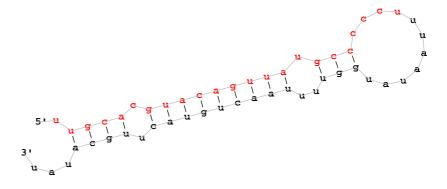
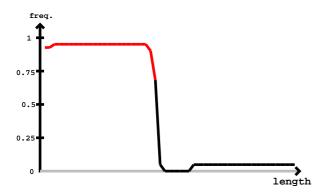
miRBase precursor : chromosome_11_51894

Total read count : 41 chromosome_11_51894 read **29**unt remaining reads : 2





chromosome_11_51894

5'- uugcacguacaguuaugcccccuuuaauaugguuuuaacuguacuugcauau	-3'	exp	
.((((.(((((((((())))))))))))))	reads	mm	sample
uugcacguacaguuaugcccA	2	1	seq
uugcacguacaguuaugccccU	2	1	seq
uugcacguacaguuaugccccc	2	0	seq
uugcacguacUguuaugccccc	1	1	seq
uugcacguacaguuaugccccA	3	1	seq
uugcacguacaguuaugcccccA	1	1	seq
uugcacguacaguuaugcccccu	21	0	seq
uugcacguacaguuaugcccccC	4	1	seq
uugcacguacaguuaugccccAuu	1	1	seq
uugcacguacaguuaugcccccuu	1	0	seq
.ugcacguacaguuaugccccc	1	0	seq
gguuuuaacuguacuugcauau	2	0	seq