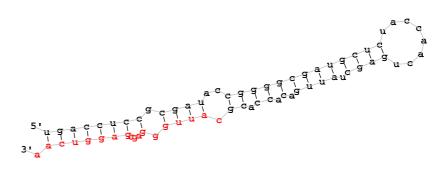
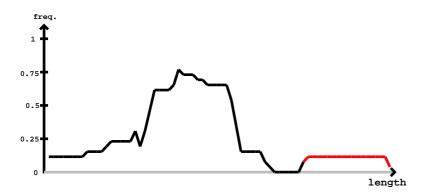
miRBase precursor : chromosome\_10\_45025

Total read count : 26 chromosome\_10\_45025 read &ount remaining reads : 23





## chromosome\_10\_45025

	0.11 0.110 0.				
5'-	ugaccuccgcgauaccggggcgaugcucuaccaacugagcuauugacaccacgcauugggagcgaggucaa	-3'	exp		
	(((((((((((((((((((((((((((((((((((((((	read	s	mm.	sample
	ugaccuccgcgauaccgg	1		0	seq
	ugaccuccgcgauaccgU	1		1	seq
	ugaccuccgcgauaccgA	1		1	seq
	cgcgauaccggUgcgaugcu	1		1	seq
	auaccggUgcgaugcucua	1		1	seq
	uaccggUgcgaugcucuacc	1		1	seq
	gggAgaugcucuaccaacuga	1		1	seq
	gggAgaugcucuaccaacugag	1		1	seq
	Ccgaugcucuaccaacug	1		1	seq
	gAgaugcucuaccaacug	2		1	seq
	Agaugcucuaccaacuga	2		1	seq
		1		1	seq
	cgaugcucuaccaacuga	1		0	seq
	gaCgcucuaccaacugag	2		1	seq
	gCugcucuaccaacugag	2		1	seq
	CucuaccaacugagcuauCga	1		1	seq
	ucuaccaacugagcuauC	2		1	seq
	ucuaccaacugagcuauug	1		0	seq
	Acauugggagcgagguca.	2		1	seq
	cauugggagcgaggucaa	1		0	seq