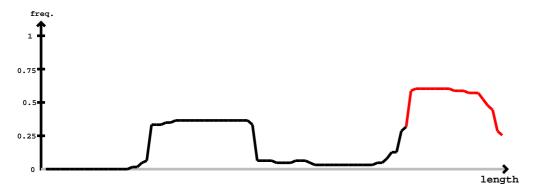
miRBase precursor : chromosome_11_51809

Total read count : 63 chromosome_11_51809 read **30**unt remaining reads : 33





chromosome_11_51809

٠-	a au au accea cuu gea cuu guau uu uu cuu cguau acaca cuc gecuaa ag geu cgu gu gu au acc ga ga aa au ac ga gu ga cuc gu gg ga uu au accea cuu gea cuu gu gu ga cuc gu ga ga aa au accea cuu gea cuu gu ga ga aa au accea cuu gea cuu gu ga ga aa au accea cuu gea cuu gu ga ga aa au accea cuu ga ga ga aa aa accea cuu ga ga ga accea cuu ga ga accea cuu ga accea cuu ga ga ga accea cuu ga ga ga accea cuu ga ga accea cuu ga ga ga accea cuu ga ga ga accea cuu ga ga accea cuu ga ga accea cuu ga ga accea cuu accea cuu ga accea cuu ga accea cuu accea	-3'	exp		
	((((((((((((((((((((((((((((((((((((reads	3	mm	sample
	uguauuuucuCcguauacacacucgccuaa	1		1	seq
	uauuuucuucguauacacacucA	2		1	seq
	auuuucuucguauacacacucgc	1		0	seq
	uuuucuuGguauacacacucgc	8		1	seq
	uuuucuucguauacacacucgA	2		1	seq
	uuuucuucguauacacacucgc	4		0	seq
	uuuucuucguauacacacucgU	2		1	seq
	uuuucuucguauacacacucgcA	1		1	seq
	ucuuGguauacacacucgccuaaaggcucg	1		1	seq
	uucguauacacacucgccuaaaggcucgu	1		0	seq
	uaaaggcucguguguauaccA	1		1	seq
	ucAuguguauaccgagaaa	1		1	seq
	agaaaauacAagugacucg	1		1	seq
	aaauacgagugacucgugA	1		1	seq
	auacAagugacucgugggau	3		1	seq
	uacAagugacucgugggaua	3		1	seq
		2		1	seq
	cgGgugacucgugggauauu	1		1	seq
		6		1	seq
		1		1	seq
		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