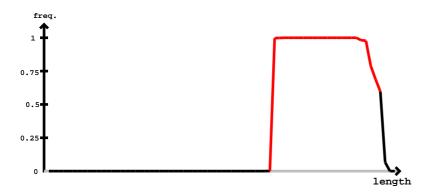
miRBase precursor : hsa-mir-941-3

Total read count : 711 hsa-miR-941 read count : 711 remaining reads : 0





hsa-miR-941

IISa-MIR-J41			
5'- ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca	-3'	exp	
((.((((((((((((((((((((((((((((((((((((reads	s mm	sample
Ccacccggcugugcacaugugc	1	1	seq
cacccggcugugcaca	13	0	seq
cacccggcugugcacau	3	0	seq
cacccggcugugcaUaug	1	1	seq
Uacccggcugugcacaug	1	1	seq
cCcccggcugugcacaug	1	1	seq
cacccggcugugcacaug	129	0	seq
caUccggcugugcacaug	1	1	seq
cacccggcugugcacaugA	4	1	seq
Aacccggcugugugcacaugu	1	1	seq
cacccggcugugUacaugu	1	1	seq
cacccggcugugcacaugu	65	0	seq
caccUggcugugcacaugug	1	1	seq
cacccggcugugugcacauguA	7	1	seq
cacccggcugugcacaugug	55	0	seq
cacccggcugugcacauguU	1	1	seq
cacccggcuguguAcacaugug	1	1	seq
cacccggcugugcacaugugU	79	1	seq
cacccggcugugugcacauAugc	1	1	seq
cUcccggcugugcacaugugc	1	1	seq
cacccggcuguguguacaugugc	1	1	seq
cacccggcugugcacaugugc	208	0	seq
cacccggcuAugugcacaugugc	1	1	seq
cacccggcugCgugcacaugugc	1	1	seq
cacccggcugugugcacaugugG	21	1	seq
cacccggcugugugGacaugugc	1	1	seq
cacccggcugugugcacaugugA	60	1	seq
cacccggcugugcacaugGgc	1	1	seq
cacccggcugugugcacauUugc	1	1	seq
cacccggcugugugcacaugugcU	47	1	seq
acccggcugugugcacaug	1	0	seq
cccggcugugugcacaugug	1	0	seq