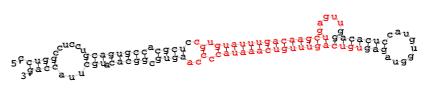
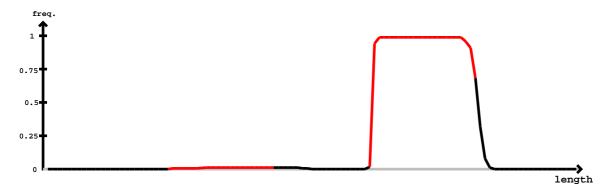
miRBase precursor : hsa-mir-223

Total read count : 340 hsa-miR-223-5p read count : 2 hsa-miR-223-3p read count : 336 remaining reads : 2





hsa-miR-223-5p

hsa-miR-223-3p

5'- ccuggccuccugcagugccacgcucc <mark>guguauuugacaagcugaguu</mark> ggacacuccaugugguagagu <mark>gucaguuugucaaauacccca</mark> agugcggcacaugcuuacca	ıg -3'	exp	
.((((((((((((((((((((((((((((((((() rea	nds mm	sample
.cguguauuugacaagcugaguuggaca	. 2	0	seq
.uuugCcaagcugaguuggacacu	. 1	1	seq
uugacaagcugaguuggacacu	. 1	0	seq
gugucaguuugucaaauacccc	. 2	0	seq
gugucaguuugucaaauacccca	. 3	0	seq
gugucaguuugucaaauaccccaU	. 1	1	seq
.ugucaguuugucaaauacc	. 10	0	seq
.ugucaguuugucaaauaccc	. 14	0	seq
.ugucCguuugucaaauaccc	. 2	1	seq
.ugucaguuugucaaauacccc	. 63	0	seq
.ugucCguuugucaaauacccc	. 7	1	seq
.ugucagGuugucaaauacccc	. 1	1	seq
.ugucCguuugucaaauacccca	. 10	1	seq
.ugucaguuugucaaauaccccG.	. 3	1	seq
.ugucaguuugucaaauaccccU	. 18	1	seq
.ugucGguuugucaaauacccca	. 1	1	seq
.ugucaguuugucaaauacccca	. 82	0	seq
.ugucaguuuAucaaauacccca	. 1	1	seq
.ugucaguuugucaaauaccccaU	. 11	1	seq
.ugucCguuugucaaauaccccaa.	. 11	1	seq
.ugucaguuugucaaauaccccaG	. 8	1	seq
.ugucaguuugucaaauaccccaa.	. 45	0	seq
	. 1	1	seq
	. 10	0	seq
.ugucaguuugucCaauaccccaag	. 1	1	seq
.ugucaguuugucaaauaccccaaU	. 6	1	seq
.ugucaguuugucaaauaccccaaC	. 2	1	seq
.ugucaguuugucaaauaccccaaA	. 2	1	seq
ugucaguuugucaaauaccccaUg	. 1	1	seq
ugucaguuugucaaauaccccaagu	. 1	0	seq
ugucaguuugucaaauaccccaagA	. 2	1	seq
.ugucaguuugucaaauaccccaaUu	. 1	1	seq
gucaguuugucaaauacc	. 1	0	seq
queaquuuqueaaauacccc.	. 4	0	sea

hsa-miR-223-5p

hsa-miR-223-3p

ccuggccuccugcagugccacgcuc<mark>cguguauuugacaagcugaguu</mark>ggacacuccaugugguagag<mark>ugucaguuugucaaauacccca</mark>agugcggcacaugcuuaccag

seq	1	1	gucaguuugucaaauaccccG
seq	0	5	.gucaguuugucaaauacccca
seq	0	4	.gucaguuugucaaauaccccaa
seq	1	1	gucaguuugucaaauaccccaaU