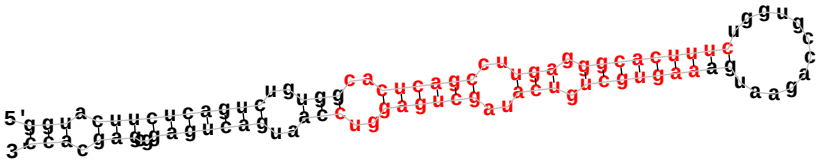
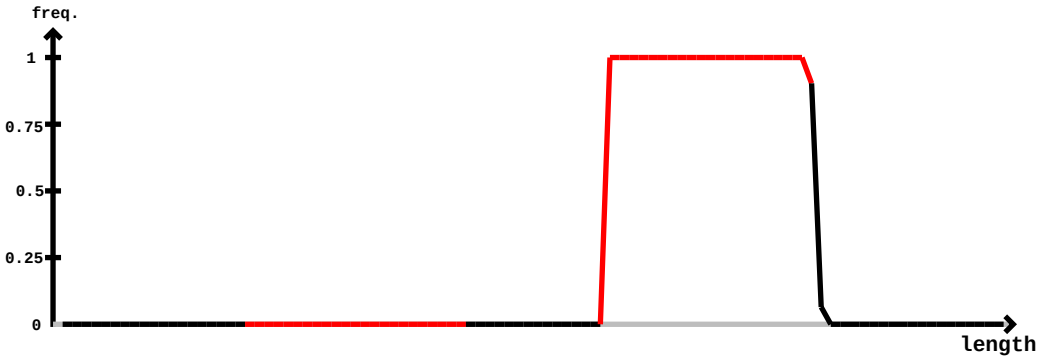
miRBase precursor	: hsa-mir-512-2
Total read count	: 31
hsa-miR-512-5p read count	: 0
hsa-miR-512-3p read count	: 31
remaining reads	: 0





hsa-miR-512-5p

hsa-miR-512-3p

```
5'- gguacuucucagucugugg cacucagccuugagggcacuuuc uggugccagaauga aagugcugucauagcugagguc caaugacugaggcgagcacc  -3'   exp
    (((.((((((((((..(((..((((((..(((.(((((((((.............)))))))))..)))..))))))..)))..)))))))..))).)))   reads     mm        sample
    ..........................................................aagugcugucauagcugagguU.....................   1         1         Z1_
    ..........................................................aagugcugucauagcugagguUc....................   1         1         Z1_

    ..........................................................aagugcugucauagcugaggC......................   2         1         Z2_
    ..........................................................aagugcugucauagcugaggu......................   1         0         Z2_
    ..........................................................aagugcugucauaAcugagguc.....................   1         1         Z2_
    ..........................................................aagugcugucauagcugagguc.....................   4         0         Z2_

    ..........................................................aagugcugucauagcugagguc.....................   11        0         GDM
    ..........................................................aagugcugucauagcugGgguc.....................   1         1         GDM
    ..........................................................aagugcugucauagcugagguc.....................   11        0         GDM
    ..........................................................aagugcugucauagcugaggucA....................   1         1         GDM

    ..........................................................aagugcugucauagcugagguc.....................   3         0         Z3_
    ..........................................................aagugcugucauagcugagguU.....................   3         1         Z3_
    ..........................................................aagugcugucauagcugagguA.....................   1         1         Z3_
    ..........................................................aagugcuCucauagcugagguc.....................   1         1         Z3_
```