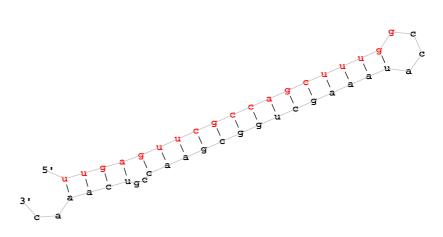
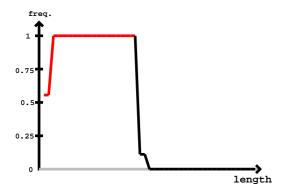
miRBase precursor : chromosome_10_47183

Total read count : 9 chromosome_10_47183 read \(\text{ Qount} \) remaining reads : 0





chromosome_10_47183

- uugaguucgeeageuuuggeeauaaageuggegaacegueaaae	-3. exp		
(((((((((((((((()))))))))))))))))))	reads	mm	sample
uugaguucgccagcuuugg	5	0	seq
.ugaguucgccagcuuugg	3	0	seq
.ugaguucgccagcuuuggcc	1	0	seq