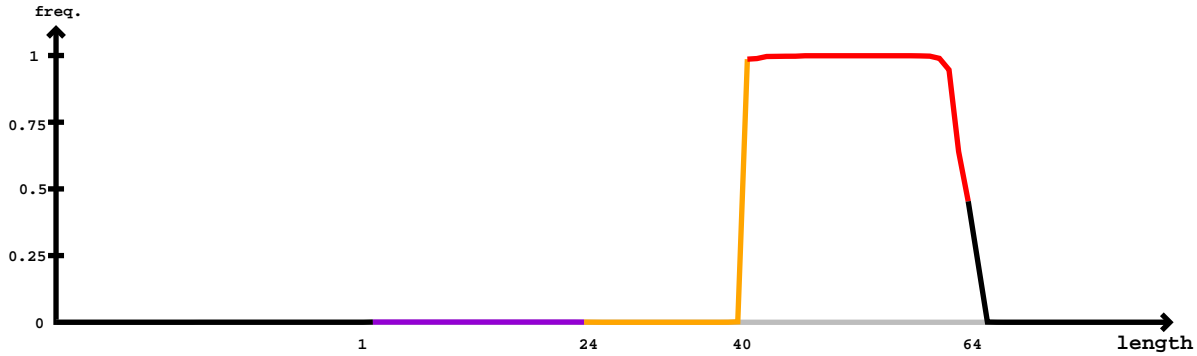
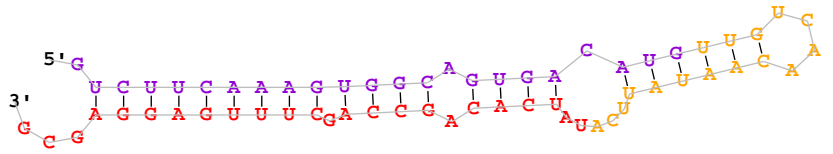


Provisional ID : 2L_dna:chromosome_chromosome:BDGP6:2L:1:23513712:1_REF_2242
Score total : 14685.6
Score for star read(s) : 3.9
Score for read counts : 14676
Score for mfe : 1.1
Score for randfold : 1.6
Score for cons. seed : 3
Total read count : 28798
Mature read count : 28775
Loop read count : 0
Star read count : 23



Star

Mature

5' -	obs	exp	known	reads	mm	sample
gugcaaaucuaaaacuaaaaaacucaacugucuucaaaaguggcagugacauguuugucaacaauuucauaucacagccagcuuugaggagcgguugcgguuuuucuguccgg	-3'	obs		1	0	seq
gugcaaaucuaaaacuaaaaaacucaacugucuucaaaaguggcagugacauguuugucaacaauuucauaucacagccagcuuugaggagcgguugcgguuuuucuguccgg	exp			2	0	seq
gugcaaaucuaaaacuaaaaaacucaacugucuucaaaaguggcagugacauguuugucaacaauuucauaucacagccagcuuugaggagcgguugcgguuuuucuguccgg	known			1	0	seq
.....(((((((.....((((((((((((.....)))))).....)))))).....)))))).....)))))).....)))))).....)))))).....)))))).....				1	1	seq
.....uaacuaaaaaacucaacu.....				1	1	seq
.....gucuucaaaaguggcagugaca.....				1	1	seq
.....gucuucaaaagugAcagugacau.....				1	1	seq
.....gucuucaaaaguggcagugaca.....				2	0	seq
.....gucuucaaaaguggcagugGcau.....				1	1	seq
.....gucuucaaaaguggcagGgacaug.....				1	1	seq
.....gucuucaaaaguggcagGgacaug.....				1	1	seq
.....gucuucaaaaguggcagugacaug.....				12	0	seq
.....gucuucaaaaguggcagugaGaug.....				1	1	seq
.....gucuucaaaaguggcagugGcaug.....				2	1	seq
.....gucuucaaaagugAgcagugacaug.....				1	1	seq
.....uucauaucaacagccagcuuugaggg.....				1	0	seq
.....Aauaucaacagccagcuuugaggag.....				1	1	seq
.....Aauaucaacagccagcuuugaggagcg.....				3	1	seq
.....Guaucaacagccagcuuugaggga.....				1	1	seq
.....auaucaacagccagcuuugaggag.....				6	0	seq
.....auaucaacagccagcuuugaggagA.....				1	1	seq
.....auaucaacagcAagcuuugaggag.....				1	1	seq
.....auaucaacagccagcuuugaggagc.....				7	0	seq
.....Guaucaacagccagcuuugaggagc.....				3	1	seq
.....auaucaacagccagcuuugaggagG.....				1	1	seq
.....auaucaacagccagcuuugaggagcg.....				20	0	seq
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.....uaucaUagccagcuuuga.....				1	1	seq
.....uaucaacagccagcuGugag.....				1	1	seq
.....uaucaacagccagcuuugaA.....				1	1	seq
.....uaCcaacagccagcuuugag.....				1	1	seq

