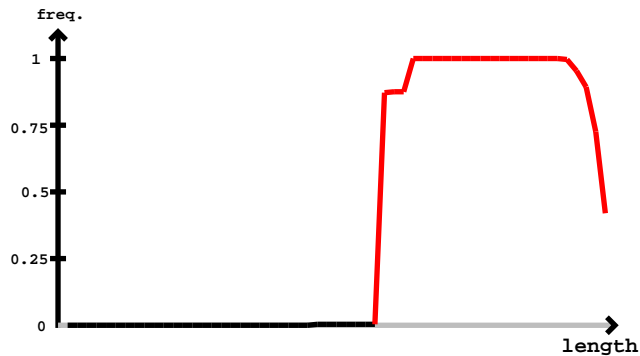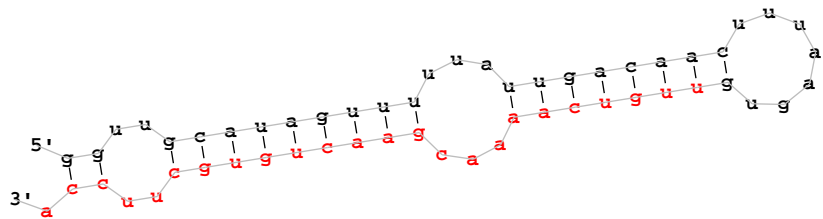miRBase precursor   : chromosome_2_4017
Total read count   : 281
chromosome_2_4017 read count : 280
remaining reads   : 1

chromosome_2_4017

```
5'- gguugcauaguuuuuauugacaacuuuaaguguugucaaaacgaacugugcuucca  -3'     exp
    ((..(((((((((...(((((((((.......)))))))))...)))))))))..)).     reads    mm      sample
    .......................uuUaguguugucaaaacgaacugugcuucc.         1        1       seq
    .......................uugucaaaacgaacugugc.....               1        0       seq
    .......................uugucaaaaUgaacugugcu....               4        1       seq
    .......................uugucaaaacgaacugugcu....               8        0       seq
    .......................uugucaaaacgaacugugcuC...               2        1       seq
    .......................uugucaaaacgaacugugcuu...               10       0       seq
    .......................uugucaaaaUgaacugugcuu...               4        1       seq
    .......................uugucaaaacgaacugugcuuA..               9        1       seq
    .......................uugucaaaacgaacugugcuuc..               26       0       seq
    .......................uugucaaaaUgaacugugcuuc..               7        1       seq
    .......................uugucaaaacgaacugugcuuU..               3        1       seq
    .......................uugucaaaacgaacugugUuuc..               1        1       seq
    .......................Cugucaaaacgaacugugcuucc.               1        1       seq
    .......................uugucaaaacgaacugugUuucc.               1        1       seq
    .......................uGgucaaaacgaacugugcuucc.               1        1       seq
    .......................uugucaaaacgaacugugcuucA.               13       1       seq
    .......................uugucaaaacgaacGgugcuucc.               1        1       seq
    .......................uugGcaaaacgaacugugcuucc.               1        1       seq
    .......................uugucaaaacgaacugugcuucc.               28       0       seq
    .......................uugucaaaacgaacugugcuucU.               20       1       seq
    .......................uugucaaaaUgaacugugcuucc.               17       1       seq
    .......................uugucaaaacgaacugugcGucc.               1        1       seq
    .......................uugucaaaacgaacugugcuucca              39       0       seq
    .......................uugucaaaaUgaacugugcCucca              2        1       seq
    .......................uugucaaaaUgaacugugcuucca              29       1       seq
    .......................uuguUaaaacgaacugugcuucca              2        1       seq
    .......................uugucaaaacgaacugugcuuccC              3        1       seq
    .......................uugucaaGacgaacugugcuucca              1        1       seq
    .......................uuguAaaaacgaacugugcuucca              1        1       seq
    .......................uugucaaaacgaacugugcuucAa              2        1       seq
    .......................Gugucaaaacgaacugugcuucca              1        1       seq
    .......................uugucaaaacgaacugugcuuccU              5        1       seq
    .......................ugucaaaacgaacugugcuC...               1        1       seq
    .......................ucaaaacgaacugugcuuc..                 1        0       seq
    .......................ucaaaacgaacugugcuucc.                 1        0       seq
```

gguugcauaguuuuuauugacaacuuuaaguguugucaaaacgaacugugcuucca

```
...................................ucaaaacgaacugugcuuccU    3    1    seq
...................................ucaaaacgaacugugcuuccC    1    1    seq
...................................Ccaaaacgaacugugcuucca    1    1    seq
...................................ucaaaacgaacCgugcuucca    1    1    seq
...................................ucaaaacgaacugugcuucca   21    0    seq
...................................ucaaaaUgaacugugcuucca    6    1    seq
```