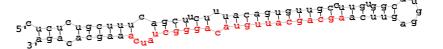
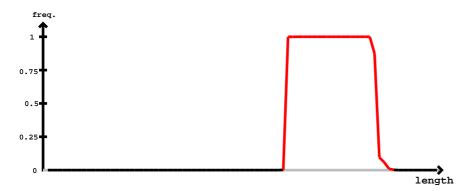
miRBase precursor : hsa-mir-107
Total read count : 2721
hsa-miR-107 read count : 2721
remaining reads : 0





hsa-miR-107

- cucucugcuuucagcuucuuuacaguguugccuuguggcauggaguuca <mark>agcagcauuguacagggcuauca</mark> aagcacaga	-3' exp		
.(((.((((.(((.(((.(((.(((.(())))))))	reads	mm.	sample
aagcagcauuguacagggc	2	0	seq
aagcagcauuguacagggcu	3	0	seq
aagcagcauuguacagggcua	1	0	seq
agcagcauuguaUagggc	3	1	seq
agcagcaCuguacagggc	1	1	seq
agcagcauuguacagggc	305	0	seq
aAcagcauuguacagggc	1	1	seq
Cgcagcauuguacagggc	27	1	seq
agcagcauugAacagggc	1	1	seq
agcagcaAuguacagggc	1	1	seq
agcagcauuguGcagggcu	2	1	seq
agcagcauuguacagAgcu	1	1	seq
agcagAauuguacagggcu	2	1	seq
agcagcauuguaUagggcu	2	1	seq
agcagcauuguacagggUu	4	1	seq
agcagcauuUuacagggcu	1	1	seq
agcagcauugAacagggcu	1	1	seq
agcagUauuguacagggcu	1	1	seq
agcagcauuguacUgggcu	1	1	seq
Cgcagcauuguacagggcu	177	1	seq
agcGgcauuguacagggcu	1	1	seq
agcagcauuAuacagggcu	4	1	seq
agUagcauuguacagggcu	2	1	seq
Ugcagcauuguacagggcu	2	1	seq
agcagcauuguacagggcG	17	1	seq
agcagcauAguacagggcu	2	1	seq
agcCgcauuguacagggcu	5	1	seq
agcagcauCguacagggcu	3	1	seq
Ggcagcauuguacagggcu	1	1	seq
agcagcauuguacCgggcu	3	1	seq
agcagcauuguacagggcu	1883	0	seq
agcagcauuguacagggcA	1	1	seq
agcagcauugCacagggcu	2	1	seq
agcagcaAuguacagggcua	1	1	seq
Cgcagcauuguacagggcua	10	1	seq

hsa-miR-107

cucucugcuuucagcuucuuuacaguguugccuuguggcauggaguucaagcagcauuguacagggcuaucaaagcacaga			
agcagcauuguacagggcua	83	0	seq
agcagcauuguacagggcuU	1	1	seq
agcagcauuCuacagggcuau	1	1	seq
Cgcagcauuguacagggcuau	11	1	seq
agcagcCuuguacagggcuau	1	1	seq
agcagcauuguacagggcuaA	3	1	seq
agcagcauuguacagggcuau	117	0	seq
agcagcauuguacagggcuauU	17	1	seq
agcagcauuguacagggcuauc	1	0	seq
agcagcauuguacagggcuauA	5	1	seq
agcagcauuguacagggcuaucU	1	1	seq
agcagcauuguacagggcuauAa	4	1	seq
agcagcauuguacagggcuauca	1	0	seq
cagcauuguacagggcua	1	0	seq