucu	ex	.p	
((((((().))))))(((((((((((((	reads	mm	sample
ugcucaacuuuccaaaugaacug	1	0	Fm1
gacauuuggccaguugggu	1	0	Fm1
ugcCcaacuuuccaaaugaacug	1	1	Fm2
gacauuuggccaguuggguaaG	5	1	Mm1
gacauuuggccaguuggguaac	5	0	Mm1
ugcucaacuuuccaaaugaacu	8	0	Mm2
ugcCcaacuuuccaaaugaacug	3	1	Mm2
ugcucaacuuuccaaaugaacug	5	0	Mm2
gacauuuggccaguuggg	1	0	Mm2
gacauuuggccaguugggua	2	0	Mm2
gacauuuggccaguuggUua	1	1	Mm2
gacauuuggccaguuggguaa	2	0	Mm2
gacauuuggccaguuggguaac	7	0	Mm2
gacauuuggccaguuggguaacu	3	0	Mm2
ugcucaacuuuccaaaugaacu	1	0	Fm3
gacauuuggccaguugggu	3	0	Fm3
gacauuuggccaguugggua	4	0	Fm3
gacauuuggccaguuggguaGc	1	1	Fm3
gacauuuggccaguuggguaac	7	0	Fm3
gacauuuggccaguuggguaacu	1	0	Fm3

Mm3 Mm3

gacauuuggccaguugggua.....

Star