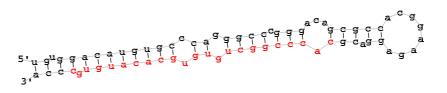
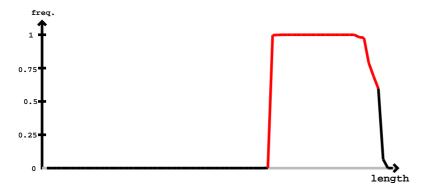
miRBase precursor : hsa-mir-941-1

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





hsa-miR-941

	IISA-MIR-941			
5'- uguggacaugugcccagggcccggg	acagcgccacggaagaggacg <mark>cacccggcugugugcacaugugc</mark> cca	-3'	exp	
((.((((((((.((.(((.(((	(((((()).))).))))))))))))))))))	read	s mm	sample
	Ccacccggcugugugcacaugugc	1	1	seq
	cacccggcugugugcaca	13	0	seq
	cacccggcugugugcacau	3	0	seq
	Uacccggcugugugcacaug	1	1	seq
	cacccggcugugugcaUaug	1	1	seq
	cCcccggcugugugcacaug	1	1	seq
	cacccggcugugugcacaug	129	0	seq
	caUccggcugugugcacaug	1	1	seq
	Aacccggcugugugcacaugu	1	1	seq
	cacccggcugugugUacaugu	1	1	seq
	cacccggcugugugcacaugA	4	1	seq
	cacccggcugugugcacaugu	65	0	seq
	cacccggcuguguAcacaugug	1	1	seq
	cacccggcugugugcacauguA	7	1	seq
	caccUggcugugcacaugug	1	1	seq
	cacccggcugugugcacaugug	55	0	seq
	cacccggcugugugcacauguU	1	1	seq
	cacccggcugugugUacaugugc	1	1	seq
	cacccggcugugugcacaugugU	79	1	seq
	cacccggcugugugcacaugugA	60	1	seq
	cacccggcugCgugcacaugugc	1	1	seq
	cacccggcugugugcacaugugG	21	1	seq
	cacccggcugugugcacauAugc	1	1	seq
	cacccggcugugugcacaugugc	208	0	seq
	cUcccggcugugugcacaugugc	1	1	seq
	cacccggcugugugcacaugGgc	1	1	seq
	cacccggcugugugGacaugugc	1	1	seq
	cacccggcuAugugcacaugugc	1	1	seq
	cacccggcugugugcacauUugc	1	1	seq
	cacceggcugugugcacaugugcU	47	1	seq
	acccggcugugugcacaug	1	0	seq
	cccggcugugugcacaugug	1	0	seq