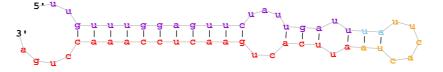
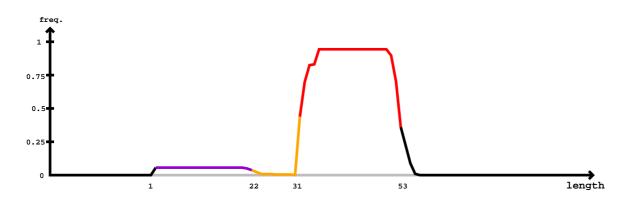
Provisional ID : chromosome\_4\_15243

Score total : 315.3
Score for star read(s) : 3.9
Score for read counts : 309.1
Score for mfe : 0.7
Score for randfold : 1.6
Score for cons. seed : 618
Matura read count : 585



Mature read count : 585 Loop read count : 0 Star read count : 33



5'- aaaguuugcgacaaggaaucuuguuuggaguucuauugauuuauucacuaauucacugaacuccaaaccugauuuccuguuugcuguaggagaccaucaauggccguua

Star	Mature

aaaguuugcgacaaggaaucuuguuuggaguucuauugauuuauucacua <mark>auucacugaacuccaaaccugau</mark> uuccuguuugcuguaggagagacaucaauggccguua	exp	•	
(((((.((((((((((((((((())),.)),.	reads	mm.	sample
uuguuggaguucuauuga	4	0	seq
uuguuggaguucuauGgau	1	1	seq
uuguuggaguucuauugau	5	0	seq
uguuuggaguucuauugaC	2	1	seq
uguuuggaguucuauugauu	15	0	seq
uuguuggaguucuauugauAu	1	1	seq
uguuuggaguucuauugauuu	2	0	seq
uuguuggaguucuauugauuua	1	0	seq
uuguuggaguucuauugauuuaAu	2	1	seq
uguuuggaguucuauugauuuauucacu	2	0	seq
uucacuaauucacugaacuccaaaccugau	1	0	seq
auucacugaacuccaaacc	4	0	seq
auucacugaacuccaaaeU	3	1	seq
auucacugaacuccaaacA	14	1	seq
aCucacugaacuccaaaccu	1	1	seq
auucacugaacuccaaaccu	51	0	seq
auucacugaacuccaaaccC	9	1	seq
auucacCgaacuccaaaccu	1	1	seq
auucacugaacuccaaaccug	53	0	seq
auucacugaacuccaaaccuU	2	1	seq
auucacugaacuccaaaccuC	2	1	seq
auucacCgaacuccaaaccug	3	1	seq
auucacugaacuccaaaccuA	7	1	seq
auCcacugaacuccaaaccug	2	1	seq
Guucacugaacuccaaaccug	3	1	seq
auucacugaacuccaaaccuUa	1	1	seq
auucacugaacuccaaaccugU	7	1	seq
auucacugaacuccaaaccugG	2	1	seq
auucacugaacuccaaaccuga	80	0	seq
auucGcugaacuccaaaccuga	1	1	seq
auucacugaacuccaaaccGga	2	1	seq
auucacGgaacuccaaaccuga	1	1	seq
auucacugaacuccaaaccugC	1	1	seq

Star Mature

Star Matu	re			
aaaguuugcgacaaggaaucuuguuuggaguucuauugauuuauucacuaauucacuga	acuccaaaccugauuuccuguuugcuguaggagagaccaucaauggccguua			
auucacCga	acuccaaaccuga	1	1	seq
auucacuga	acuccaaaccugGu	1	1	seq
auucacuga		2	1	seq
auucacuga		14	0	seq
	acuccaaaccugauA	1	1	seq
auucacuga	acuccaaaccugauAu	1	1	seq
auucacuga	acuccaaaccugauuA	1	1	seq
uucacuga	acuccaaacU	4	1	seq
uucacuga	acuccaaacA	1	1	seq
uucacuga	acuccaaacG	3	1	seq
uucacuga	acuccaaaccu	38	0	seq
uucacuga		2	1	seq
uucacuga		5	1	seq
			1	
uucacuCa		1		seq
uucacAga		1	1	seq
uucacuga	acuccaaaccuU	2	1	seq
	acuccaaaccug	1	1	seq
uucacuga	acuccaaaccuA	8	1	seq
uucacuga	Gcuccaaaccug	1	1	seq
uucacuga	acuccaaaccug	70	0	seq
uAcacuga	acuccaaaccug	1	1	seq
uucacuga		2	1	seq
uucacuga		1	1	seq
uucacugg	•	16	0	seq
		3	0	_
	acuccaaaccugau			seq
	acuccaaaccugauu	1	0	seq
ucacuga		12	0	seq
	acuccaaaccuA	11	1	seq
ucacuga		1	1	seq
ucacuga		2	1	seq
ucacCga	acuccaaaccug	1	1	seq
ucGcuga	acuccaaaccug	1	1	seq
ucaUuga	acuccaaaccug	2	1	seq
ucacuga	acuccaaaccug	35	0	seq
ucacuga	acuccaaaccuga	7	0	seq
ucacuga	acuccaaaccugU	4	1	seq
ucacuga	acuccaaaccugau	1	0	seq
ucacuga	acuccaaaccugauA	2	1	seq
cacuga	acuccaaaccuA	2	1	seq
cacuga		2	0	seq
		3	1	seq
	acuccaaaccugC	9		_
	acuccaaaccuga		0	seq
	acuccaaaccGga	1	1	seq
	acuccaaaccugau	7	0	seq
		1	1	seq
acuga		40	0	seq
acuga	acuccaaacUugauu	1	1	seq
acuga	acCccaaaccugauu	1	1	seq
acuga	acuccaaaccugauA	1	1	seq
acuga	acuccaaaccugauC	2	1	seq
acuga	acuccaaaccugauGu	1	1	seq
acuga	acuccaaaccugauuA	2	1	seq
acuga		1	1	seq
_				_