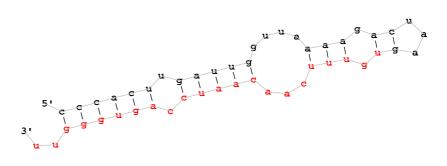
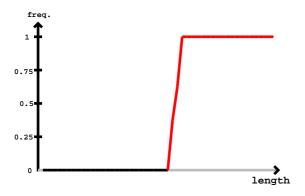
miRBase precursor : chromosome\_4\_13439

Total read count : 8 chromosome\_4\_13439 read x8unt remaining reads : 0





## chromosome\_4\_13439

5'- cccacuugauugguuaaaagacuaaguguuucaacaauccaguggguu	-3' exp		
(((((((((((()).)))))))))))))	reads	mm	sample
uguuucaacaauccaguggguu	3	0	seq
guuucaacaauccaguggguu	2	0	seq
uuucaacaauccaguggguu	3	0	seq