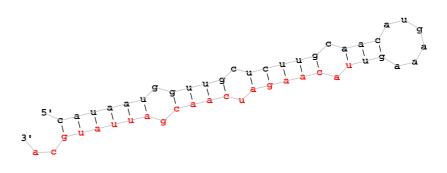
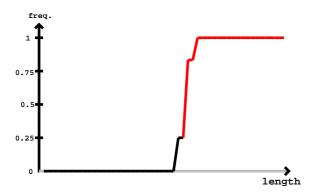
miRBase precursor : chromosome\_12\_57427

Total read count : 12 chromosome\_12\_57427 read tount remaining reads : 0





## chromosome\_12\_57427

5'- cauaaugguugcucuugcaacaugaaaguuacaagaucaacgauuaugca	-3' exp	<b>&gt;</b>	
(((((((((((((((((((((((((((((((((((((((	reads	mm	sample
guuacaGgaucaacgauuaugca	1	1	seq
Auuacaagaucaacgauuaugca	1	1	seq
guuacaagaucaacgUuuaugca	1	1	seq
uacaagaucaacgauuaugca	7	0	seq
Uaagaucaacgauuaugca	1	1	seq
caagaucaacgauuaugca	1	0	seq