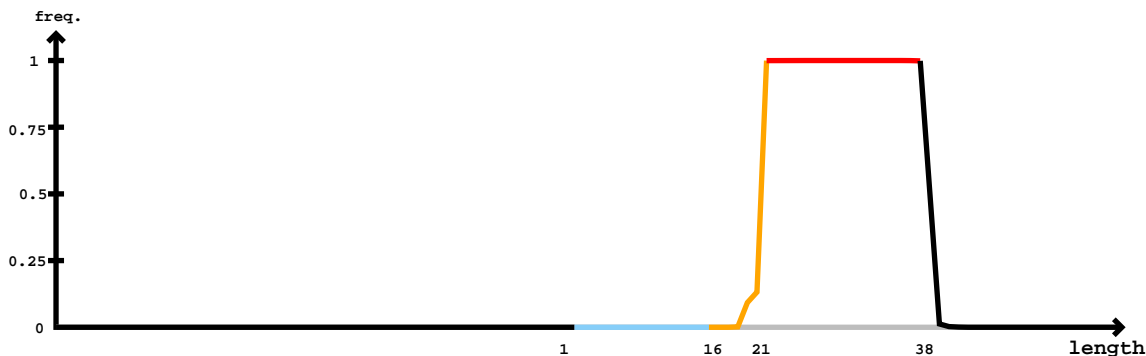
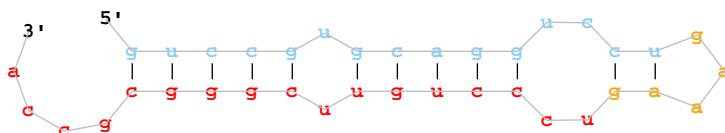


Provisional ID : DRO_25037_4505
 Score total : 2.5
 Score for star read(s) : -1.3
 Score for read counts : 0
 Score for mfe : 2.2
 Score for randfold : 1.6
 Score for cons. seed :
 Total read count : 25871
 Mature read count : 25871
 Loop read count : 0
 Star read count : 0



Star Mature

5' - cucccgguucgccauggaugaagccguggcgaugguggcucgaagccucug guccgugcagguccugaaguccugucggggcgcca gaugucgggggcuuaaug -3'	exp		
(((((((.....)))))).....((((.....))))(((((((.....)))))).....)))).....	reads	mm	sample
.....aaguccugucggggcgc.....	1	0	seq
.....Caguccugucggggcgcca.....	15	1	seq
.....aaguccugucggggcgcca.....	20	0	seq
.....Cguccugucggggcgc.....	1	1	seq
.....aguccugucggggcgc.....	7	0	seq
.....aguccugucggggcgcc.....	1	0	seq
.....Cguccugucggggcgcca.....	10	1	seq
.....aguccugucggggcgcca.....	2330	0	seq
.....aguccugucggggcgccG.....	3	1	seq
.....agGccugucggggcgcca.....	2	1	seq
.....aguccugucggggcAcca.....	1	1	seq
.....aguccugucggggcgUa.....	1	1	seq
.....aguccugucggggcgGcGa.....	1	1	seq
.....Cuccugucggggcgcc.....	1	1	seq
.....guccugucggggcgcc.....	22	0	seq
.....guccugucggggcgcca.....	981	0	seq
.....guccugGucggggcgcca.....	1	1	seq
.....guccugucggggcCcca.....	1	1	seq
.....Cuccugucggggcgcca.....	2	1	seq
.....guccugucggggcgGca.....	1	1	seq
.....guccAguucggggcgcca.....	1	1	seq
.....guccugucgUgcgcca.....	1	1	seq
.....Auccugucggggcgcca.....	2	1	seq
.....guccugucgggAcgcca.....	1	1	seq
.....guccugucggggcUcca.....	1	1	seq
.....guccugucggggcgccC.....	1	1	seq
.....guccugucggggcgccG.....	7	1	seq
.....guccuguuGggggcgcca.....	1	1	seq
.....guccugucggggcgccaA.....	1	1	seq
.....guccugucggggcgccaU.....	1	1	seq
.....uccugucUggggcgcca.....	1	1	seq
.....uccugCucggggcgcca.....	4	1	seq
.....uUccugucggggcgcca.....	3	1	seq
.....uccugucgggAcgcca.....	1	1	seq

