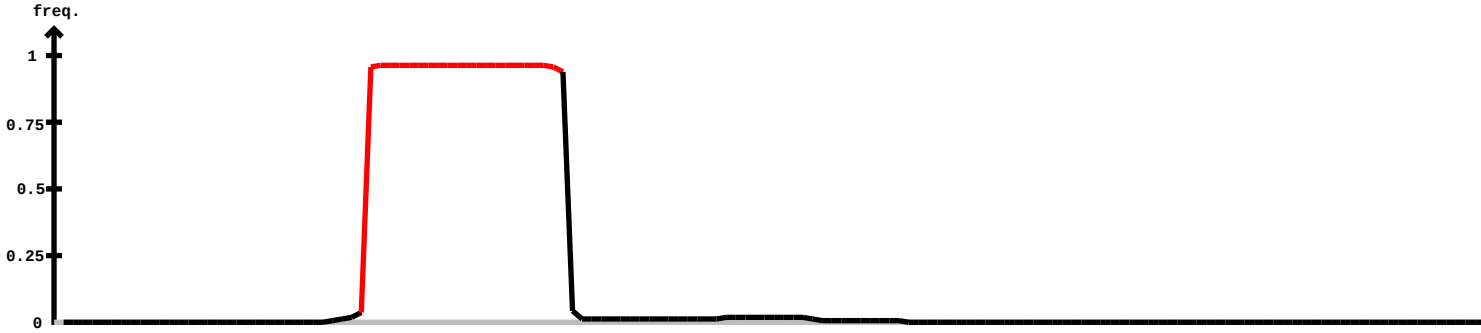
miRBase precursor : dme-mir-997_MI0005859
Total read count : 163
dme-miR-997-5p_MIMAT0005517 : 155 read count
dme-miR-997-3p_MIMAT0020897 : 2 read count
remaining reads : 5

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                                   dme-miR-997-5p_MIMAT0005517
5'- uuauggauccucuuucaaugaauuuaguaugcccaaacucgaaggaguuucaccuccauaagagcgacaguccuggagaaguuaucagagccaaaaaaauucauaugaugaugcauuuuccgucucugaaaacgucuucagcagaagu
    .....(((((...(((((((...(((((((..((((.(((((.(((((((((.((..((((((((((..(((((((((.(((.((((((.(...(((((.(((.....))).)))))..).)))))).))).)))))))..)).))
    ........................uaugcccaaacucgaaggaguuu.......................................................................................................
    .........................augcccaaacucgaaggaguuuca.....................................................................................................
    ........................ugcccaaacucgaaggaguuuc......................................................................................................
    ...........................Acccaaacucgaaggaguuuca....................................................................................................
    ............................cccaaacucgaaggaguuuc.....................................................................................................
    ............................cccaaacucgaaggCguuuc.....................................................................................................
    ............................cccaaacucCaaggaguuuca....................................................................................................
    ............................cccaaacucgaaggaguuCca....................................................................................................
    ............................cccaaacucgaaggGguuuca....................................................................................................
    ............................cccaaacucgaaggaguuuca....................................................................................................
    ............................cccaaacucgaaggaguCuca....................................................................................................
    ............................cccaaacucgaaggagAuuca....................................................................................................
    ............................cccaaacucgaaggaCuuuca....................................................................................................
    ............................cUcaaacucgaaggaguuuca....................................................................................................
    ............................cccaaacCcgaaggaguuuca....................................................................................................
    ............................cccaaacucgaaggaguuucaA...................................................................................................
    ............................cccaaacucgaaggaguuucac...................................................................................................
    ...........................ccaaacucgaaggaguuuca.....................................................................................................
    ...................................................ccuccauaagagcgacaguccugga.....................................................................
    ..................................................ccuccauaagagcgacaguccuggag.....................................................................
    ...................................................................aguccuggagaaguuauca.....................................................
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