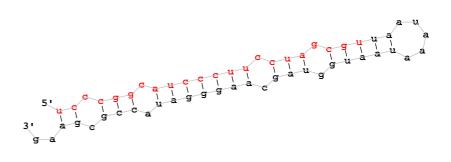
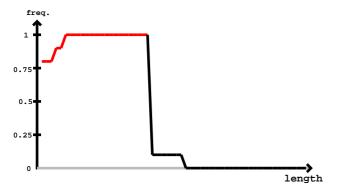
miRBase precursor : chromosome_13_64300

Total read count : 10 chromosome_13_64300 read \(\text{ Qount} \) remaining reads : 1





chromosome_13_64300

٠-	ucccggcaucccuuccuagcguuaauaaauagguagcaagggauaccgcgaag	-3' exp	P	
	((.(((.(((((((((((((()))))).))))))))	reads	mm.	sample
	ucccggcaucccuuccuagcgu	7	0	seq
	ucccggcaucccuuccuagcguuaauaaa	1	0	seq
	ccggcaucccuuccuagcgu	1	0	seq
	Agcaucccuuccuagcgu	1	1	seq