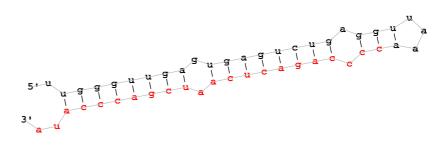
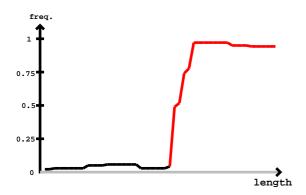
miRBase precursor : chromosome\_6\_23574

Total read count : 139 chromosome\_6\_23574 read xbaht remaining reads : 8





## chromosome\_6\_23574

5'- uuggguugagugagucugagguuaaaccccagacucaaucgacccaua	-3' ex	rp q	
.(((((((((((((((((())).))))))))))))	reads	mm	sample
uuggguugagugagucuga	3	0	seq
.uggguugaAugagucuga	1	1	seq
agugagucugagguuaaaccccagacucaa	3	0	seq
Ugucugagguuaaaccccagacucaaucga	1	1	seq
accccagacucaaucgacccaua	2	0	seq
ccccagacucaaucgacccUua	1	1	seq
ccccagaGucaaucgacccaua	2	1	seq
ccccagGcucaaucgacccaua	2	1	seq
Ucccagacucaaucgacccaua	6	1	seq
ccccagacucaaucgaAccaua	1	1	seq
ccccagacucaGucgacccaua	1	1	seq
ccccagacucaaucgacccGua	1	1	seq
ccccagacucaaucgacccaua	46	0	seq
ccccagCcucaaucgacccaua	1	1	seq
ccccagacucaaucgaUccaua	1	1	seq
cccagacucaaucgacccaua	4	0	seq
ccagacucaaucgacccaua	28	0	seq
Ucagacucaaucgacccaua	3	1	seq
cagacucaaucgacccaua	4	0	seq
Uagacucaaucgacccaua	1	1	seq
agacucaaucgacccaua	22	0	seq
Ggacucaaucgacccaua	4	1	seq
agacucaaucgacAcaua	1	1	seq