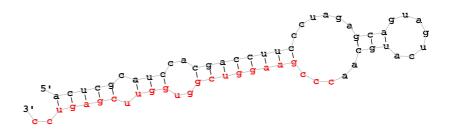
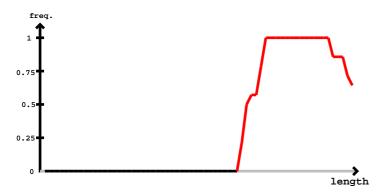
miRBase precursor : chromosome_12_63240

Total read count : 14 chromosome_12_63240 read &\u00e9unt remaining reads : 0





chromosome_12_63240

		-3' exp	5'- acucgcauccacgaccuucccuagagcaguagucaugcaacccgaaggucggugguucgagucc	5
sample	mm	reads	(((((((((((((((((((((((((((((((((((((((
seq	1	3	cccgaaggucgCugguucgagucc	
seq	1	2	ccgaaggucgUugguucg	
seq	1	1	ccgaaggucgCugguucgagu	
seq	1	1	ccgaaggucgCugguucgagucc	
seq	1	1	cgaaggucgCugguucgagu	
seq	1	1	Uaggucggugguucgaguc.	
seq	1	2	aaggucgCugguucgagucc	
seq	1	1	aggucggGgguucgagucc	
seq	1	1	aggucggugguucAagucc	
sea	0	1	aggucggugguucgagucc	