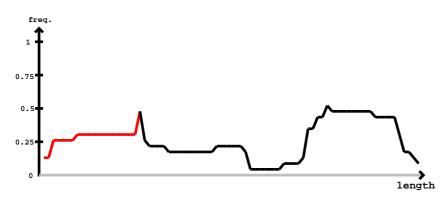
miRBase precursor : chromosome_2_6754

Total read count : 23 chromosome_2_6754 read count remaining reads : 16





chromosome_2_6754

	chromosome_2_6754			
5'-	${\tt gcgaauacagccgccucucau} {\tt ucgcucccgcuacuacuuaguaacgggagcgaaugagaggcggcuguauucgcaggc}$	-3'	exp	
	(((((((((((((((((((((((((((((((((((((((reads	s mm	sample
	Acgaauacagccgccucuca	2	1	seq
	Acgaauacagccgccucucau	1	1	seq
	.Ugaauacagccgccucuca	3	1	seq
	acagecgccucucauucCc	1	1	seq
	auucgcucccgcuacuacGuag	1	1	seq
	auucgcGcccgcuacuagu	1	1	seq
	auucgcAcccgcuacuagu	1	1	seq
	auucgcucccgcuacuagC	1	1	seq
	acuuaguaacgggagcgaaugaga	1	0	seq
	gGgaaugagaggcggcugu	1	1	seq
	augagaggcggcuguauucgU	1	1	seq
	ugagaggcggcuguauucA	3	1	seq
	ugagaggcggcuguauucgU	2	1	seq
	agaggcggcuguauucgcaU	1	1	seq
	agaggcggcuguauucgcagU.	1	1	seq
	aggcggcuguauucgcaggU	2	1	seq