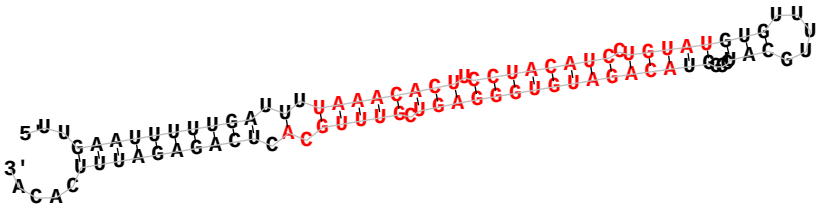
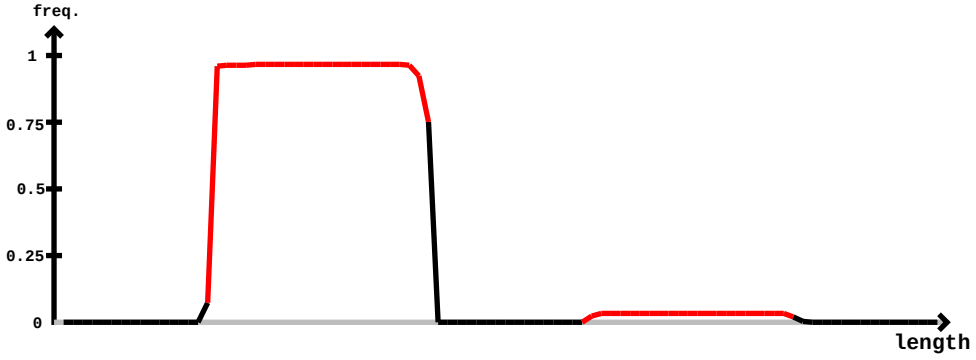
miRBase precursor : dme-mir-975_MI0005832
Total read count : 301
dme-miR-975-3p_MIMAT0020872 read count
dme-miR-975-5p_MIMAT0005491 read count
remaining reads : 0



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                                   dme-miR-975-5p_MIMAT0005491
                                                                   dme-miR-975-3p_MIMAT0020872
5'- uugaauuuuugauuuuaaacacuuccuacauccuguauguguuuugcauccgguacagauguggagucguuugcacucagagauuucaca  -3'    exp
    ..((((((((((.(.(((((((.(((((((.(((((((.....)))...)))))))))))))))).)))))).).))))))))))....   reads     mm        sample
    .............uuaaacacuuccuacauccug.......................................................   1         0         seq
    .............uuaaacacuuccuacauccugu......................................................   4         0         seq
    .............Auaaacacuuccuacauccugu......................................................   1         1         seq
    .............uuaaacacuuccuacauccugua.....................................................   14        0         seq
    .............uuaaacacuuccuacauccuguau....................................................   2         0         seq
    ..............uaaacacuuccuacauccugu......................................................   7         0         seq
    ..............Aaaacacuuccuacauccugua.....................................................   3         1         seq
    ..............uaaacacuuccuacauccugua.....................................................   33        0         seq
    ..............uaaacacuuccAacauccugua.....................................................   1         1         seq
    ..............uaaacacuuccAacauccuguau....................................................   1         1         seq
    ..............uaaacacuuccuacGuccuguau....................................................   1         1         seq
    ..............uaaacacAuccuacauccuguau....................................................   1         1         seq
    ..............uaaacacuuccuacaAccuguau....................................................   2         1         seq
    ..............uaaacacuuccuacauccuAuau....................................................   1         1         seq
    ..............uaaacacuuccuacauccuguau....................................................   210       0         seq
    ..............uaaacacuuccuacauccuguaG....................................................   1         1         seq
    ..............uaaacacuuccuacauAcuguau....................................................   1         1         seq
    ..............uaaacacuuccuGcauccuguau....................................................   1         1         seq
    ..............Gaaacacuuccuacauccuguau....................................................   2         1         seq
    ..............uaaGcacuuccuacauccuguau....................................................   1         1         seq
    ..............uaaacacuuUcuacauccuguau....................................................   1         1         seq
    ...............Uaacacuuccuacauccugua.....................................................   1         1         seq
    ................cacuuccuacauccuguau......................................................   1         0         seq
    ...................................................acagauguggagucguuugc................   3         0         seq
    ...................................................acagaugGggagucguuugc................   1         1         seq
    ...................................................acagauguggagucguuugca...............   3         0         seq
    ..................................................cagauguggagucguuugca................   2         0         seq
    ..................................................cagauguggagucguuugcaA...............   1         1         seq
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