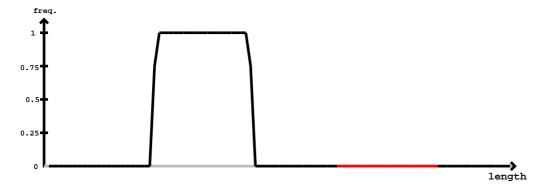
miRBase precursor : hsa-mir-607

Total read count : 4 hsa-miR-607 read count : 0 remaining reads : 4

uug ceuaaaaucaeaeaaauuauaaaucuagaeeuaa<mark>aeuu</mark>gaaaceeaag<sup>a</sup>ag a 5'3<sup>9</sup>99uuucaguguguuc<del>aauaucuagaeeuaa</del>euuggguce<sub>guc</sub>ag



## hsa-miR-607

5'- uugccuaaagucacacagguuauagaucuggauuggaacccagggagccagacugccuggguucaaauccagaucuauaacuugugugacuuuggg	-3' exp		
((((((((((((((((((((((((((((((((((((	reads	mm	sample
auagaucuggauuggaaccc	1	0	seq
auagaucuggauuggaaceca	2	0	seq
uagaucuggauuggaaccca	1	0	seq