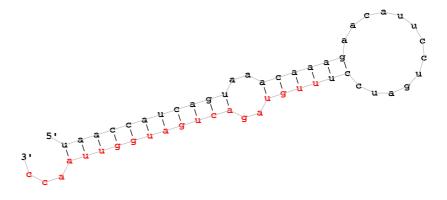
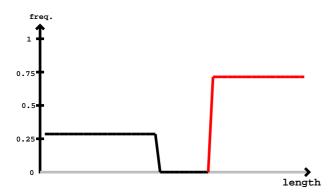
miRBase precursor : chromosome\_12\_60878

Total read count : 7 chromosome\_12\_60878 read **5** ount remaining reads : 2





## chromosome\_12\_60878

- uaaccaucaguaaacaaagaacauuccugauccuuuguagacugaugguuaacc	-3' exp		
(((((((((((((((())))))))))))))))	reads	mm	sample
uaaccaucaguaaacaaagaaca	2	0	seq
uuguagacugaugguuaacA	5	1	seq