Provisional ID	: chr2_3685
Score total	: 12151.4
Score for star read(s)	: 3.9
Score for read counts	: 12146.2
Score for mfe	: -0.2
Score for randfold	: 1.6
Score for cons. seed	:
Total read count	: 23836
Mature read count	: 23807
Loop read count	: 0
Star read count	: 29

Mature   **Star**

```
5'- ccucucaucagaagcaucgaucugaagauuccagucaccaguaugugaccggucacugcaaucuacaguucgauuuucucgacaucgcgcaguuuauguuauuuauucuau  -3'    obs
    ccucucaucagaagcaucgaucugaagauuccagucaccaguaugugaccggucacugcaaucuacaguucgauuuucucgacaucgcgcaguuuauguuauuuauucuau        exp
    .......(((((((.(((((.(((.(((((.((((.(((.((.....)).))).)))).))))).))).))))).))))).))......(((.....)))...........      reads   mm   sample
```

read	reads	mm	sample
.................ucugaagauuccagucaA..	1	1	seq
.................ucugaagauuccaAucac..	14	1	seq
.................ucugaagaAuccagucac..	1	1	seq
.................ucugaagauuccagucac..	56	0	seq
.................ucugaagauuccagucacc...	117	0	seq
.................ucugaagauuccaAucacc...	3	1	seq
.................ucugaagUuuccagucacc...	1	1	seq
.................ucugaagUuuccagucacca..	1	1	seq
.................ucugaagauuccagucacca..	250	0	seq
.................ucugaagaAuccagucacca...	4	1	seq
.................uAugaagauuccagucacca...	1	1	seq
.................ucugaagauucGagucacca...	1	1	seq
.................ucGgaagauuccagucaccag..	3	1	seq
.................ucugaagauuccaUucaccag..	2	1	seq
.................ucAgaagauuccagucaccag..	2	1	seq
.................ucugaagaAuccagucaccag..	10	1	seq
.................ucugaagauuccCgucaccag..	1	1	seq
.................ucugaagauuccUgucaccag..	1	1	seq
.................ucugaagauucAagucaccag..	1	1	seq
.................ucugaagUuuccagucaccag..	11	1	seq
.................Acugaagauuccagucaccag..	1	1	seq
.................uAugaagauuccagucaccag..	1	1	seq
.................ucugaagauuccagucaccag..	1128	0	seq
.................ucuUaagauuccagucaccagu...	17	1	seq
.................uUugaagauuccagucaccagu...	4	1	seq
.................ucugaGgauuccagucaccagu...	3	1	seq
.................ucugaagauuccagucacAagu...	1	1	seq
.................ucugaagauuccagucacAagu...	1	1	seq
.................ucugaagauuccagucGagu...	1	1	seq
.................ucugaagauuccagCcaccagu...	1	1	seq
.................ucAgaagauuccagucaccagu...	17	1	seq
.................ucugaagauuccagucUccagu...	11	1	seq
.................ucugaagauuccagucaccagu...	16383	0	seq
.................ucugaagaCuccagucaccagu...	4	1	seq

cccucaucagaagcaucgaucugaagauuccagucaccaguauguggaccggucacugcaaucuacaguucgauuuuucucgacaucgcgcaguuuauguuauuuauucuau

Sequence			
....................ucugaagauuccCgucaccagu..	15	1	seq
....................ucugaagaAuccagucaccagu..	113	1	seq
....................uAugaagauuccagucaccagu..	6	1	seq
....................ucugaagGuuccagucaccagu..	1	1	seq
....................ucugaagauuccagucaAcagu..	4	1	seq
....................ucugaagUuuccagucaccagu..	126	1	seq
....................ucugaagauuccaUucaccagu..	22	1	seq
....................ucugaagauCccagucaccagu..	4	1	seq
....................ucugaagauuUcagucaccagu..	1	1	seq
....................ucugaagauuccagucaccagG..	25	1	seq
....................ucugaCgauuccagucaccagu..	1	1	seq
....................ucugGagauuccagucaccagu..	1	1	seq
....................ucugaagauuccagAcaccagu..	4	1	seq
....................ucugaagauuccagucaccaUu..	1	1	seq
....................ucugaagauuccagucaccagA..	10	1	seq
....................ucugaagauucAagucaccagu..	4	1	seq
....................ucugaagauuccagcacUagu..	23	1	seq
....................ucugaagauuccUgucaccagu..	1	1	seq
....................ucugaaAauuccagucaccagu..	1	1	seq
....................Acugaagauuccagucaccagu..	17	1	seq
....................ucugaagauuccagucaccagC..	1	1	seq
....................Ccugaagauuccagucaccagu..	2	1	seq
....................ucugaagauuccagucaccGgu..	960	1	seq
....................ucugaagauAccagucaccagu..	5	1	seq
....................ucugaagauuccagucaccUgu..	8	1	seq
....................ucGgaagauuccagucaccagu..	18	1	seq
....................ucugaagauccGgucaccagu..	5	1	seq
....................ucugaagauuccaguUaccagu..	25	1	seq
....................ucAgaagauuccagucaccagua..	1	1	seq
....................ucugaagUuuccagucaccagua..	2	1	seq
....................ucugaagauuccagucaccGgua..	6	1	seq
....................ucugaagauuccagucaccagua..	109	0	seq
....................ucugaagauuccagucaccagCa..	1	1	seq
....................ucugaagauuccaUucaccagua..	1	1	seq
....................ucugaagauuccagucaccaguU..	2644	1	seq
....................ucugaagauuccagucaccaguC..	4	1	seq
....................ucugaagauuccagucaccaguUu...	1545	1	seq
....................ucugaagauuccagucaccaguCu...	8	1	seq
....................ucugaagauuccagucaccaguaA...	1	1	seq
....................ucugaagauuccagucaccaguau...	4	0	seq
....................ucugaagauuccagucaccaguauU..	15	1	seq
....................ucugaagauuccagucaccaguaug..	7	0	seq
....................ucugaagUuuccagucaccaguaug..	1	1	seq
....................ucugaagauuccagucaccaguaugugaA..	1	1	seq
....................ucugaagauuccagucaccaguaugugaU..	3	1	seq
....................ucugaagauuccagucaccaguaugugacc..	2	0	seq
....................ucugaagauuccagucaccaguaugugacU..	1	1	seq
....................gaagauuccagucaccag..	1	0	seq
....ccggucacugcaaucuacaguuc.............................	25	0	seq
....cggucacugcaaucuacaguuc..............................	3	0	seq
....cggucacugcaaucuacaguucU.............................	1	1	seq