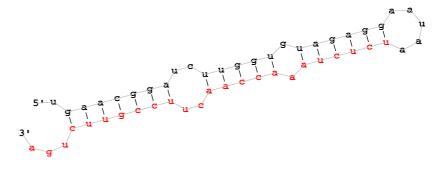
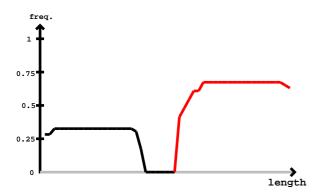
miRBase precursor : chromosome_4_15141

Total read count : 46 chromosome_4_15141 read x3 unt remaining reads : 15





chromosome_4_15141

'- ugaacggaucuugguguagaggaauaaucucuaaaccaacuuccguucuga	-3'	exp	
.((((((((((((((())))))).)))))	reads	s mm	sample
ugaacggaucuuggugua	1	0	seq
ugaacggaucuugguguag	5	0	seq
ugaacggaucuugguguUga	1	1	seq
ugaacggaucuugguguaga	5	0	seq
Agaacggaucuugguguaga	1	1	seq
.Aaacggaucuugguguag	1	1	seq
.gaacggaucuugguguagG	1	1	seq
ucucuaaaccaacuuccguucu	1	0	seq
Acucuaaaccaacuuccguucug.	1	1	seq
ucucuaaaccaacuuccguucuga	4	0	seq
uAucuaaaccaacuuccguucuga	2	1	seq
Gcucuaaaccaacuuccguucuga	2	1	seq
Acucuaaaccaacuuccguucuga	9	1	seq
cucuaaaccaacuuccguucuga	3	0	seq
ucuaaGccaacuuccguucuga	1	1	seq
ucuaaaccaacuuccguucuga	2	0	seq
cuaaaccaacuuccguucuga	3	0	seq
aaaccaacuuccguucuga	3	0	seq
	ugaacggaucuuggugua ugaacggaucuugguguaga ugaacggaucuugguguaga Agaacggaucuugguguaga .Aaacggaucuugguguagg .gaacggaucuugguguagg .ucucuaaaccaacuuccguucuga .ucucuaaaccaacuuccguucuga .ukucuaaaccaacuuccguucuga .ukucuaaaccaacuuccguucuga .ucucuaaaccaacuuccguucuga .ucucuaaaccaacuuccguucuga .ucucuaaaccaacuuccguucuga .ucucuaaaccaacuuccguucuga .ucucuaaaccaacuuccguucuga	. ((((((((((((((((((((((((((((((((((((.((((((((((((((((((((((((((((((((((((