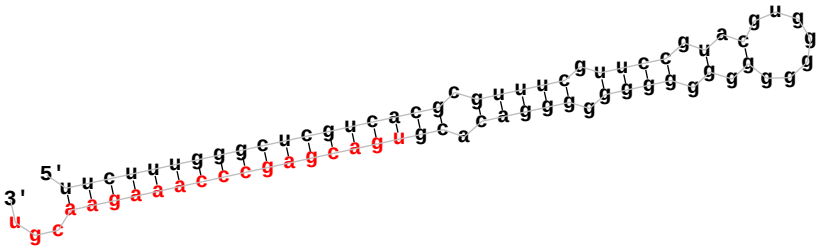
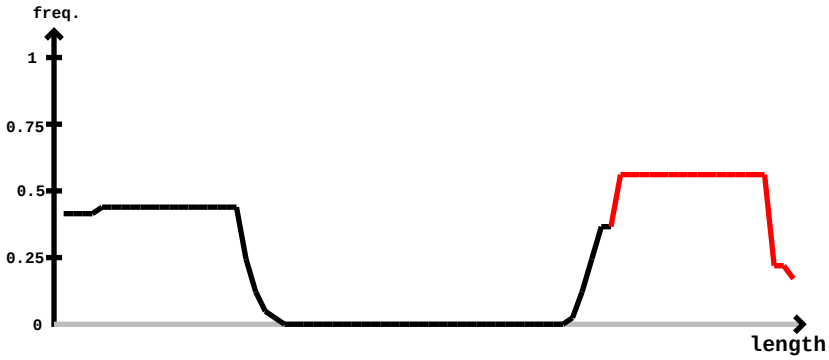
miRBase precursor : chromosome_5_20362
Total read count : 41
chromosome_5_20362 read count : 13
remaining reads : 28





chromosome_5_20362

```
5'- uucuuugggcucgucacgcguuucguuccguacgugggggggggggggggggacacgugacgagcccaaagaacgu  -3'    exp
    ((((((((((((((((((.(((((.((((.(.(.......).).)))).)))))).))))))))))))))))))...   reads      mm         sample
    uucuuugggcucgucacA.................................................................  4          1          seq
    uucuuugggcucgucacg.................................................................  3          0          seq
    uucuuugggcucUucacg.................................................................  1          1          seq
    uucuuugggcucgucacgc................................................................  4          0          seq
    uucuuugggcucgucacgU................................................................  1          1          seq
    uucuuugggcucgucacgcg...............................................................  3          0          seq
    uucuuugggcucgucacgcgu..............................................................  1          0          seq
    ...uuugggcucgucacgcguA.............................................................  1          1          seq
    ...................................................acGcgugacgagcccaaagaa...  1          1          seq
    ...................................................cGcgugacgagcccaaagaa...  3          1          seq
    ...................................................cGcgugacgagcccaaagaacg.  1          1          seq
    ...................................................acgugacgagcccaaagaa...  1          0          seq
    ...................................................Gcgugacgagcccaaagaa...  4          1          seq
    ...................................................cgugacgagcccaaagaa...  3          0          seq
    ...................................................Ugugacgagcccaaagaa...  2          1          seq
    ...................................................ugacgagcccaaagaacg.  1          0          seq
    ...................................................ugaUgagcccaaagaacgu  1          1          seq
    ...................................................ugacgagcccaaagaacgu  6          0          seq
```