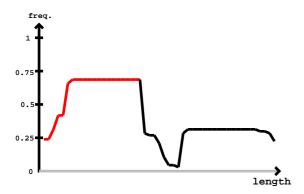
miRBase precursor : chromosome_13_70720

Total read count : 67 chromosome_13_70720 read **44**unt remaining reads : 23





chromosome_13_70720

ugcguugcagacgaacaaaaccccagggcuuuguucgucugcaacgca	-3' e:	кp	
(((((((((((((((((((((((((((((((((((((((reads	mm	sample
ugcguugcagacgaGcaaaa	1	1	seq
ugcguugcagacgaacaaaG	5	1	seq
Agcguugcagacgaacaaaa	2	1	seq
ugcguugcagacgaacaaaa	6	0	seq
ugcguuAcagacgaacaaaa	1	1	seq
ugcguugcagacgaaAaaaa	1	1	seq
.gcguugcagacgaacaaaa	4	0	seq
.Acguugcagacgaacaaaa	1	1	seq
Aguugcagacgaacaaaa	5	1	seq
Gguugcagacgaacaaaa	1	1	seq
cguugcagacgaacaaaac	1	0	seq
uugcagacgaacUaaaccc	2	1	seq
uugcagacgaacaaaaccc	1	0	seq
uugcagacgaacaaaacccc	1	0	seq
uugcagacgaacaaaacccA	6	1	seq
uugcagacgaacaaaacccca	3	0	seq
uugcagacgaacaaaaccccU	1	1	seq
uugcagacgaacaaaaccccGgg	1	1	seq
uugcagacgaacaaaaccccCgg	1	1	seq
ugcagacgaacaaaaccc	1	0	seq
ugcagacgaacaaaacccc	1	0	seq
agggUuuuguucgucugcaa	1	1	seq
gUuuuguucgucugcaacg	1	1	seq
	4	1	seq
Uuuuguucgucugcaacgca	8	1	seq
cuuuguucgucugcaacgca	5	0	seq
uuuguucgucugUaacgca	2	1	seq
	ugcguugcagacgacaaaa ugcguugcagacgaacaaaa ugcguugcagacgaacaaaa ugcguugcagacgaacaaaa ugcguugcagacgaacaaaa ugcguugcagacgaacaaaa ugcguugcagacgaacaaaa .Aguugcagacgaacaaaa .Aguugcagacgaacaaaa .Aguugcagacgaacaaaa .Gguugcagacgaacaaaauugcagacgaacaaaacuugcagacgaacaaaaccuugcagacgaacaaaacccuugcagacgaacaaaacccuugcagacgaacaaaacccc	((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((((