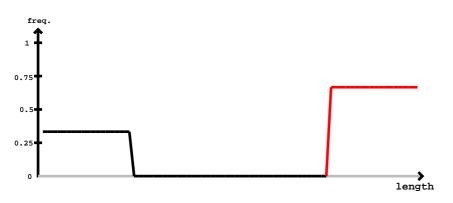miRBase precursor : chromosome_2_4261
Total read count : 3
chromosome_2_4261 read count : 2
remaining reads : 1

chromosome_2_4261

```
5'- cuggccaaacgaucgcaacauuuggacgcaacauaugccaacauugucgcguucagauguugcgaucguuuggccacc  -3'    exp
     .(((((((((((((((((((((((((((((.(((.(((....))))))).)))))))))))))))))))))))))))..    reads      mm       sample
     cCggccaaacgaucgcaa.......................................................    1          1        seq
     .....................................................uugcgaucguuuggccacc    2          0        seq
```