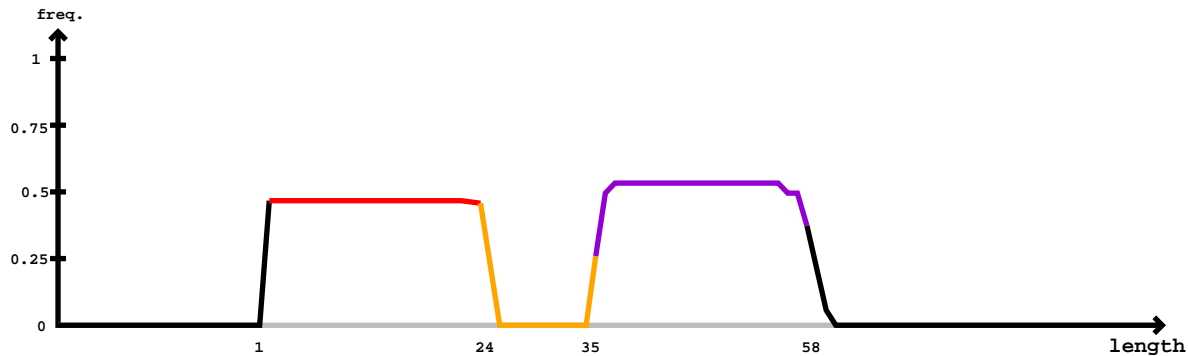
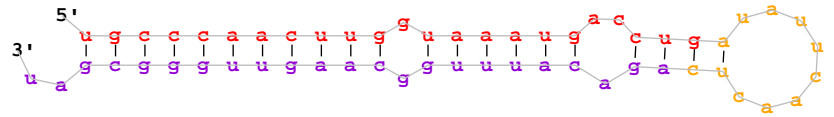


Provisional ID : NMRB01000076.1\_Notospermus\_geniculatus\_strain\_Ushimado\_scaffold76\_whole\_genome\_shotgun\_sequence\_2796  
Score total : 106.8  
Score for star read(s) : 3.9  
Score for read counts : 102.1  
Score for mfe : 1.4  
Score for randfold : 0  
Score for cons. seed : -0.6  
Total read count : 212  
Mature read count : 99  
Loop read count : 0  
Star read count : 113



## Mature Star

5'	caauuugauccuuuagugcuugcccaacuugguaaaugaccugauuucaacucagacauuuggcaaguugggcgauuucuuuggguuugaacucugcugaaacacacaaaaga	-3'	obs	
	caauuugauccuuuagugcuugcccaacuugguaaaugaccugauuucaacucagacauuuggcaaguugggcgauuucuuuggguuugaacucugcugaaacacacaaaaga		exp	
	((((((((.....))))))))).....	reads	mm	sample
	.....ugcccaacuugguaaCugacc.....	1	1	Mm3
	.....ugcccaacuugguaaaugaccuga.....	1	0	Mm3
	.....ugcccaacuugguaaCugaccuga.....	1	1	Mm3
	.....agacauuuggcaaguugggc.....	1	0	Mm3
	.....agacauuuggcaaguugggcga.....	1	0	Mm3
	.....agacauuuggcaaguugggcgau.....	2	0	Mm3
	.....ugcccaacuugguaaCugaccug.....	2	1	Fm3
	.....ugcccaacuugguaaCugaccuga.....	7	1	Fm3
	.....agacauuuggcaaguugggcgau.....	6	0	Fm3
	.....agacauuuggcaaguugggcgaAu.....	2	1	Fm3
	.....agacauuuggcaaguugggcgauA.....	2	1	Fm3
	.....gacauuuggcaaguugggc.....	6	0	Fm3
	.....gacauuuggcaaguugggcga.....	1	1	Fm3
	.....gacauuuggcaaguugggcga.....	12	0	Fm3
	.....gacauuuggcaaguugggcgau.....	15	0	Fm3
	.....gacauuuggcaaguugggcgauu.....	3	0	Fm3
	.....acauuuggcaaguugggcga.....	5	0	Fm3
	.....ugcccaacuugguaaaugaccu.....	1	0	Mm2
	.....ugcccaacuugguaaaugaccug.....	43	0	Mm2
	.....ugcccaacuugguaaCugaccug.....	5	1	Mm2
	.....ugcccaacuugguaaCugaccuga.....	5	1	Mm2
	.....ugcccaacuugguaaaugaccugU.....	8	1	Mm2
	.....ugcccaacuugguaaaugaccuga.....	11	0	Mm2
	.....agacauuuggcaaguugggcgU.....	1	1	Mm2
	.....agacauuuggcaaguugggcga.....	6	0	Mm2
	.....agacauuuggcaaguugggcgau.....	2	0	Mm2
	.....agacauuuggcaaguugggcgauu.....	1	0	Mm2
	.....gacauuuggcaaguugggc.....	1	0	Mm2
	.....gacauuuggcaaguugggcgau.....	3	0	Mm2
	.....ugcccaacuugguaaaugaccug.....	8	0	Mm1

Mature

Star

caauuugauccuuuagugcuugccccaacuugguaaaugaccugauauucaacucagacauuugggcaaguugggcgaauuucuuuggguugaacuugcugaaacacacaaaaga			
.....agacauuugggcaaguugAgcgau.....	1	1	Mm1
.....agacauuugggcaaguugCgcgau.....	1	1	Mm1
.....agacauuugggcaaguugggcgau.....	8	0	Mm1
.....agacauuugggaGguugggcgau.....	4	1	Mm1
.....gacauuugggcaaguugggcgaAu.....	9	1	Mm1
.....ugcccaacuugguaaaugaccugU.....	6	1	Fm2
.....agacauuugggcaaguugggcgauiu.....	12	0	Fm2
.....acauuugggcaaguugggcgaUA.....	3	1	Fm2
.....agacauuugggcaaguugggcgau.....	5	0	Fm1