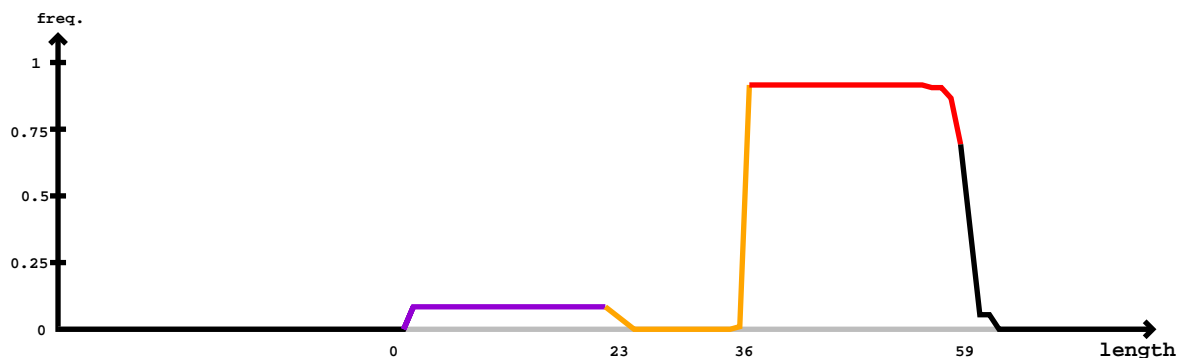
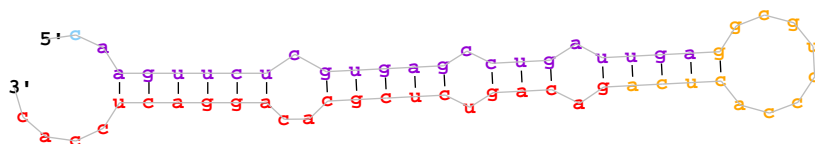


Provisional ID : NMRB01002087.1_Notospermus_geniculatus_strain_Ushimado_scaffold2087,_whole_genome_shotgun_sequence_35667
Score total : 96.8
Score for star read(s) : 3.9
Score for read counts : 97
Score for mfe : -3.4
Score for randfold : 0
Score for cons. seed : -0.6
Total read count : 202
Mature read count : 185
Loop read count : 0
Star read count : 17



Star

Mature

5'	obs	exp	reads	mm	sample
uuugaauaaaauccuuacuccagucagucaguuacaaacagguucucgugagccgugauugaggcguccacucagacagucucgcacaggacuccacacuuaaaaaaagauaaa					
uuugaauaaaauccuuacuccagucagucaguuacaaacagguucucgugagccgugauugaggcguccacucagacagucucgcacaggacuccacacuuaaaaaaagauaaa					
.....((((((((.....)))))).....((((((((((((((((((((.....)))))).....)))))).....)))))).....					
.....aaguucucgugagccgugauug.....			1	0	Mm3
.....gacagucucgcacaggacuc.....			1	0	Mm3
.....gacagucucgcacaggacuccac.....			1	0	Mm3
.....gacagucucgcacaggacuccacU.....			5	1	Mm3
.....aaguucucgugagccgugauug.....			7	0	Mm2
.....aaguucucgugagccgugauuga.....			6	0	Mm2
.....aaguucucgugagccgugauugU.....			2	1	Mm2
.....aaguucucgugagccgugauugaA.....			1	1	Mm2
.....agacagucucgcacaggacucca.....			1	0	Mm2
.....gacagucucgcacaggacuc.....			1	0	Mm2
.....gacagucucgcacaggacucc.....			2	0	Mm2
.....gacagucucgcacaggacucca.....			9	0	Mm2
.....gacagucucgcacaggacuccac.....			6	0	Mm2
.....gacagucucgcacaggacuccacU.....			7	1	Mm2
.....gacagucucgcacaggacuccaca.....			3	0	Mm2
.....agacagucucgcacaggacuccacU.....			1	1	Fm1
.....gacagucucgcacaggacucc.....			6	0	Fm1
.....gacagucucgcacaggacucca.....			25	0	Fm1
.....gacagucucgcacaggacuccac.....			1	1	Fm1
.....gacagucucgcacaggacuccac.....			1	1	Fm1
.....gacagucucgcacaggacuccaU.....			5	1	Fm1
.....gacagucucgcacaggacuccac.....			67	0	Fm1
.....gacagucucgcacaggacuccac.....			1	1	Fm1
.....gacagucucgcacaggacuccaca.....			7	0	Fm1
.....gacagucucgcacaggacuccacU.....			15	1	Fm1
.....gacagucucgcacaggacuccacacu.....			2	0	Fm1
.....gacagucucgcacaggacuccac.....			9	0	Fm2
.....gacagucucgcacaggacuccacacu.....			9	0	Fm2