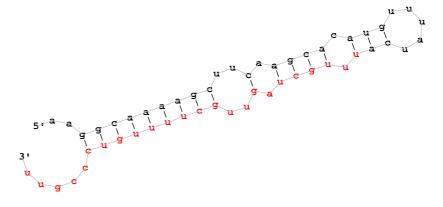
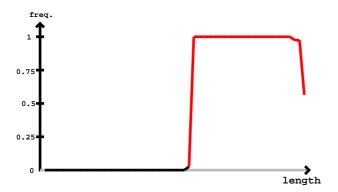
miRBase precursor : chromosome\_12\_60348

Total read count : 39 chromosome\_12\_60348 read **29**unt remaining reads : 0





5 '

## chromosome\_12\_60348

٠-	aaggcaaaagcuucaagcacauguuuaucauuugcuaguugcuuuugucccguu	-3'	exp	
	(((((((((((((((((((((((((((((((((((((	reads	s mm	sample
	Cuuugcuaguugcuuuugucccguu	1	1	seq
	uuugcuaguugcuuuuguccc	1	0	seq
	uuugcuaguugcuuuugucccgC.	2	1	seq
	uuugcuGguugcuuuugucccgu.	1	1	seq
	uuugcuaguugcuuuugucccgu.	9	0	seq
	uCugcuaguugcuuuugucccgu.	3	1	seq
	uuugcuaguugcuuuugucccgA.	1	1	seq
	Cuugcuaguugcuuuugucccguu	2	1	seq
	uuugcuaguugcuuuugucccguC	1	1	seq
	uuugcuaguugcuuuugucccguG	1	1	seq
	uuugcuaguugcCuuugucccguu	1	1	seq
	uCugcuaguugcuuuugucccguu	3	1	seq
	uuugcuaguugcuuuugucccguu	13	0	seq