Provisional ID : chromosome_4_15211

0

33

Star

 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

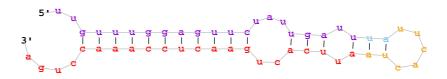
 Score for cons. seed
 :

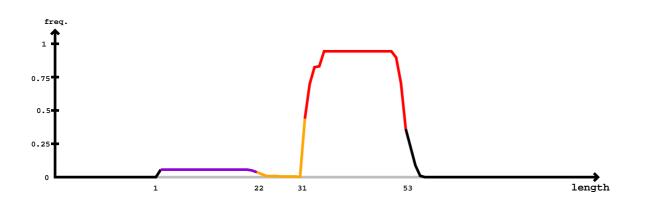
 Total read count
 : 618

 Mature read count
 : 585

Loop read count

Star read count





Mature

.....auucacugaacuccaaaccuUa......

Э.	- aaaguuugcgacaaggaaucuuguuuggaguucuauugauuuauucacua <mark>auucacugaacuccaaaccuga</mark> uuuccuguuugcuguaggagagaccaucaauggccguua	-3'	obs	
	aaaguuugcgacaaggaaucuuguuuggaguucuauugauuua <mark>uucacuaauucacugaacuccaaaccuga</mark> uuuccuguuugcuguaggagagaccaucaauggccguua		exp	
	(((((.(((((((((((((((((())),.)),.	read	ls mm	sample
	uuguuuggaguucuauuga	4	0	seq
	uuguuuggaguucuauugau	5	0	seq
	uuguuuggaguucuauugaC	2	1	seq
	uuguuggaguucuauGgau	1	1	seq
	uuguuggaguucuauugauu	15	0	seq
	uuguuuggaguucuauugauAu	1	1	seq
	uuguuggaguucuauugauuu	2	0	seq
	uuguuggaguucuauugauuua	1	0	seq
	uuguuuggaguucuauugauuuaAu	2	1	seq
	uguuuggaguucuauugauuuauucacu	2	0	seq
		1	0	seq
	auucacugaacuccaaacU	3	1	seq
	auucacugaacuccaaacc	4	0	seq
	auucacugaacuccaaacA	14	1	seq
	auucacugaacuccaaaccC	9	1	seq
	aCucacugaacuccaaaccu	1	1	seq
	auucacCgaacuccaaaccu	1	1	seq
	auucacugaacuccaaaccu	51	0	seq
	auucacugaacuccaaaccuC	2	1	seq
	auucacugaacuccaaaccug	53	0	seq
	auCcacugaacuccaaaccug.	2	1	seq
	auucacugaacuccaaaccuU	2	1	seq
	Guucacugaacuccaaaccug.	3	1	seq
	auucacugaacuccaaaccuA	7	1	seq
	auucacCgaacuccaaaccug	3	1	seq
	auucacugaacuccaaaccugG	2	1	seq
	auucacugaacuccaaaccugC	1	1	seq
	auucGcugaacuccaaaccuga	1	1	seq
	auucacCgaacuccaaaccuga	1	1	seq
	auucacugaacuccaaaccugU	7	1	seq
	auucacugaacuccaaaccuga	80	0	seq
	auucacugaacuccaaaccGga	2	1	seq

Star Mature

Star	Mature			
aaaguuugcgacaaggaaucuuguuuggaguucuauugauuuauucacu	aauucacugaacuccaaaccugauuuccuguuugcuguaggagagaccaucaauggccguua			
	.auucacGgaacuccaaaccuga	1	1	seq
	.auucacugaacuccaaaccugGu	1	1	seq
		2	1	seq
	.auucacugaacuccaaaccugaA			_
	.auucacugaacuccaaaccugau	14	0	seq
•••••	.auucacugaacuccaaaccugauA	1	1	seq
•••••	.auucacugaacuccaaaccugauuA	1	1	seq
•••••	.auucacugaacuccaaaccugauAu	1	1	seq
	uucacugaacuccaaacG	3	1	seq
	uucacugaacuccaaacU	4	1	seq
	uucacugaacuccaaacA	1	1	seq
	uucacugaacuccaaacUu	2	1	seq
	uucacugaacuccaaaccu	38	0	seq
	uucacugaacuccaaaccC	5	1	seq
	-			_
	uucacuCaacuccaaaccug	1	1	seq
•••••	uucacugaGcuccaaaccug	1	1	seq
•••••	uAcacugaacuccaaaccug	1	1	seq
	uucacAgaacuccaaaccug	1	1	seq
	Cucacugaacuccaaaccug	1	1	seq
	uucacugaacuccaaaccuU	2	1	seq
	uucacugaacuccaaaccuA	8	1	seq
	uucacugaacuccaaaccug	70	0	seq
	uucacugaacuccaaaccugU	1	1	seq
	uucacugaacuccaaaccuga	16	0	seq
	uucacugaacuccaaGccuga	2	1	seq
	uucacugaacuccaaaccugau	3	0	seq
	uucacugaacuccaaaccugauu	1	0	seq
	ucacugaacuccaaaccu	12	0	seq
		35	0	_
	ucacugaacuccaaaccug	1	1	seq
	ucacugaUcuccaaaccug	11	1	seq
	ucacugaacuccaaaccuAucGcugaacuccaaaccug	1	1	seq
				seq
	ucacCgaacuccaaaccug	1	1	seq
	ucacugaacuccaaaccuU	2	1	seq
	ucaUugaacuccaaaccug	2	1	seq
•••••	ucacugaacuccaaaccuga	7	0	seq
•••••	ucacugaacuccaaaccugU	4	1	seq
	ucacugaacuccaaaccugau	1	0	seq
	ucacugaacuccaaaccugauA	2	1	seq
	cacugaacuccaaaccug	2	0	seq
	cacugaacuccaaaccuA	2	1	seq
	acugaacuccaaaccGga	1	1	seq
	acugaacuccaaaccugC	3	1	seq
	acugaacuccaaaccuga	9	0	seq
	acugaacuccaaaccugau	7	0	seq
	Ccugaacuccaaaccugau	1	1	seq
	acugaacuccaaaccugauC	2	1	seq
	acugaacuccaaaccugauA	1	1	seq
	acugaacuccaaaccugauu	40	0	seq
		1	1	_
	acugaacCccaaaccugauuacugaacuccaaacUugauu	1	1	seq
		1	1	seq
	acugaacuccaaaccugauuC			seq
	acugaacuccaaaccugauGu	1	1	seq
	acugaacuccaaaccugauuA	2	1	seq