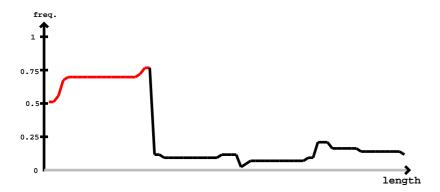
miRBase precursor : chromosome_14_108998

Total read count : 43 chromosome_14_108998 rea \mathfrak{G} 0ount remaining reads : 13





chromosome_14_108998

- gaaucucagaccucggccuugauguauugaccgagcgacagcgaggucaauacaucaaggccgaggucugagaa	-3'	exp	
((((((((((((((((((((((((((((((((((((reads	mm	sample
Caaucucagaccucggccuug	22	1	seq
.aaucucagaccucggccu	1	0	seq
.aaucucagaccucggccuug	1	0	seq
Cucucagaccucggccuug	1	1	seq
aucucagaccucggccuug	4	0	seq
ucucagaccucgAccuugau	1	1	seq
uugauguauugaccgagcgac	1	0	seq
ugauguauugaccgagcgaU	1	1	seq
ugauguauugaccgagcgac	2	0	seq
gacagcgaggucaauacaucaa	1	0	seq
gcgaggucaauacaucaa	1	0	seq
gaggucaauacaucaaggUcga	1	1	seq
aucaaggUcgaggucugaga.	1	1	seq
caaggccgaggucugagaG	1	1	seq
caaggccgaggucugagaU	4	1	seq