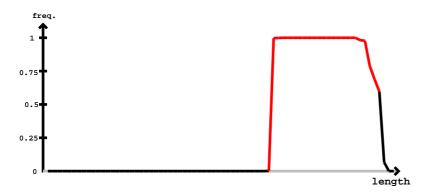
miRBase precursor : hsa-mir-941-5

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





hsa-miR-941

	iisa-mir-541			
5'- ugugca	caugugcccagggcccgggacagcgccacggaagaggacg <mark>cacccggcugugugcacaugugc</mark> cca	-3'	exp	
((.(((((((((((((((((((((((((((((((((((((((((reads	s mm	sample
	Ccacccggcugugcacaugugc	1	1	seq
	cacccggcugugcaca	13	0	seq
	cacccggcugugcacau	3	0	seq
	cacccggcugugcacaug	129	0	seq
	Uacccggcugugcacaug	1	1	seq
	cCcccggcugugcacaug	1	1	seq
	cacccggcugugcaUaug	1	1	seq
	caUccggcugugcacaug	1	1	seq
	cacccggcugugUacaugu	1	1	seq
	cacccggcugugcacaugu	65	0	seq
	Aacccggcugugugcacaugu	1	1	seq
	cacccggcugugcacaugA	4	1	seq
	cacccggcugugcacaugug	55	0	seq
	cacccggcuguguAcacaugug	1	1	seq
	cacccggcugugugcacauguA	7	1	seq
	cacccggcugugugcacauguU	1	1	seq
	caccUggcugugcacaugug	1	1	seq
	cacccggcugugugcacaugugU	79	1	seq
	cUcccggcugugcacaugugc	1	1	seq
	cacccggcugugcacaugugG	21	1	seq
	cacccggcugugugcacauUugc	1	1	seq
	cacccggcugugGacaugugc	1	1	seq
	cacccggcugugugcacaugugA	60	1	seq
	cacccggcuguguguacaugugc	1	1	seq
	cacccggcuAugugcacaugugc	1	1	seq
	cacccggcugCgugcacaugugc	1	1	seq
	cacccggcugugugcacaugGgc	1	1	seq
	cacccggcugugugcacauAugc	1	1	seq
	cacccggcugugugcacaugugc	208	0	seq
	cacccggcugugugcacaugugcU	47	1	seq
	acccggcugugugcacaug	1	0	seq
	cccggcugugugcacaugug	1	0	seq