Provisional ID: NMRB01002131.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold2131,\_whole\_genome\_shotgun\_sequence\_36178

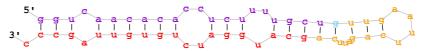
108.3 Score total Score for star read(s) 3.9 103.6 Score for read counts Score for mfe 1.4 Score for randfold 0 Score for cons. seed -0.6 Total read count 215 193 Mature read count

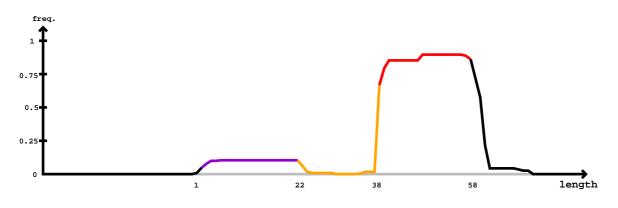
0

22

Loop read count

Star read count





gcagccauccuguagaaaucucaaaccuuugggucaacacaccucuuuugcuguugaauucagauucagcauggaucuguguuagcccacaggauugaguuccuucaccua	exp	•	
(((()))).(((((((.((((.((((.(((((((	reads	mm	sample
gcauUgaucuguguuagcccacgcauUgaucuguguuagcccac	12	1	Fm2
gucaacaccucuuuugcug	5	0	Fm1
ucagcauUgaucuguguuagcc	2	1	Fm1
ucagcauUgaucuguguuagccc	1	1	Fm1
gcauUgaucuguguuagccca	2	1	Fm1
gcauggaucuguguuagcccac	10	0	Fm1
gcauUgaucuguguuagcccac	1	1	Fm1
gcauggaucuCuguuagcccac	1	1	Fm1
gcauggaucuguguuagcccaca	4	0	Fm1
cauggaucuguguuagcccacU	4	1	Fm1
.auggaucuguguuagcccac	2	0	Fm1
.auggaucuguguuGgccac	1	1	Fm1
.auggaucuguguuagcccaca	1	0	Fm1
ggucaacaccucuuuugcu	5	0	Mm1
gcauggaucuguguuGgccca	8	1	Mm1
gcauggaucuguguuGgcccac	5	1	Mm1
.uguguuagccGacaggauugagu	4	1	Mm1
gggucaacaccucuuuugcuguuCaau	1	1	Mm2
ggucaacaccucuuuugcu	2	0	Mm2
ggucaacaccucuuuugcug	2	0	Mm2
gucaacaccucuuuugcugu	1	0	Mm2
aacacaccucuuuugcuguuCaau	1	1	Mm2
.uucagcauUgaucuguguuagcc	1	1	Mm2
gcauUgaucuguguuagccc	3	1	Mm2
gcauggaucuguguuagcUc	3	1	Mm2
gcauUgaucuguguuagccca	3	1	Mm2
gcauggaucuguguuagcUca	1	1	Mm2
gcauggaucuguguuagcUcac	2	1	Mm2
gcauUgaucuguguuagcccac	11	1	Mm2
.cauUqaucuququuaqcc.	2	1	Mm2

Star Mature

gcagccauccuguagaaaucucaaaccuuugggucaacaccucuuuugcuguugaauucagauucagcauggaucuguguuagcccacaggauugaguuccuucaccua			
cauUgaucuguguuagccc	1	1	Mm2
cauUgaucuguguuagcccac	1	1	Mm2
cauUgaucuguguuageccaca	2	1	Mm2
auUgaucuguguuagcccac	2	1	Mm2
gggucaacaccucuuuugcug	1	0	Fm3
gucaacacaccucuuuugcugu	1	0	Fm3
ucaacacaccucuuuugcug	5	0	Fm3
.gcauUgaucuguguuagc	2	1	Fm3
.gcauUgaucuguguuagcc	1	1	Fm3
geauggaucuguguuagece	12	0	Fm3
.gcauUgaucuguguuagccc	25	1	Fm3
.gcauUgaucuguguuagccca	1	1	Fm3
gcauggaucuguguuagcUca	6	1	Fm3
gcauggaucuguguuagcccac	6	0	Fm3
gcauUgaucuguguuagcccac	20	1	Fm3
.gcauggaucuguguuGgcccac	2	1	Fm3
gcauggaucuguguuagcUcac	2	1	Fm3
gcauggaucuguguuagcccacU	6	1	Fm3
cauUgaucuguguuageccac	4	1	Fm3
cauggaucuguguuagcccaca	5	0	Fm3
cauggaucuguguuagcccacU	4	1	Fm3
cauggaucuguguuagcUcaca	6	1	Fm3
auggaucuguguuageccaeU	2	1	Fm3
auggaucuguguuagcccaca	1	0	Fm3
auUgaucuguguuagcccaca	5	1	Fm3
.gcauggaucuguguuagcccac	1	0	Mm3
.uguguuagccGacaggauug	2	1	Mm3
.uguguuagccGacaggauuga.	2	1	Mm3
.uguguuagccGacaggauugagu	2	1	Mm3