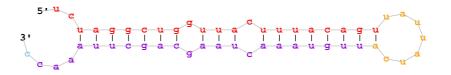
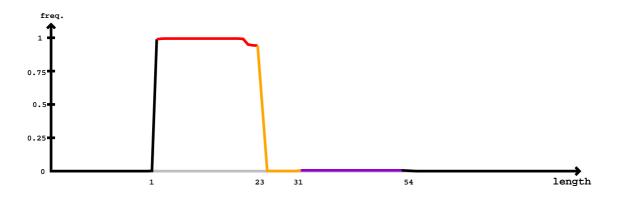
Provisional ID : chromosome_10_46388

Score total : 682.2 Score for star read(s) : 3.9 Score for read counts : 675.6 Score for mfe : 1.1 Score for randfold : 1.6 Score for cons. seed :

Total read count : 1337
Mature read count : 1330
Loop read count : 0
Star read count : 7





Mature	Star
--------	------

uuaugccauugcugauucuaggcugguuacuuuacaguuauuaucauuguaaacuaagcagcuuaaaccucaguuguagugauauuauuacaguuuacuuagga	exp		
((((((((((((((((((((((((((((((((((reads	mm	sample
Cucuaggcugguuacuuuacagu.	2	1	seq
ucuaggcugguuacuuua	4	0	seq
ucuaggcugguuaUuuuac	1	1	seq
ucuaggcugguuacuCuac	3	1	seq
ucuaggcugguuacuuuac	51	0	seq
ucuaggcugguuacuuuaU	1	1	seq
ucuaggcugguuacuuuaca	7	0	seq
ucuaggcugguuacuuuacag	2	0	seq
ucuaggcugguuacCuuacagu	4	1	seq
ucuGggcugguuacuuuacagu	9	1	seq
ucuaggcugguuaUuuuacagu	1	1	seq
ucuaggcugguuacuuuacCgu	1	1	seq
ucuaggcugguuacuuGacagu	1	1	seq
ucuaggcugguAacuuuacagu	1	1	seq
ucuaggcuUguuacuuuacagu	1	1	seq
ucuaggcugguuacuuuacagG	5	1	seq
ucCaggcugguuacuuuacagu.	1	1	seq
ucuaggcugguuacuCuacagu.	6	1	seq
ucuaggcugguuUcuuuacagu.	1	1	seq
ucuaggcugguuacuuuacagA	17	1	seq
ucuaggcugguuacuAuacagu	2	1	seq
ucAaggcugguuacuuuacagu.	1	1	seq
ucuaggcGgguuacuuuacagu	2	1	seq
ucuaggcugguuacuuuacGgu	2	1	seq
	2	1	seq
ucuaggcugAuuacuuuacagu	3	1	seq
ucuaggcugguuaGuuuacagu	1	1	seq
	4	1	seq
	1	1	seq
	4	1	-
			seq
ucuaggcuAguuacuuuacagu	1	1	seq
ucuaggcugguuaAuuuacagu	1	1	seq
	587	0	seq

Mature Star

Mature	Star			
uuaugccauugcugaguucuaggcugguuacuuuacaguuauuau	cauuguaaacuaagcagcuuaaaccucaguuguagugauauuauuacaguuuacuuagga			
Ccuaggcugguuacuuuacagu		2	1	seq
UcuaggcugUuuacuuuacagu		1	1	seq
ucuaggcugguuacuuuacagC		94	1	seq
ucuaggcuggCuacuuuacagu		2	1	seq
ucuUggcugguuacuuuacaguu		1	1	seq
ucuaggcugCuuacuuuacaguu		1	1	seq
ucuaggcugguuacuuuacaguu		396	0	seq
ucuaggcAgguuacuuuacaguu		1	1	seq
ucuaggcugguuacuuuacaguG		2	1	seq
Gcuaggcugguuacuuuacaguu		1	1	seq
ucCaggcugguuacuuuacaguu		2	1	seq
ucuGggcugguuacuuuacaguu		1	1	seq
Gazaguu		1	1	seq
ucuaggcugguuacuuuacagCu		2	1	seq
ucuagAcugguuacuuuacaguu		1	1	seq
ucuaggcugguuacuAuacaguu		1	1	seq
ucuaggcugguuacuuuacaguC		35	1	seq
GuaggcugguuaGuuuacaguu		2	1	seq
ucuaggcugguuUcuuuacaguu		2	1	seq
Acuaggcugguuacuuuacaguu		2	1	seq
ucuaggcugguuacuuuacagGu		6	1	seq
ucuaggcugguuacuuuacGguu		3	1	seq
acAaggcugguuacuuuacaguu		1	1	seq
ucuaggcugguuacuuuacagAu		1	1	seq
ucuaggcugguuGcuuuacaguu		1	1	seq
		1	1	seq
ucuaggcugguuacuCuacaguu		2	1	seq
uGuaggcugguuacuuuacaguu		1	1	seq
ucuaggcugguuacuuuacaguA		13	1	seq
ucuaggcugguuacuuuUcaguu		1	1	seq
ucuaggcugguuacuGuacaguu		2	1	seq
ucuaggcCgguuacuuuacaguu		2	1	seq
		3	1	seq
CuaggcugguuacCuuacaguu		3	1	seq
ucuaggcugguuacuuuacaUuu		2	1	seq
ucuaggcugguuacuuuacaguuU		1	1	seq
cuaggcugguuacuuuaca		1	0	seq
cuaggcugguuacuuuacag		1	0	seq
		1	1	seq
cuaggcugguuacuuuacagu		1	0	seq
cuaggcugguuacuuuacaguu		2	0	seq
uaggcugguuacuuuacaguC		1	1	seq
	cauuguaaacuaagcagcuuaaacU	1	1	seq
	uuguaaacuaagcagcuuaaaU	1	1	seq
	uuguaaacuaagcagcuuaaac	2	0	seq
	uuguaaacuaagcagcuuaaacc	2	0	seq
	uuguaaacuaagcagcuuaaaccA	2	1	seq
		-	-	224