Provisional ID : chromosome_14_73226

3286

0

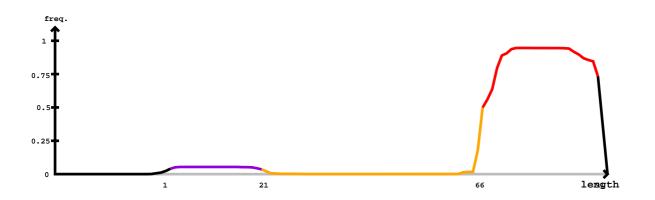
109

Score total : 0.7
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 0.4
Score for randfold : 1.6
Score for cons. seed :
Total read count : 3395

Mature read count Loop read count

Star read count





Star Mature

5'- aaauuauuuucuuaaaaaagugaggucaaguggagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccucg
aaauuauuuucuuaaaaaaagugaggucaaguggagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccucg

und	CAL	•	
((((((((((((((((((((((((((((((((reads	mm	sample
aaUugaggucaagugagug	1	1	seq
aaUugaggucaagugaguggaag	2	1	seq
aCugaggucaagugagug	2	1	seq
agugaggucaagugagug	2	0	seq
Ugugaggucaagugaguga	3	1	seq
aUugaggucaagugaguga	1	1	seq
Ugugaggucaagugaggaag.	1	1	seq
aUugaggucaagugaguggaag.	2	1	seq
aUugaggucaagugaguggaagag.	1	1	seq
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aUugaggucaagugaggaagagg.	1	1	seq
agugaggucaagugagggaaug	2	0	seq
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	2	1	seq
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	6	1	seq
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	2	1	seq
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	1	1	seq
ugaggucaagugGguggaa	1	1	seq
ugaggucaagugaguggaa	12	0	seq
uAaggucaagugagugaa	3	1	seq
ugagguUaagugagugaa	1	1	seq
ugaggucaagugaguggaag.	9	0	seq
	1	1	seq
Cgaggucaagugaggaag.	1	1	seq
uAaggucaaguggaag	1	1	seq
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Star Mature			
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	19	1	seq
	16	1	seq
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.gaggucaagugaguggaaga.	12	0	seq
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Gggucaagugaguggaaga.	2	1	seq
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	2	1	seq
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Ugucaaguggaguggaagag	1	1	seq
ggucaagugaguggaagaggg	3	0	seq
Ugucaagugaguggaagaggg	2	1	seq
	1	1	seq
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aacGgauaacccucaaccguccggaccu	1	1	seq
aGcugauaacccucaaccguccggaccu.	1	1	seq
	5	1	seq
	12	1	_
	1	1	seq
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aaAugauaacccucaaccguccggaccuc.	1	1	seq
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	2	1	seq
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cAgauaacccucaaccguccggaccuc.	2	1	seq
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	1 1 9 1	0 1 1	sec
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	1 9 1 1 2	0 1 1 1	sec sec
	1 1 9 1 1 2 1	0 1 1 1 1	seq seq seq
	1 9 1 1 2 1 1	0 1 1 1 1 1	seq seq seq seq
	1 1 9 1 1 2 1 1 9	0 1 1 1 1 1 0	seq seq seq seq seq
	1 1 9 1 1 2 1 1 9 4 67 2	0 1 1 1 1 1 0 1 1 1 1 1 1 1 1 1 1 1 1 1	sec sec sec sec sec sec
gauaacccucaCccguccgga	1 1 9 1 1 2 1 1 9 4 67 2 1	0 1 1 1 1 1 0 1 1 1 1 1 1 1 1 1 1 1 1 1	sec sec sec sec sec sec sec
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	iture		
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	aacccucaaccgucGggaccuc. 1	1	seq
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	aacccucaaccguccggaccAc. 2		seq
	aacccucaaccguccgAaccuc. 1		seq
gaua			seq
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Talia A			504
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gaua	aacccucaaccguccggaccucg 3 aacccucaaccguccg	0 1 0 1 0	pee pee pee pee pee pee pee pee pee pee
	aacccucaaccguccggaccucg 3 aacccucaaccguccg	0 1 0 1 0 1 1	pea pea pea pea pea pea pea pea pea pea
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gaua Uua aua aua aua Cua aua aua	aacccucaaccguccggaccucg aacccucaaccguccg	0 1 1 0 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	seq
gaua Uua aua aua Cua aua aua aua	aacccucaaccguccggaccucg aacccucaaccguccg	0 1 1 0 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	seq
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gaua Uua aua aua aua Cua aua aua	aacccucaaccguccggaccucg aacccucaaccguccg	0 1 1 0 0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	seq
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Star Mature			
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		1	seq
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		1	seq
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		1	seq
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		1	seq seq
		1	seq
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		1	seq
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.aacceGeaaccguceggaccuc		1	seq
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aacccucaacGguccggaccuc		1	seq
		1	seq
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Star	Mature			
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	acccucaaccguccggac	15	0	seq
	acccucaaccguccggacc	3	0	seq
	acccucaaGcguccggacc	1	1	seq
		1	1	seq
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		1	1	seq
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	acccucaaccguccggaccu	21	0	seq
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		2	1	
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		1	1	seq
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		1	1	seq
		21	0	seq
•••••••••••••••••••••••••••••••••••••••	ccucaaccAuccggaccu	2	1	seq
••••••	ccucaaccguccggaccCc.	1	1	seq
	Acucaaccguccggaccuc.	2	1	seq
	ccucaaccCuccggaccuc.	1	1	seq
	Ucucaaccguccggaccuc.	9	1	seq
	Gcucaaccguccggaccuc.	2	1	seq
	ccucaaccAuccggaccuc.	2	1	seq
	ccucaaccguccgAaccuc.	2	1	seq
	cUucaaccguccggaccuc.	1	1	seq
	ccucaaccUuccggaccuc.	1	1	seq
	ccucaaccguccggaccuc.	65	0	seq
	ccucaaccguccggGccuc.	1	1	seq
	ccucaaccguccggaccucg	1	0	seq
	Uucaaccguccggaccuc.	1	1	seq
	cucaaccguccggaccuc.	27	0	seq
		1	1	seq
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