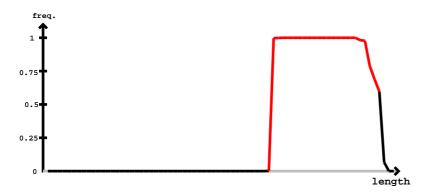
miRBase precursor : hsa-mir-941-4

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





hsa-miR-941

- ugugcacaugugcccagggcccgggacagcgccacggaagaggacg <mark>cacccggcugugugcacaugugc</mark> cca	-3' exp		
((.((((((((((((((((((((((((((((((((((((	reads	mm	sample
Ccacccggcugugugcacaugugc	1	1	seq
cacccggcugugugcaca	13	0	seq
cacccggcugugcacau	3	0	seq
Uacccggcugugugcacaug	1	1	seq
cacccggcugugugcaUaug	1	1	seq
cacccggcugugugcacaug	129	0	seq
caUccggcugugugcacaug	1	1	seq
cCcccggcugugugcacaug	1	1	seq
Aacccggcugugugcacaugu	1	1	seq
cacccggcugugugcacaugu	65	0	seq
cacccggcugugugcacaugA	4	1	seq
cacccggcugugUacaugu	1	1	seq
cacccggcugugugcacaugug	55	0	seq
cacccggcugugugcacauguA	7	1	seq
caccUggcugugugcacaugug	1	1	seq
cacccggcugugugcacauguU	1	1	seq
cacccggcuguguAcacaugug	1	1	seq
cacccggcugugugcacaugGgc	1	1	seq
cacccggcugugugcacaugugA	60	1	seq
cacccggcuAugugcacaugugc	1	1	seq
cacccggcugugugcacauUugc	1	1	seq
cacccggcugugugGacaugugc	1	1	seq
cacccggcugugugcacaugugG	21	1	seq
cacccggcugugugcacauAugc	1	1	seq
cacccggcugugugcacaugugc	208	0	seq
cacccggcugugugUacaugugc	1	1	seq
cUcccggcugugugcacaugugc	1	1	seq
cacccggcugCgugcacaugugc	1	1	seq
cacccggcugugugcacaugugU	79	1	seq
cacccggcugugugcacaugugcU	47	1	seq
acccggcugugugcacaug	1	0	seq
cccggcugugugcacaugug	1	0	seq