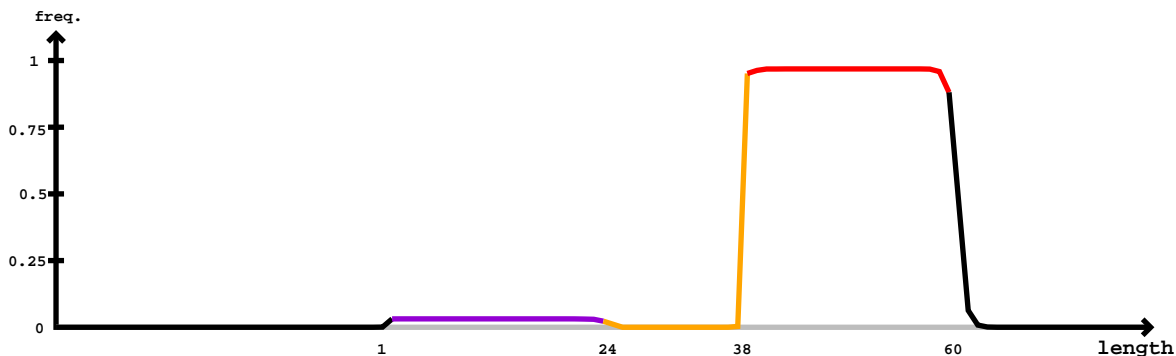
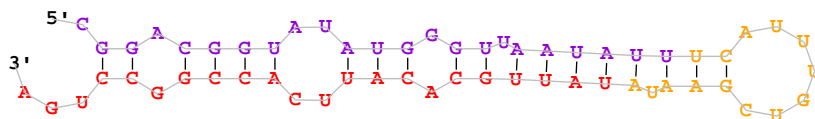


Provisional ID : 2R_dna:chromosome_chromosome:BDGP6:2R:1:25286936:1_REF_11456
 Score total : 3491.8
 Score for star read(s) : 3.9
 Score for read counts : 3488.9
 Score for mfe : -1.6
 Score for randfold : -2.2
 Score for cons. seed : 3
 Total read count : 6855
 Mature read count : 6639
 Loop read count : 0
 Star read count : 216



Star

Mature

5'	obs	exp	known	reads	mm	sample
acggugaaaaucgaaaauucagaucauuuuucggacgguaauuggguuaauuuucauuuugucgaaauauuugcacaauccacggccuga	1	1		1	1	seq
acggugaaaaucgaaaauucagaucauuuuucggacgguaauuggguuaauuuucauuuugucgaaauauuugcacaauccacggccuga	3	0		3	0	seq
.....ucggacgguaauuggguCaau.....	1	0		1	0	seq
.....cggacgguaauuggguuaa.....	2	0		2	0	seq
.....cggacgguaauuggguCaau.....	4	1		4	1	seq
.....cggacgguaauuggguCaau.....	21	1		21	1	seq
.....cggacgguaauuggguuaauuu.....	25	0		25	0	seq
.....cggacgguaauuggguCaau.....	3	1		3	1	seq
.....cggacgguaauuggguCaau.....	2	1		2	1	seq
.....cggacgguaCauggguuaauuu.....	1	1		1	1	seq
.....cggacgguaauuggguCaau.....	81	1		81	1	seq
.....cggacgguaauuggguuaauuu.....	1	1		1	1	seq
.....cggacgguaauuggguuaauuu.....	59	0		59	0	seq
.....cggacgguaauuggguuaauuu.....	7	1		7	1	seq
.....cggacgguaauuggguCaau.....	1	1		1	1	seq
.....cggacgguaauuggguuaauuu.....	1	0		1	0	seq
.....Gauauugcacaauccacggccu.....	1	1		1	1	seq
.....auauugcacaauccacggccug.....	4	0		4	0	seq
.....auCuugcacaauccacggccuga.....	1	1		1	1	seq
.....auauugcacaauccacggcUuga.....	1	1		1	1	seq
.....auauugcacaauccacggccuga.....	8	0		8	0	seq
.....auuuCcauuaccacggccuga.....	1	1		1	1	seq
.....auauugcacaauccacggccugaU.....	1	1		1	1	seq
.....auauugcacaauccacggccugaa.....	3	0		3	0	seq
.....auauugcacaauccacggccugaaaa.....	1	0		1	0	seq
.....uauugcacaauccacggcG.....	1	1		1	1	seq
.....uauugcacaauccacggcc.....	4	0		4	0	seq
.....uauugcacaauccacgUccu.....	1	1		1	1	seq
.....uauugcacaauccacgAgccu.....	1	1		1	1	seq
.....Auauugcacaauccacggccu.....	1	1		1	1	seq

Star

Mature

acggugaaaaucgaaauuucgagcauuuuucggacgguaauaggguuauuuucauuugucgaaauuauugcacaauccaccggccugaaaauucaagaauuaacauc

