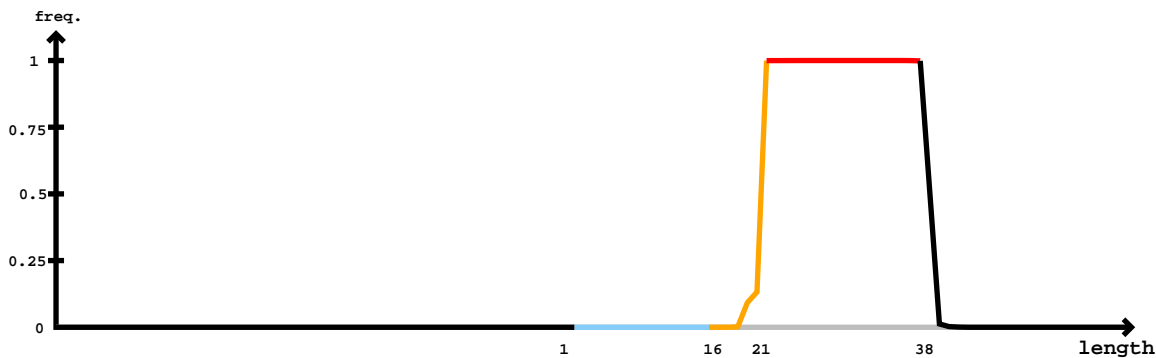
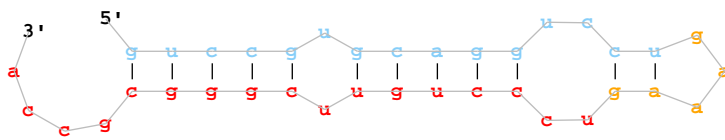


Provisional ID : DRO_25037_4509
 Score total : 2.5
 Score for star read(s) : -1.3
 Score for read counts : 0
 Score for mfe : 2.2
 Score for randfold : 1.6
 Score for cons. seed :
 Total read count : 25871
 Mature read count : 25871
 Loop read count : 0
 Star read count : 0



Star
Mature

5' -	-3'	exp	reads	mm	sample
cccccgggucugccauggaugaagccgugggcgaugguggcugaagccucuguccgugcaggucuccugaagucccugucggggcgccagaugucggggggcuuuaacac					
.....aagucccugucggggcgcc.....			1	0	seq
.....Cagucccugucggggcgcc.....			15	1	seq
.....aagucccugucggggcgcc.....			20	0	seq
.....agucccugucggggcgcc.....			7	0	seq
.....Cgucccugucggggcgcc.....			1	1	seq
.....agucccugucggggcgcc.....			1	0	seq
.....agucccugucggggcgccUa.....			1	1	seq
.....agucccugucggggcgcc.....			2330	0	seq
.....agucccugucggggcgccGa.....			1	1	seq
.....Cgucccugucggggcgcc.....			10	1	seq
.....agGcccgucggggcgcc.....			2	1	seq
.....agucccugucggggcgccG.....			3	1	seq
.....agucccugucggggcgccA.....			1	1	seq
.....Cucccugucggggcgcc.....			1	1	seq
.....gucccugucggggcgcc.....			22	0	seq
.....gucccugucgggAcgcca.....			1	1	seq
.....gucccugucggggcgcc.....			1	1	seq
.....gucccugucggggcgccG.....			7	1	seq
.....Cucccugucggggcgcc.....			2	1	seq
.....gucccugGuccggggcgcc.....			1	1	seq
.....gucccugucggggcgcc.....			981	0	seq
.....Aucccugucggggcgcc.....			2	1	seq
.....gucccugucggggcgccC.....			1	1	seq
.....guccAuccggggcgcc.....			1	1	seq
.....gucccugucgggUgcca.....			1	1	seq
.....gucccugucggggcgcc.....			1	1	seq
.....gucccugucggggcgccUcca.....			1	1	seq
.....gucccugucggggcgccCcca.....			1	1	seq
.....gucccugucggggcgccU.....			1	1	seq
.....gucccugucggggcgccA.....			1	1	seq
.....Ccccgucggggcgcc.....			1	1	seq
.....ucccuAuccggggcgcc.....			1	1	seq
.....ucccNguccggggcgcc.....			2	1	seq
.....ucccugCuccggggcgcc.....			4	1	seq

Star Mature

ccccgggucugccaugggaagccgugggcgaugggcuugaagccucug	uccUguuucgggcgcca	1	1	seq
.....
.....	ucccuguucggNcgcca	1	1	seq
.....	ucccuguucgggcccC	2	1	seq
.....	uccAguuucgggcgcca	3	1	seq
.....	ucccuguucAgggcgcca	8	1	seq
.....	ucccGguucgggcgcca	3	1	seq
.....	ucccuguuAgggcgcca	3	1	seq
.....	ucccuguucgggcccG	89	1	seq
.....	Acccuguucgggcgcca	2	1	seq
.....	uUccuguucgggcgcca	3	1	seq
.....	ucccCguucgggcgcca	2	1	seq
.....	ucccGguucgggcgcca	2	1	seq
.....	ucccuguucgggAcca	4	1	seq
.....	ucccuguucgggUcca	1	1	seq
.....	ucccuguucgggcccU	70	1	seq
.....	ucccuguucgggcccGa	8	1	seq
.....	ucccugGuucgggcgcca	2	1	seq
.....	uccGguucgggcgcca	11	1	seq
.....	ucccuguucgggAcgcca	1	1	seq
.....	ucccuguucUggcgcca	1	1	seq
.....	ucccuguGcgggcgcca	5	1	seq
.....	uAccuguucgggcgcca	2	1	seq
.....	ucccuguucCggcgcca	3	1	seq
.....	ucccuguucggUcgcca	1	1	seq
.....	Gcccuguucgggcgcca	8	1	seq
.....	ucccuguucgAgcgcca	4	1	seq
.....	ucccuUuucgggcgcca	1	1	seq
.....	ucGeuguucgggcgcca	1	1	seq
.....	ucccuguuUgggcgcca	2	1	seq
.....	ucccuguucgggcccTa	21741	0	seq
.....	ucccuguucgggcccUa	2	1	seq
.....	ucUcuguucgggcgcca	1	1	seq
.....	ucccAguucgggcgcca	2	1	seq
.....	ucccuguucgggcccag	17	0	seq
.....	ucccuguucgggcccU	65	1	seq
.....	ucccuguucgggcccA	81	1	seq
.....	ucccuguucgggcccagC	4	1	seq
.....	ucccuguucgggcccUa	8	1	seq
.....	ucccuguucgggcccaga	140	0	seq
.....	ucccuguucgggcccAa	101	1	seq
.....	ucccuguucgggcccUga	3	1	seq
.....	ucccuguucgggcccUau	5	1	seq
.....	ucccuguucgggcccagaA	10	1	seq
.....	ucccuguucgggcccagau	5	0	seq
.....	ucccuguucgggcccAau	25	1	seq
.....	ucccuguucgggcccagCu	1	1	seq
.....	ucccuguucgggcccagauA	5	1	seq
.....	ucccuguucgggcccagaAg	10	1	seq
.....	cccuguucgggcccU	1	1	seq
.....	cccuguucgggcccA	2	1	seq
.....	ccuguucgggcccUa	3	1	seq
.....	cuguucgggcccUau	1	1	seq
.....	cuguucgggcccUaug	1	1	seq
.....	uguucgggcccUaug	1	1	seq
.....	uguucgggcccagaAg	1	1	seq