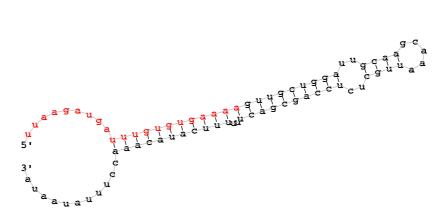
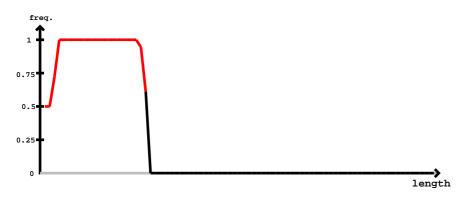
miRBase precursor : chromosome_4_12328

Total read count : 18 chromosome_4_12328 read x\dot{\dot{\dot{8}}nt} remaining reads : 0





chromosome_4_12328

5'-	${\color{blue} uuaagaugauuugugugaaaa} guugcuggauugcaagcaaauugcucuccagcgacuuuuuucauacaaaccuuuauaaua}$	-3' exp		
	((((((((((((((((((((((((((((((((((reads	mm	sample
	uuaagaugauuugugugaa	1	0	seq
	uuaagaugauuugugugaaG	1	1	seq
	uuaagaugauuugugugaaa	2	0	seq
	uuaagaugauuugugugaaGa	5	1	seq
	.uaagaugauuugugugaaG	1	1	seq
	.uaagaugauuugugugaaC	1	1	seq
	.uaagaugauuugugugaaGa	2	1	seq
	aagaugauuugugugaaG	1	1	seq
	.aagaugauuugugugaaGa.	4	1	seq