Provisional ID: NMRB01001134.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold1134,\_whole\_genome\_shotgun\_sequence\_25287

Score total : 3.1

 Score for star read(s)
 : -1.3

 Score for read counts
 : 0

 Score for mfe
 : 1.4

 Score for randfold
 : 0

 Score for cons. seed
 : 3

 Total read count
 : 606

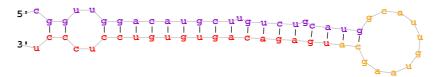
Mature read count Loop read count

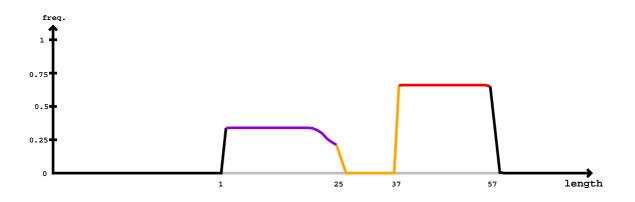
Star read count

400

0

206





Star

5'- ggaauuuuugaucagagaauccaaauuucuuccccgguuggacaugcuugucugcauggcauuguaagca <mark>ugagacaguguguccucccua</mark> ggaaugaaaguua	iugucau	-3' ol	os	
ggaauuuuugaucagagaauccaaauuucuucccegsuuggacaugcuugucugcauggcauuguaagca <mark>ugagacaguguguccucccua</mark> ggaaugaaaguua	lugucau	e:	кр	
(((((((()))))))((((((((((((((((	· • • • • •	reads	mm	sample
.cgguuggacaugcuugucu		9	0	Fm3
.cgguuggacaugcuugu		13	0	Fm3
Ggguuggacaugcuugu		1	1	Fm3
.cgguuggacaugcuuguUugc		1	1	Fm3
.cgguuggUcaugcuuguc		1	1	Fm3
cgguuggacaugcuugucugc		13	0	Fm3
cgguuggacaugcuugA		8	1	Fm3
cgguuggacaugcuugucugcau		10	0	Fm3
		88	0	Fm3
.cgguuggacaugcuuguAugcaug		1	1	Fm3
.ugagacaguguccuccc		4	0	Fm3
.ugagacagugucucucu		1	1	Fm3
.ugagacaguguccucAcu		5	1	Fm3
ugagacagugAguccucccu		1	1	Fm3
.ugagacaguguccucGcu		1	1	Fm3
Agagacaguguccucccu.		1	1	Fm3
.ugagacaguguccuUccu		1	1	Fm3
.ugagacaguguccuAccu		2	1	Fm3
.Ngagacaguguccucccu		1	1	Fm3
ugagacaguguccucccu		281	0	Fm3
.ugagacaguguccucccuU		2	1	Fm3
.ugagacaguguccucccuUg		2	1	Fm3
gagacaguguccucccu		1	0	Fm3
.cgguuggacaugcuuguc		2	0	Fm1
cgguuggacaugcuugucugcaug		3	0	Fm1
.ugagCcaguguccucccu		1	1	Fm1
		1	1	Fm1
ugagacaguguguccuUccu		1	1	Fm1
.ugagacaguguccucccG.		1	1	Fm1
ugagacaguguguccucccu		35	0	Fm1
		1	1	Fm1

......cgguuggacaugcuugucugc.....

Mature

Star Mature

ggaauuuuugaucagagaauccaaauuucuuccccgguuggacaugcuugucugcauggcauuguaagca <mark>ugagacaguguguccucccua</mark> ggaaugaaaguuaugucau			
	1	0	Mm3
	12	0	Mm3
.augagacaguguguccucccuU	1	1	Mm3
ugagacaguguguccuccc	1	0	Mm3
ugagacaguguguccucccu	31	0	Mm3
ugagacaguguguccucUcu	1	1	Mm3
ugagacaguguguccucccuU	4	1	Mm3
.cgguuggacaugcuugAcugca	1	1	Mm1
	14	0	Mm1
Ugguuggacaugcuugucugcaug	1	1	Mm1
	17	0	Mm1
Cguuggacaugcuugucugcaug.	1	1	Mm1
ugagacaguguguccucccu.	3	0	Mm1
ugagacaguguguccucccuU	1	1	Mm1
ugagacaguguguccucccuUg	1	1	Mm1
	2	0	Mm2
.cgguuggacaugcuugucugGau	1	1	Mm2
	5	0	Mm2
ugagacaguguguccuccc	1	0	Mm2
ugagacagugCguccucccu	1	1	Mm2
ugagacaguguguccucccG	1	1	Mm2
ugagacaguguguccucccu.	10	0	Mm2
ugagacaguguguccucccuU	1	1	Mm2