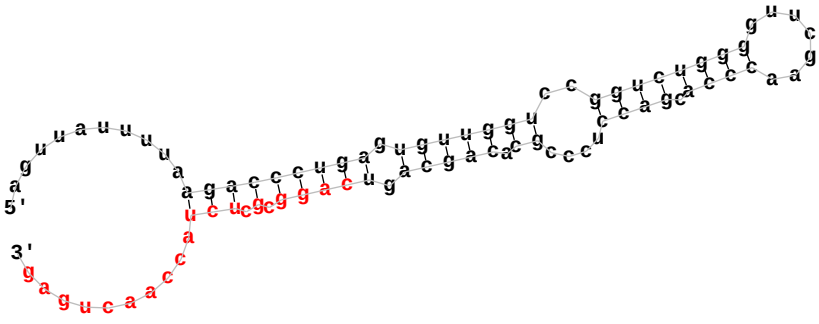
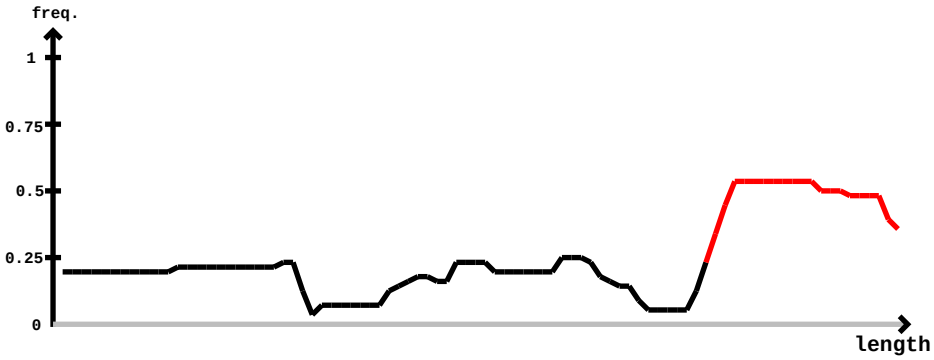
miRBase precursor : chromosome_12_58177
Total read count : 56
chromosome_12_58177 read count : 27
remaining reads : 29





chromosome_12_58177

```
5'- aguuauuuuaagacccugaguguugguccggucuggggguucgaacccacgaccucccgcacagcagucaggcgcucuaccaacugag  -3'   exp
    ..........(((((((((.(((((((..((((((((.......)))).))))....)).))))).))))).).))).........   reads     mm        sample
    aguuauuuuaagacccugaguguu.................................................................   5         0         seq
    aguuauuuuaagGcccugaguguu.................................................................   1         1         seq
    aguuauuuuaagacccugaguguug................................................................   3         0         seq
    aguuaCuuuaagacccugaguguug................................................................   1         1         seq
    aguuauuuuGagacccugaguguug................................................................   1         1         seq
    ...........gacccugaguguugguccggucugggA...................................................   1         1         seq
    ......................uugguccggucuggggguuc...............................................   1         0         seq
    ........................uccggucuggggguucgaU.............................................   2         1         seq
    ................................ugggAuucgaacccacgaccuc..................................   3         1         seq
    ................................gggguucgaacccGcgaccucc..................................   1         1         seq
    ................................ggguucgaacccacgaccuccA..................................   1         1         seq
    ................................ggCucgaacccacgaccu......................................   1         1         seq
    ......................................cgaacccacgaccuUccgc................................   2         1         seq
    ......................................cgaacccacgaccucUcgca...............................   2         1         seq
    ......................................gaacccacgaccucGcgc.................................   1         1         seq
    ..................................................ccucccgcacagcagucCggcgcucua.........   2         1         seq
    ..................................................ccucccgcacagcagucCggcgcucuacca......   1         1         seq
    ..................................................................gGcaggcgcucuaccaacug..   3         1         seq
    ..................................................................gGcaggcgcucuaccaacuga.   1         1         seq
    ..................................................................Gcaggcgcucuaccaacug..   1         1         seq
    ..................................................................Acaggcgcucuaccaacug..   1         1         seq
    ..................................................................Gcaggcgcucuaccaacugag   4         1         seq
    ..................................................................caggcgcucuaccaacugag   6         0         seq
    ..................................................................aCgcgcucuaccaacuga.   1         1         seq
    ..................................................................aggcgcucuaccaacugag   3         0         seq
    ..................................................................Uggcgcucuaccaacugag   2         1         seq
    ..................................................................ggcgcucuaccGacugag   1         1         seq
    ..................................................................gCcgcucuaccaacugag   1         1         seq
    ..................................................................gAcgcucuaccaacugag   2         1         seq
    ..................................................................ggcgcucuCccaacugag   1         1         seq
```