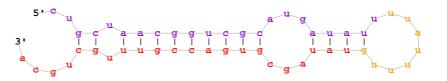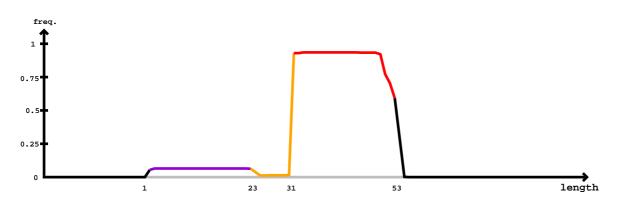Provisional ID   : chromosome_5_137408
Score total   :   287.4
Score for star read(s) :   3.9
Score for read counts :   281
Score for mfe   :   0.8
Score for randfold   :   1.6
Score for cons. seed   :
Total read count   :   563
Mature read count   :   526
Loop read count   :   0
Star read count   :   37

<pre>
                    Star   Mature

5'- acgacccaaugggguccuuuccugcuaacggucgcaugauauuuuauuuuguauagcgugaccguuugcugcaaagugccuauuaacggcuaugaaugaacguaacaaacgu   -3'   obs
    acgacccaaugggguccuuuccugcuaacggucgcaugauauuuuauuuuguauagcgugaccguuugcugcaaagugccuauuaacggcuaugaaugaacguaacaaacgu   exp
    ..(((((...)))))((((.(.((.(((((((((...((((........)))).(((.(((((((((...))))))))).))).).(((((((((...)))).........((((.....)))) reads   mm   sample
    ....cugcuaacggucUcaugauauu..   1   1   s07
    ....cugcuaacggucgcaugauauC..   2   1   s07
    ....ugcuaacggucgcaugauauu..   1   0   s07
    ....uuauuuuguauUgcgugaccguuugcugca..   2   1   s07
    ....guauUgcgugaccguuugcugc...   1   1   s07
    ....uauUgcgugaccguuugc...   2   1   s07
    ....uauUgcgugaccguuugcu..   7   1   s07
    ....uauUgcgugaccguuugcug...   5   1   s07
    ....uauUgcgugaccguuugcugc..   2   1   s07
    ....uauUgcgugaccguuugcugca...   5   1   s07

    ....cugcuaacggucgcaugauauu..   5   0   s12
    ....cugcuaacggucgcaugauauC..   1   1   s12
    ....ugcuaacggucgcaugauau...   1   0   s12
    ....uuauuuuguauUgcgugaccguuugcugca..   5   1   s12
    ....uauUgcgugaccguuugcu..   4   1   s12
    ....uauUgcgugaccguuugcug...   4   1   s12
    ....uauUgcgugaccguuugcugc..   3   1   s12
    ....uauUgcgugaccguuugcugca... 13   1   s12

    ....uauUgcgugaccguuugcu..   3   1   s09
    ....uauUgcgugaccguuugcug...   3   1   s09
    ....uauUgcgugaccguuugcugc..   6   1   s09
    ....uauUgcgugaccguuugcugca... 14   1   s09

    ....uauCgcgugaccguuugcugca...   1   1   s14

    ....uauagcgugaccguuugcugca...   1   0   s08

    ....uauagcgugaccguuugcug...   2   0   s10
    ....uauagcgugaccguuugcugca...   1   0   s10

    ....uauUgcgugaccguuugcug...   2   1   s17
</pre>

Star   **Mature**

acgacccaaugggguccuuuccugcuaacggucgcaugauauuuuauuuuguauagcgugaccguuugcugcaaagugccuauuaacggcuaugaaugaacguaacaaacgu

Sequence			
....uauUgcgugaccguuugcugc...	1	1	s17
....uauUgcgugaccguuugcugca..	1	1	s17
..................cugcuaacggucgcaugauaCu..	1	1	s03
..................cugcuaacggucgcaugauauu..	3	0	s03
....uauUgcgugaccguuugcu...	2	1	s03
....uauUgcgugaccguuugcug..	3	1	s03
....uauUgcgugaccguuugcugc...	1	1	s03
....uauUgcgugaccguuugcugca..	22	1	s03
....uUgcgugaccguuugcugc...	2	1	s03
....uauagcgugaccguuugcugU..	1	1	s06
....uauCgcgugaccguuugcugc..	1	1	s06
....uauCgcgugaccguuugcu...	1	1	s11
..................cugcuaacggucgcaugauauu..	1	0	s02
..................cugcuaacggucgcaugauauG..	17	1	s02
...................ugcuaacggucgcaugauauGu...	4	1	s02
....uauCgcgugaccguuugc..	5	1	s02
....uauCgcgugaccguuugcu...	68	1	s02
....uauCgcgugaccguuugcug..	20	1	s02
....uauCgcgugaccguuugcugc...	46	1	s02
....uauCgcgugaccguuugcugca..	270	1	s02
....uauCgcgugaccguuugcugcaa.......................................	1	1	s02
....uauCgcgugaccguuugcugcaaa......................................	1	1	s02
....uCgcgugaccguuugcugca..	1	1	s02
....uuuuguauagcgugaccAu..	1	1	s05