Star

Mature

gugcaaaucuuuaacuaaaacuuaacugucuucaaguggcagugacauguuugucaacaauuucuaucacagccagcuuugaggagcguuugcgguuuuucucgucgg

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.....uaucaacagccagcuuugag.....	24	0	seq
.....Aaucaacagccagcuuugagg.....	5	1	seq
.....uaucaacagccagcCuuugagg.....	3	1	seq
.....uaucaacagccagcuuugagg.....	208	0	seq
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.....uCucaacagccagcuuugagg.....	7	1	seq
.....uaucaacagccagcuuugaCg.....	1	1	seq
.....uaucaacagccGgcuuugagg.....	2	1	seq
.....uaucaUagccagcuuugagg.....	1	1	seq
.....uaucaacagccagcuuugCgg.....	1	1	seq
.....uaucaacagccagcCuuugagg.....	1	1	seq
.....uaucaacagcUagcuuugagg.....	1	1	seq
.....uaucaacagccaCuuugagg.....	1	1	seq
.....uaucaacagccagcuuugaAg.....	2	1	seq
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.....uaucaacGgcccagcuuugagga.....	1	1	seq
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Star

Mature

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.....uaucaacagccagcUuugaggagC.....	6	1	seq
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Star

Mature

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.....uaucaagccagcCuuugaggagc.....	2	1	seq
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Star

Mature

gugcaaaucuaaaacuaaaacucaacugucuucaaguggcagugacauguuugucaacaauuucuaucacagccagcuuugaggagcguuugcgguuuuucucgucgg

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.....uUucacagccagcuuugaggagcg.....	1	1	seq
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.....uaucaagccagcuuugagUagcg.....	26	1	seq
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.....aucagccagcuuugaggag.....	14	0	seq

Star

Mature

Sequence	Count	Count	Label
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.....aucacagccagcuuugaggagU.....	1	1	seq
.....aucacagccagcuuugaggagcg.....	43	0	seq
.....aucacagccagcuuugaggagcgu.....	2	0	seq
.....ucacagccagcuuugaAga.....	1	1	seq
.....ucacagccagcuuugaggga.....	8	0	seq
.....ucacagccagcuuugagggaA.....	1	1	seq
.....ucacagccagcuuugaAgag.....	1	1	seq
.....ucacagccagcuuugagggaU.....	1	1	seq
.....Acacagccagcuuugaggag.....	1	1	seq
.....ucacagccagcuuugaggag.....	50	0	seq
.....ucacagccagcuuugGggag.....	1	1	seq
.....ucacagccagcuuugaggagc.....	42	0	seq
.....Acacagccagcuuugaggagc.....	2	1	seq
.....ucacagccagcuuugaCgagc.....	1	1	seq
.....ucacagccagcuuugaAgagc.....	5	1	seq
.....ucGcagccagcuuugaggagcg.....	1	1	seq
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.....ucacagccagcuuugaggagcU.....	4	1	seq
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.....ucacagccagcuuugCggagcg.....	2	1	seq
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.....ucacagccagcuuugaggagcg.....	70	0	seq
.....ucacagccagcuuCgaggagcg.....	1	1	seq
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.....ucacaUccagcuuugaggagcgu.....	1	1	seq
.....ucacagccagcuuugaggagcgAu.....	1	1	seq
.....ucacagccagcuuugaggagcguu.....	1	0	seq
.....cacagccagcuuugaAgag.....	1	1	seq
.....cacagccagcuuugaggag.....	4	0	seq
.....cacagccagcuuugaggagc.....	1	0	seq
.....cacagccagcuuugaggagcg.....	8	0	seq
.....acagccagcuuugaggagc.....	1	0	seq
.....acagccagcuuugaggagA.....	1	1	seq
.....acagccagcuuugaggagcg.....	9	0	seq
.....acagccagcuuugaggagcguA.....	1	1	seq
.....acagccagcuuugaggagcguuU.....	1	1	seq
.....cagccagcuuugaggagc.....	2	0	seq
.....caAccagcuuugaggagcg.....	1	1	seq
.....cagccagcuuugaggagcg.....	1	0	seq
.....agccagcuuugaggagcg.....	46	0	seq
.....agccagcuuugaggagCgcg.....	2	1	seq
.....agccagcuuugaggagcgu.....	1	0	seq