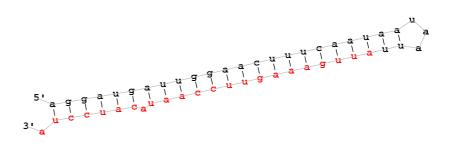
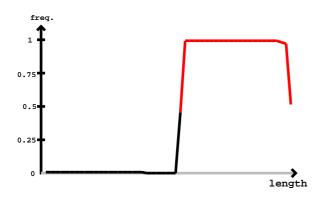
miRBase precursor : chromosome_12_61632

Total read count : 270 chromosome_12_61632 read **268**nt remaining reads : 2





chromosome_12_61632

aggaugauuggaacuuucaauaauaauuauugaaaguuccaauacauccua	-3' exp		
(((((((((((((((((((((((((((((((((((((((reads	mm	sample
aggaugauuggaacuuucaa	1	0	seq
Gggaugauuggaacuuucaa	1	1	seq
uauugaaaguuccaauacauccu.	55	0	seq
uauugaaaguuccaauacaucUu.	1	1	seq
uauugaaaguCccaauacauccu.	1	1	seq
uauCgaaaguuccaauacauccu.	1	1	seq
uauugaaaguuccGauacauccu.	1	1	seq
uCuugaaaguuccaauacauccu.	1	1	seq
uauugaaaguuccaCuacauccu.	1	1	seq
uauugaaaguuccaaCacauccu.	1	1	seq
uauugaaaguuccaauacaucAu.	1	1	seq
uauugaaaguucAaauacauccua	2	1	seq
uauuAaaaguuccaauacauccua	1	1	seq
uauugaaaguuccaauacauccuC	1	1	seq
uCuugaaaguuccaauacauccua	1	1	seq
uauugaaaguCccaauacauccua	1	1	seq
uaAugaaaguuccaauacauccua	1	1	seq
uauugaaaguuccaauacauccCa	1	1	seq
uGuugaaaguuccaauacauccua	1	1	seq
uauugaaaguuccaauacauccua	50	0	seq
uauugaaaguuccaauacaGccua	1	1	seq
auugaaaguuccaauacauc	3	0	seq
auugaaaguuccaauacaucc	3	0	seq
Guugaaaguuccaauacauccu.	4	1	seq
auugaaaguucGaauacauccu.	1	1	seq
auugaaGguuccaauacauccu.	1	1	seq
aCugaaaguuccaauacauccu.	1	1	seq
auugaaaguuccaauacauccu.	53	0	seq
auugaGaguuccaauacauccua	1	1	seq
aCugaaaguuccaauacauccua	1	1	seq
auugaaaguucUaauacauccua	1	1	seq
auugaaagCuccaauacauccua	3	1	seq
auugaaaguuccaauacauccua	61	0	seq
auugaaaguucAaauacauccua	1	1	seq
auugaaaguuccaGuacauccua	2	1	seq

chromosome_12_61632

aggaugauuggaacuuucaauaauaauu <mark>auugaaaguuccaauacauccua</mark>			
Guugaaaguuccaauacauccua	8	1	seq
auugaaGguuccaauacauccua	1	1	seq