.....uauugcacaauccacUggccu.....	1	1	seq
.....Cauugcacaauccaccggccu.....	2	1	seq
.....uauugcacaauccaccggccu.....	55	0	seq
.....uauugcacaauAaccggccu.....	1	1	seq
.....uauugcacaauccaccggccuA.....	2	1	seq
.....uauugcacaauccacUggccug.....	1	1	seq
.....uauugcacaauccaccAgccug.....	2	1	seq
.....Cauugcacaauccaccggccug.....	1	1	seq
.....uauugcacaauccaccgUccug.....	4	1	seq
.....uauugcacaauccaccggccuU.....	1	1	seq
.....uauugcacaaucaUcggccug.....	1	1	seq
.....uauugcacaauAaccggccug.....	1	1	seq
.....uauugcacaauccaccggAcug.....	1	1	seq
.....uauugcacaauccaccAgggccug.....	2	1	seq
.....uauugcacaauccaccUggccug.....	1	1	seq
.....uauugcacaauccaccggccug.....	1	1	seq
.....uauugcGcauuccaccggccug.....	1	1	seq
.....uauugcAauuccaccggccug.....	1	1	seq
.....uauuAcacaauccaccggccug.....	1	1	seq
.....uauCgcacaauccaccggccug.....	1	1	seq
.....uauugcacaauccaccggccug.....	1	1	seq
.....uauugcacaauccaccggccug.....	498	0	seq
.....uauuUcacaauccaccggccug.....	4	1	seq
.....uauugcacaauccaccggcAug.....	1	1	seq
.....uauugcaUauuccaccggccuga.....	7	1	seq
.....uCuugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccgCccuga.....	3	1	seq
.....uauugcacaauccaccggcGuga.....	3	1	seq
.....uauugcacaauccaccggccuga.....	7	1	seq
.....uauuUcacaauccaccggccuga.....	2	1	seq
.....uauugcacaauUaccggccuga.....	4	1	seq
.....uaGugcacaauccaccggccuga.....	5	1	seq
.....Cauugcacaauccaccggccuga.....	8	1	seq
.....uauugcacaauccaccggcAuga.....	12	1	seq
.....uauugcUcauuccaccggccuga.....	1	1	seq
.....uauugcacaauccaccAgggccuga.....	5	1	seq
.....uauugcacaauccaccggccuga.....	2	1	seq
.....uauugcacaauCcaccggccuga.....	4	1	seq
.....uauugcacaauGcaccggccuga.....	2	1	seq
.....uauuAcacaauccaccggccuga.....	3	1	seq
.....uauugcacaauccaccggccAuga.....	1	1	seq
.....uauugcacaauccaccggccuga.....	12	1	seq
.....uauugcacaauccaccgAccuga.....	2	1	seq
.....Gauugcacaauccaccggccuga.....	2	1	seq
.....uauugcacaCucaccggccuga.....	8	1	seq
.....uauugcacaaucaUcggccuga.....	10	1	seq
.....uauugcacCuuccaccggccuga.....	1	1	seq
.....uauugcacaauccaccUggccuga.....	1	1	seq
.....uauugcacaauccaccCggccuga.....	5	1	seq
.....uauugcacaauccaccggccGga.....	5	1	seq
.....uauugcacGuuccaccggccuga.....	4	1	seq
.....uauGgcacaauccaccggccuga.....	1	1	seq
.....uauugUacaauccaccggccuga.....	2	1	seq
.....uauugcacaauccaccgUccuga.....	12	1	seq
.....uauugcacaauccaccUggccuga.....	4	1	seq
.....uauAgcacaauccaccggccuga.....	2	1	seq
.....uauugcacaauccaccggccuCa.....	2	1	seq
.....uauugcacaauccaccggccugG.....	8	1	seq
.....uGuugcacaauccaccggccuga.....	3	1	seq
.....uauugcacaauccaccggUcuga.....	5	1	seq
.....uauugcacaauccaccggccuUa.....	8	1	seq
.....uauugcacaauccaccggccugC.....	9	1	seq
.....uaCugcacaauccaccggccuga.....	5	1	seq
.....uauugcacaauccaccggccuga.....	4429	0	seq
.....uauugcacaauccaccggccuAa.....	5	1	seq
.....uauugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccggccugU.....	30	1	seq
.....uauCgcacaauccaccggccuga.....	9	1	seq
.....uauugcacaauccaccggAcuga.....	25	1	seq
.....uauugcacaauccaccggcUuga.....	7	1	seq

Star

Mature

acggugaaaaucgaaauucugaucuuuuucggacggguauaggguuauuuucauuugucgaaauuauugcacaauccaccggccuga

