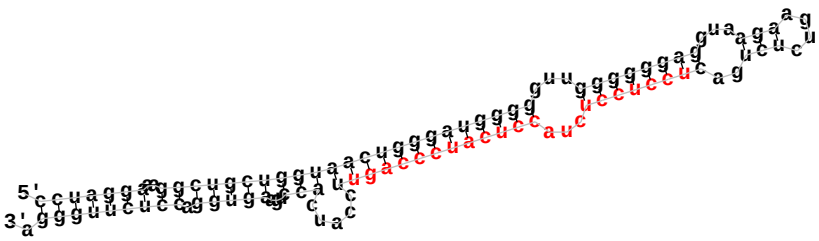
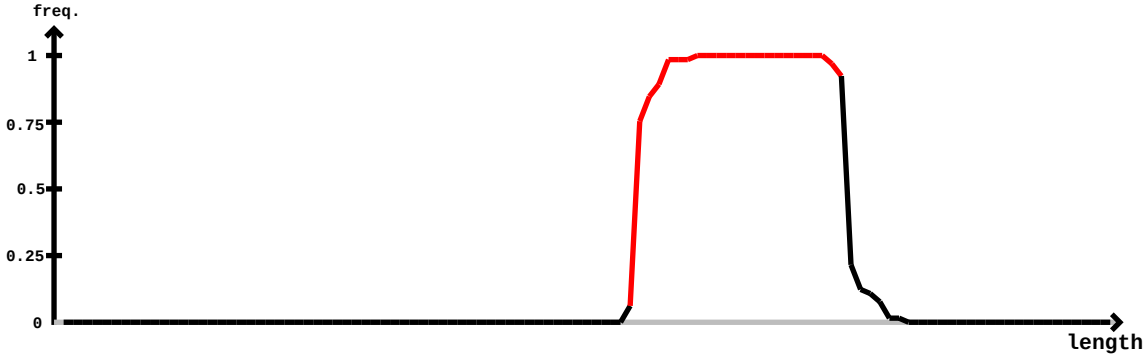
miRBase precursor	: hsa-mir-5193
Total read count	: 65
hsa-miR-5193 read count	: 64
remaining reads	: 1





hsa-miR-5193

```
5'- ccuaggaaaggcugcugguaacugggaugggggguuggggggagguaagaagucucugacuccuccucuaccucaucccaguuccaucaccugaaguggaccucuuggga  -3'    exp
    (((((((..((((((((((((((((((((((...((((((((...(((....)))..))))))))...))))))))))))).....)))...))))).))))))))).   reads     mm        sample
    ..................................................................uccuccucuaccucauccca.............................   2         0         Z3_
    ..................................................................uccuccucuaccucaucccagu...........................   2         0         Z3_
    ..................................................................ccuccucuaccucaucccagu............................   2         0         Z3_

    ..................................................................cuccuccucuaccucaucccagu...........................   4         0         GDM
    ..................................................................uccuccucuaccucaucccagA............................   7         1         GDM
    ..................................................................uccuccucuaccucaucccagu...........................   13        0         GDM
    ..................................................................uccuccucuaccucUucccagu...........................   1         1         GDM
    ..................................................................uccuccucuaccucaucccagu...........................   13        0         GDM
    ..................................................................uccuccucuaccucaucccagu...........................   13        0         GDM
    ..................................................................uccuccucuaccucaucccagA............................   7         1         GDM
    ..................................................................uccuccucuaccucaucccaguu..........................   1         0         GDM
    ..................................................................ccuccucuaccucaucccagu............................   1         0         GDM
    ..................................................................ccuccucuaccucaucccaguu..........................   1         0         GDM
    ...................................................................uccucuaccucaucccaguCcc.........................   1         1         GDM
    ...................................................................uccucuaccucaucccaguucc.........................   1         0         GDM
    ...................................................................uccucuaccucaucccaguucca........................   4         0         GDM
    ....................................................................ucuaccucaucccaguuccauA.......................   1         1         GDM

    ..................................................................uccuccucuaccucaucccagu...........................   5         0         Z2_
    ..................................................................ccuccucuaccucaucccaguu..........................   2         0         Z2_

    ..................................................................uccuccucuaccucaucccag............................   2         0         Z1_
    ..................................................................uccuccucuaccucaucccaA............................   1         1         Z1_
    ..................................................................uccuccucuaccucaucccagC...........................   1         1         Z1_
    ..................................................................uccuccucuaccucaucccagu...........................   10        0         Z1_
    ...................................................................cuccucuaccucaucccaguu..........................   2         0         Z1_
    ...................................................................cuccucuaccucaucccaguuc.........................   1         0         Z1_
```