miRBase precursor : chromosome_1_2029
Total read count : 115
chromosome_1_2029 read count : 49
remaining reads : 66

```
5'- aguacgagggguguguucgaaauguuaaacauuuccaacacucccucguacucg  -3'    exp
    ((((((((((.(((((.(((((((...))))))).))))).))))))))))..    reads    mm         sample
    aguacgagggguguguucgaaaC..............................    2        1          seq
    aguacgagggguguguucgCaau..............................    1        1          seq
    aguacgagggguguguucgaaau..............................    17       0          seq
    aAuacgagggguguguucgaaaug.............................    1        1          seq
    aguacgagggguguguucgaaaug.............................    24       0          seq
    aguacgagggguguguGcgaaaug.............................    1        1          seq
    ..uacgagggguguguucgaaaau.............................    2        0          seq
    ..uacgagggguguguucgaaaug.............................    1        0          seq
    ...................................uuuccaacacucccucgu.....    5        0          seq
    ...................................uuuccaacacucccucguacu..    8        0          seq
    ...................................uuCccaacacucccucguacu..    3        1          seq
    ...................................uuuccaacacucccucguacC..    1        1          seq
    ...................................uuuccaacacucccucguacuc.    3        0          seq
    ...................................uuuccCacacucccucguacuc.    1        1          seq
    ...................................uuuccaacacucccucguacuA.    1        1          seq
    ...................................uuuccaacacCcccucguacuc.    1        1          seq
    ...................................uuuccaacacucccucguacucA    8        1          seq
    ...................................uuuccaacacucccucguacucg    20       0          seq
    ....................................uuccaacacucccucguac...    1        0          seq
    ....................................uuccaacGcucccucguacu..    4        1          seq
    ....................................uuccaacacucccucguacC..    2        1          seq
    ....................................uuccaacacucccucguacu..    4        0          seq
    ....................................uuccaacacucccucguacucg    4        0          seq
```