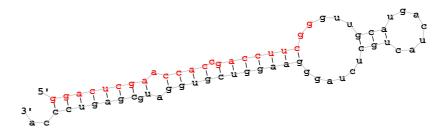
miRBase precursor : chromosome_12_59290

Total read count : 11 chromosome_12_59290 read &bunt remaining reads : 0





chromosome_12_59290

; · -	ggacucgaaccaccgaccuucggguugcaugacuacugcucuagggaaggucguggaugcgaguccca	-3'	exp		
	(((((((((((((((((((((((((((((((((((((((reads	s	mm	sample
	ggacucgaaccCccgaccu	1		1	seq
	ggacucgaaccaccgaccu	1		0	seq
	ggacucgaaccaGcgaccuu	1		1	seq
	ggacucgaaccaGcgaccuucgg	2		1	seq
	ggacucgaaccaGcgaccuucggg	2		1	seq
	.gacucgaaccaccgaccuA	1		1	seq
	acucgaaccaGcgaccuucgg	1		1	seq
	cgaaccaAcgaccuucgg	2		1	seq