Star Mature

cucccgguucgccauggaugaagccgugggcgaugugggcucgaagccucug	guccgugcagguccu	gaaguccugucggggcgcca	gaugucgggggcuuaaug			
.....	Ccccguucggggcgcca.....	1	1	seq
.....	ucccgguucggggcUcca.....	1	1	seq
.....	ucccAguucggggcgcca.....	2	1	seq
.....	ucccgGucggggcgcca.....	2	1	seq
.....	uAccguucggggcgcca.....	2	1	seq
.....	ucccguuAgggcgcca.....	3	1	seq
.....	ucGcguucggggcgcca.....	1	1	seq
.....	ucUcguucggggcgcca.....	1	1	seq
.....	ucccgguucgggUcgcca.....	1	1	seq
.....	Gcccguucggggcgcca.....	8	1	seq
.....	ucccgguucggggcgcU.....	70	1	seq
.....	uccAuguucggggcgcca.....	3	1	seq
.....	ucccGguucggggcgcca.....	3	1	seq
.....	ucccguuUggggcgcca.....	2	1	seq
.....	ucccgguucAgggcgcca.....	8	1	seq
.....	ucccgGcggggcgcca.....	5	1	seq
.....	ucccNguucggggcgcca.....	2	1	seq
.....	ucccgguucggggcAcca.....	4	1	seq
.....	Acccgguucggggcgcca.....	2	1	seq
.....	ucccgguucggggcgcUa.....	2	1	seq
.....	ucccgguucggggcgcca.....	21741	0	seq
.....	ucccgguucggggcgcG.....	89	1	seq
.....	ucccgguucgAgcgcca.....	4	1	seq
.....	ucccAuuucggggcgcca.....	1	1	seq
.....	ucccuUuucggggcgcca.....	1	1	seq
.....	ucccgAucggggcgcca.....	2	1	seq
.....	ucccgguucCggcgcca.....	3	1	seq
.....	ucccgguucggggcgcGa.....	8	1	seq
.....	ucccgguucggggcgcC.....	2	1	seq
.....	ucccCguucggggcgcca.....	2	1	seq
.....	ucccgguucgggNogcca.....	1	1	seq
.....	uccGugucggggcgcca.....	11	1	seq
.....	uccUguucggggcgcca.....	1	1	seq
.....	ucccgguucggggcgcag.....	17	0	seq
.....	ucccgguucggggcgccaA.....	81	1	seq
.....	ucccgguucggggcgccaU.....	65	1	seq
.....	ucccgguucggggcgccaAa.....	101	1	seq
.....	ucccgguucggggcgccagC.....	4	1	seq
.....	ucccgguucggggcgccaga.....	140	0	seq
.....	ucccgguucggggcgccaUa.....	8	1	seq
.....	ucccgguucggggcgcUga.....	3	1	seq
.....	ucccgguucggggcgccagau.....	5	0	seq
.....	ucccgguucggggcgccaUau.....	5	1	seq
.....	ucccgguucggggcgccagCu.....	1	1	seq
.....	ucccgguucggggcgccagaA.....	10	1	seq
.....	ucccgguucggggcgccaAau.....	25	1	seq
.....	ucccgguucggggcgccagaAg.....	10	1	seq
.....	ucccgguucggggcgccagauA.....	5	1	seq
.....	cccguucggggcgccaA.....	2	1	seq
.....	cccguucggggcgccaU.....	1	1	seq
.....	ccguucggggcgccaUa.....	3	1	seq
.....	cguucggggcgccaUau.....	1	1	seq
.....	cguucggggcgccaUaug.....	1	1	seq
.....	uguucggggcgccaUaug.....	1	1	seq
.....	uguucggggcgccagaAg.....	1	1	seq