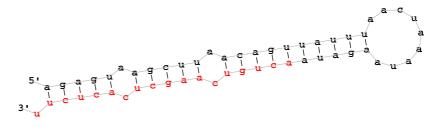
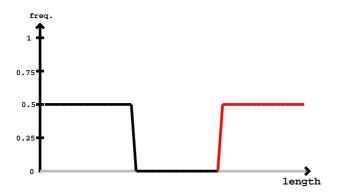
miRBase precursor : chromosome\_3\_7918

Total read count : 2 chromosome\_3\_7918 read count remaining reads : 1





## chromosome\_3\_7918

٠-	agaguaagcuuaacaguuauuuaacuaaauaagauaacugucaagcucacucuu	-3'	exp		
	(((((((((((((((((((((((((((((((((((((((	reads	3	mm	sample
	agaguaagcuuaacaguu	1		0	seq
	acugucaagcucacuUuu	1		1	seq