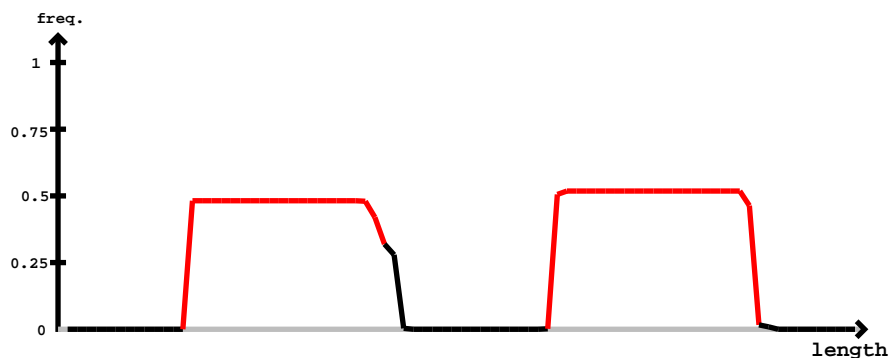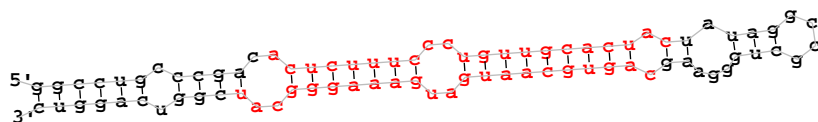miRBase precursor   : hsa-mir-130b
Total read count   : 548
hsa-miR-130b-5p read count 264
hsa-miR-130b-3p read count 284
remaining reads   : 0

hsa-miR-130b-3p

hsa-miR-130b-5p

```
5'- ggccugcccgacacucuuucccguuugcacuacuauaggccgcugggaagcagugcaaugaugaaagggcaucggucagguc   -3'    exp
    ((((((.((((..(((((((..(((((((((.((.(((....)))...)).)))))))))..)))))).)))).))))))    reads    mm         sample
    ...........acucuuucccguuugcac...................................................       1       0          seq
    ...........acucuuucccguuugcacu..................................................      29       0          seq
    ...........Ccucuuucccguuugcacu..................................................       3       1          seq
    ...........acucuuucccguuugcacA..................................................       1       1          seq
    ...........acucuuCcccguuugcacua.................................................       1       1          seq
    ...........acucuuucccguuugcacua.................................................      44       0          seq
    ...........acucuuucccguuugcacuG.................................................       1       1          seq
    ...........acCcuuucccguuugcacua.................................................       1       1          seq
    ...........Ccucuuucccguuugcacua.................................................       8       1          seq
    ...........acucuuucccguuugcacuaG................................................       9       1          seq
    ...........acucuuucccguuugcacuac................................................      13       0          seq
    ...........Ucucuuucccguuugcacuacu...............................................       1       1          seq
    ...........acucuuucccguuGgcacuacu...............................................       1       1          seq
    ...........acucuuucccguuugcacuacu...............................................     136       0          seq
    ...........acucuuucccguuugcacuacG...............................................       2       1          seq
    ...........Ccucuuucccguuugcacuacu...............................................      11       1          seq
    ...........acucuuucccguuugcacuacAa..............................................       2       1          seq
    ...........................................gcagugcaaugaugaaagggca.............         1       0          seq
    ............................................cagugcaaugaugaaagggc.............        29       0          seq
    ............................................cagugcaaugauAaaagggc.............         1       1          seq
    ............................................cagugUaaugaugaaagggca............         2       1          seq
    ............................................cagugcaaugauAaaagggca............         1       1          seq
    ............................................cagugcaaugaugaaagggcG............         1       1          seq
    ............................................cagugcaaugaugaaagggca............       230       0          seq
    ............................................Uagugcaaugaugaaagggca............         1       1          seq
    ............................................cagugcaaugaugCaagggca............         1       1          seq
    ............................................cagugcaCugaugaaagggca............         1       1          seq
    ............................................cagugcaaugaugaaagggcau...........         4       0          seq
    ............................................cagugcaaugaugaaagggcauU..........         4       1          seq
    ............................................cagugcaaugaugaaagggcauA..........         1       1          seq
    ............................................Cgugcaaugaugaaagggca.............         1       1          seq
    ............................................agugcaaugaugaaagggca.............         6       0          seq
```