Provisional ID: NMRB01001071.1_Notospermus_geniculatus_strain_Ushimado_scaffold1071,_whole_genome_shotgun_sequence_24532

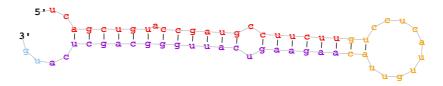
Score total : 3.4

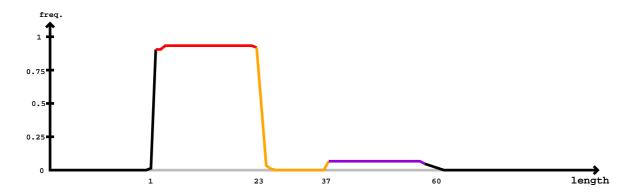
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 1.7
Score for randfold : 0

Score for randfold : 0 Score for cons. seed : 3 Total read count : 210 Mature read count : 196 Loop read count : 0

14

Star read count





5'- ucguucccccugagagaugaucagcuguaccgaugccuucuuguccucauuguuacaagaagucauugggcagcucaugaauucuuggguaaagucgccaguggaauguuua

ucguuccccugagagauga <mark>ucagcuguaccgaugccuucuu</mark> guccucauuguuacaagaagucauugggcagcucaugaauucuuggguaaagucgccaguggaaug	uuua	exp	
.((((((((((((((((((((((((((((((((((((((reads	mm	sample
ucagcuguaccgaugccuucu	1	0	Mm2
ucagcuguaccgaugccuucuu	43	0	Mm2
ucagcuguaccgaugccGucuu	1	1	Mm2
ucagcuguaccgaugccuucuug	8	0	Mm2
ucageuguacegaugceuucuuC	2	1	Mm2
ucageuguacegaugecuucuugu	2	0	Mm2
agcuguaccgaugccuucuugu	1	0	Mm2
.ucagcuguacAgaugccuucuu	1	1	Mm1
	54	0	Mm1
aagaagucauuggAcagcuc.		1	Mm1
aagaagucauuggAcagcuca	6	1	Mm1
ucagcuguaccgaugccCucuu	1	1	Fm1
		1	Fm1
		1	Fm1
		0	Fm1
agcuguaccgaugccuucuuguc		0	Fm1
ageuguaeegaugeeuueuugueaagaagucauuggAcagcuca		1	Fm1
aayaayucauuyyxcaycuca	1	-	FILL
ucagcuguaccgaugccuucuu	11	0	Fm2
aagaagucauugggcagcuca	2	0	Fm2
aucagcuguaccgaugccuucuuUu	1	1	Mm3
ucagcuquaccqauqccuucuu		0	Mm.3
ucagcuguaccgaugccuucuug		0	Mm3
aucagcuguaccgaugccuucu	1	0	Fm3
aucagcuguaccgaugcAuucu	1	1	Fm3
ucagcuguacGgaugccuucuu	1	1	Fm3
ucagcuguacAgaugccuucuu	1	1	Fm3
ucagcuguaccgaugccuGcuu	1	1	Fm3
ucagcuguaccgaugccuucuu	15	0	Fm3

Mature Star

ucguucccccugagagaug <mark>aucagcuguaccgaugccuucuu</mark> guccucauuguuacaagaagucauugggcagcucaugaauucuuggguaaagucgccaguggaauguuua			
ucagcuguaccgaugccuucuug	5	0	Fm3
ucagcuguaccgaugccuucuugG	1	1	Fm3
agcuguaccgaugccuucuug	3	0	Fm3
aagaagucauuggAcagcuca	1	1	Fm3