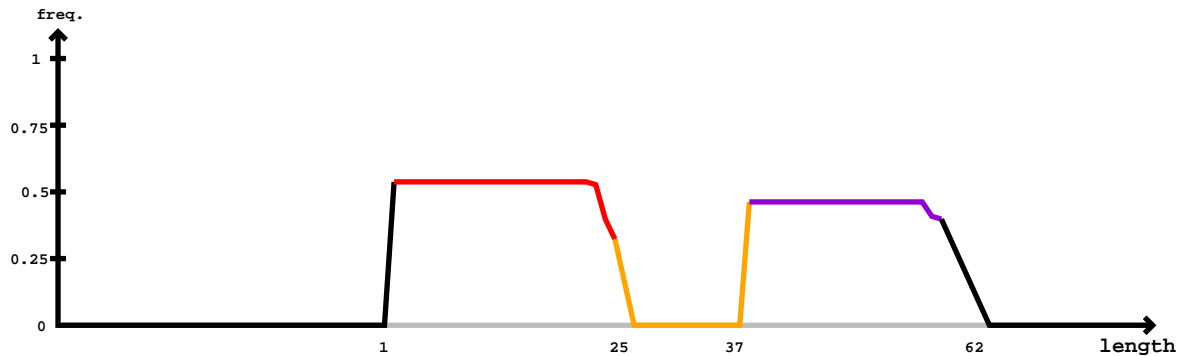
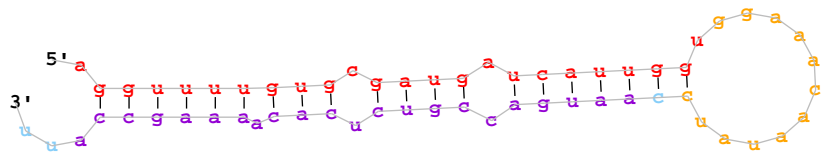


Provisional ID : NMRB01003120.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold3120,\_whole\_genome\_shotgun\_sequence\_42094  
Score total : 46  
Score for star read(s) : 3.9  
Score for read counts : 41.4  
Score for mfe : 1.2  
Score for randfold : 0  
Score for cons. seed : -0.6  
Total read count : 93  
Mature read count : 50  
Loop read count : 0  
Star read count : 43



### Mature

### Star

5' -		-3'	obs		
			exp		
ccguguguucaacaaggaaaacguucguagauuagguuuugugcgaugaucauugguggaacaauauc	aaugaccgucucacaaaagcca	uuucugagcgucaaaacaucg			
ccguguguucaacaaggaaaacguucguagauuagguuuugugcgaugaucauugguggaacaauauc	aaugaccgucucacaaaagcca	uuucugagcgucaaaacaucg			
(((((.....)))..(((((((.....(((((((.....(((((((.....)))))))))))))))))))))).....			reads	mm	sample
.....agguuuugugcgaugaucauug.....			1	0	Mm2
.....agguuuugugcgaugaucauugg.....			1	0	Mm2
.....aaugaccgucucacaaaagcca.....			1	0	Mm2
.....aaugaccgucucacaaaagcca.....			6	0	Mm2
.....agguuuugugcgaugaucauuggu.....			1	0	Mm3
.....agguuuugugcgaugaucauugguU.....			1	1	Mm3
.....aaugaccgucucacaaaagcca.....			1	0	Mm3
.....aaugaccgucucacaaaagcca.....			3	0	Mm3
.....agguuuugugcgaugaucauuggA.....			25	1	Fm2
.....agguuuugugcgaugaucauuggu.....			1	0	Fm2
.....agguuuugugcgaugaucauu.....			1	0	Fm1
.....agguuuugugcgaugaucauug.....			6	0	Fm1
.....agguuuugugcgaugaucauugg.....			2	0	Fm1
.....aaugaccgucucacaaaagcca.....			2	0	Fm1
.....aaugaccgucucacaaaagcca.....			14	0	Fm1
.....aaugaccgucucacaaaagcca.....			1	0	Fm1
.....agguuuugugcgaugaucauug.....			5	0	Fm3
.....agguuuugugcgaugaucauugg.....			4	0	Fm3
.....agguuuugugcgaugaucauuggA.....			2	1	Fm3
.....aaugaccgucucacaaaag.....			5	0	Fm3
.....aaugaccgucucacaaaagc.....			1	0	Fm3
.....aaugaccgucucacaaaagcc.....			1	0	Fm3
.....aaugaccgucucacaaaagcca.....			6	0	Fm3
.....aaugaccgucucacaaaagccau.....			2	0	Fm3