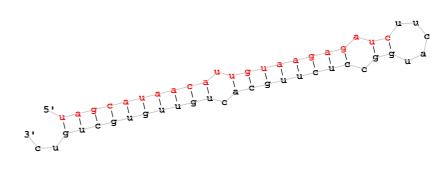
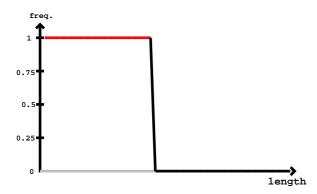
miRBase precursor : chromosome\_8\_34859

Total read count : 18 chromosome\_8\_34859 read  $x \cdot b \cdot 8$ nt remaining reads : 0





## chromosome\_8\_34859

5'-	uagcauaacauuguaagagaucuucauggccucuugcacuguugugcuguc	-3'	exp		
	(((((((((((((((((((((((((((((((((((((((	read	5	mm	sample
	uagcauaacauuguaagagauc	18		0	seq