.....uauugcacaAucaccggccuga.....	1	1	seq
.....uauugAacaauccaccggccuga.....	2	1	seq
.....uauugcacaauccaccggccCga.....	7	1	seq
.....uauugcacUuucaccggccuga.....	1	1	seq
.....uauuCcacaauccaccggccuga.....	1	1	seq
.....uUuugcacaauccaccggccuga.....	3	1	seq
.....uauugcGcauucaccggccuga.....	9	1	seq
.....Aauugcacaauccaccggccuga.....	27	1	seq
.....uauugGacaauccaccggccuga.....	3	1	seq
.....uauugcacaaucaGcggccuga.....	2	1	seq
.....uauugcaGauucaccggccuga.....	1	1	seq
.....uaAugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccAgccuga.....	9	1	seq
.....uauugcacaauccaccUgcccuga.....	1	1	seq
.....uauugcacaauccaccAgccuga.....	1	1	seq
.....uauugcacaauccaccggcGugaa.....	1	1	seq
.....uauugcacaauccaccggccuAaa.....	1	1	seq
.....uauugcacaauccaccggccugaC.....	7	1	seq
.....uauugcacaauccaccggccugCa.....	3	1	seq
.....uauugcacaauUaccggccuga.....	1	1	seq
.....uauugcacaauccaccggccugaN.....	3	1	seq
.....Aauugcacaauccaccggccuga.....	13	1	seq
.....uauugcacaauccaccggccugUa.....	11	1	seq
.....uauuAcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccggccugaU.....	74	1	seq
.....uauugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccggccuga.....	1	1	seq
.....uauCgcacaauccaccggccuga.....	3	1	seq
.....uauugcacaauccaccggccugGa.....	1	1	seq
.....Cauugcacaauccaccggccuga.....	2	1	seq
.....uauugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccAgccuga.....	2	1	seq
.....uauugcacaauccaccggccuga.....	595	0	seq
.....uauugcacaauCcaccggccuga.....	2	1	seq
.....uauuUcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccgAccuga.....	2	1	seq
.....uauugcacaauccaccCgcccuga.....	1	1	seq
.....uauugcacaauccaccggcUugaa.....	1	1	seq
.....uauugcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccggUcugaa.....	1	1	seq
.....uauugcacaauccaccggccCgaa.....	2	1	seq
.....uauugcacaauccaccgCccuga.....	1	1	seq
.....uauugcacaauccaccggccugaG.....	3	1	seq
.....uauuCcacaauccaccggccuga.....	1	1	seq
.....uauugcacaauccaccUggccuga.....	2	1	seq
.....uauugcacaCucaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccAgccugaaa.....	2	1	seq
.....uauugcacaauccaccggccugaUa.....	15	1	seq
.....uauugcacaauccaccggccugaU.....	9	1	seq
.....Aauugcacaauccaccggccugaaa.....	3	1	seq
.....uauugcGcauucaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccggccugaCa.....	2	1	seq
.....uauugcacaauccaccggccugaaa.....	312	0	seq
.....uaCugcacaauccaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccgUcccugaaa.....	1	1	seq
.....uauugcacaauccaccUgcccugaaa.....	1	1	seq
.....uauugUacaauccaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccggccugCaa.....	1	1	seq
.....uauugcacaauccaccggccugaaC.....	3	1	seq
.....uauugcacaauccaccggccugUaa.....	1	1	seq
.....uauugcacaauccaccggAcugaaa.....	1	1	seq
.....uauugcacaauccaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccggccugaaG.....	2	1	seq
.....Cauugcacaauccaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccggccuUaaa.....	2	1	seq
.....uauugcacaauccaccggccugaUaa.....	1	1	seq
.....uauugcacaauccaccggccugaaaU.....	1	1	seq
.....uauugcacaauccaccggccugaaa.....	1	1	seq
.....uauugcacaauccaccggccugaaaC.....	1	1	seq

Star

Mature

acggugaaaaucgaaaauucuaagaucauuuuucggacgguaauaggguaauuuucauuugucgaaauauugcacaucacccggccuga	aaaaaucaagaauaacauc			
.....uauugcacaucacccggAcugaaaa.....	2	1		seq
.....uauugcacaauUaccggccugaaaa.....	1	1		seq
.....uauugcacaucacccggccugaaaa.....	40	0		seq
.....uauugcacaucacccggccugaaaaA.....	3	1		seq
.....uauugcacaucacccggccugaaaaC.....	1	1		seq
.....aGugcacaucacccggccug.....	1	1		seq
.....auugcacaucacccGccuga.....	1	1		seq
.....auugcacaucacccggccuga.....	37	0		seq
.....auugcacaucacccggccuUa.....	1	1		seq
.....auugcacaucacccggccugaa.....	17	0		seq
.....auugcacaucacccggccugaU.....	7	1		seq
.....auugcacaucacccggccugaUa.....	2	1		seq
.....auugcacaucacccggccugaaa.....	10	0		seq
.....auugcacaucacccggccugaaaa.....	1	0		seq
.....auugcacaucacccggccugaaaaC.....	1	1		seq
.....auugcacaucacccggccugaaaaA.....	1	1		seq
.....uugcacaucacccggccug.....	3	0		seq
.....uugcacaucacccggccuAa.....	1	1		seq
.....uugcacaucacccggccuga.....	25	0		seq
.....uugcacaGucacccggccuga.....	1	1		seq
.....uugcacaucacccggccugaa.....	2	0		seq
.....uugcacaucacccggccugaaa.....	1	0		seq
.....uGcacaucacccggccugaaa.....	1	1		seq
.....uugcacaucacccggccugaUa.....	1	1		seq
.....uugcacaucacccggccugaUaa.....	1	1		seq
.....gcacaucacccggccuga.....	2	0		seq