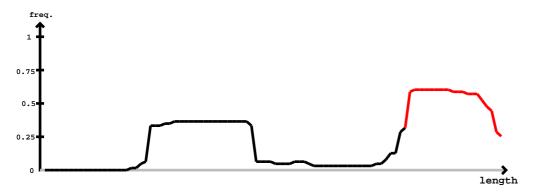
miRBase precursor : chromosome_11_51807

Total read count : 63 chromosome_11_51807 read **30**unt remaining reads : 33





chromosome_11_51807

5'	- aauaucccacuugcacuuguauuuucuucguauacacacucgccuaaaggcucguguguauaccgagaaaauacgagugacucgugggauauucc	-3'	exp		
	((((((((((((((((((((((((((((((((((((reads		mm	sample
	uguauuuucuCcguauacacacucgccuaa	1		1	seq
	uauuuucuucguauacacacucA	2		1	seq
	auuuucuucguauacacacucgc	1		0	seq
	uuuucuucguauacacacucgA	2		1	seq
	uuuucuucguauacacacucgU	2		1	seq
	uuuucuuGguauacacacucgc	8		1	seq
	uuuucuucguauacacacucgc	4		0	seq
	uuuucuucguauacacacucgcA	1		1	seq
	ucuuGguauacacacucgccuaaaggcucg	1		1	seq
	uucguauacacacucgccuaaaggcucgu	1		0	seq
	uaaaggcucguguauaccA	1		1	seq
	ucAuguguauaccgagaaa	1		1	seq
	agaaaauacAagugacucg	1		1	seq
	aaauacgagugacucgugA	1		1	seq
	auacAagugacucgugggau	3		1	seq
	uacAagugacucgugggaua	3		1	seq
	cAagugacucgugggauau	2		1	seq
		6		1	seq
		1		1	seq
		1		1	seq
	Aagugacucgugggauauu	1		1	seq
	gGgugacucgugggauauuc.	1		1	seq
	agugacucgugggauauu	1		0	seq
	agugacucgugggauauG	1		1	seq
	agugacucgugggauauucc	8		0	seq
	agugacucgugggauauucA	7		1	seq
	gugacucgugggauauucA	1		1	seq